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Juan Pablo Bello, Elaine Chew, Douglas Turnbull (eds.)

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Preface

We have 105 papers accepted as posters, of which 24 will be also presented as plenary talks. There were 173 submissions.

(ADD MAIN BODY OF PREFACE HERE.)

Juan Pablo Bello, Elaine Chew
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Session 1 - Plenary Talks

A TUNNELING-VANTAGE INDEXING METHOD FOR NON-METRICS

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ABSTRACT

We consider an instance of the Earth Mover's Distance (EMD) useful for comparing rhythmical patterns. To make searches for r-near neighbours efficient, we decompose our search space into disjoint metric subspaces, in each of which the EMD reduces to the l_1 norm. We then use a combined approach of two methods, one for searching within the subspaces, the other for searching between them. For the former, we show how one can use vantage indexing without false positives nor false negatives for solving the exact r-near neighbour problem, and find an optimum number and placement of vantage objects for this result. For searching between subspaces, where the EMD is not a metric, we show how one can guarantee that still no false negatives occur, and the percentage of false positives is reduced as the search radius is increased.

1 INTRODUCTION

Searching a database for a melody or rhythm is often equivalent to a special version of the nearest neighbour problem: one wants to find items which are similar, but not necessarily identical to a given query, and be sure to retrieve all such items up to a certain level of dissimilarity. With a distance measure which adequately captures similarity, this amounts to retrieving all items which lie inside a ball around the query, where the radius of the ball is the dissimilarity threshold up to which retrieval results are considered relevant. We will call this search radius r, and items whose distance from a query is at most r will be called *r-near neighbours*.

If the database is large, one needs a data structure which makes it possible to retrieve the r-near neighbours without comparing the query to each point in the database. For high numbers of dimensions, where exact methods such as kd-trees do not offer much improvement over a linear search, a number of approximate methods have been suggested. Andoni and Indyk [2] point to many approximate methods and describe one of the most popular among them, *locality sensitive hashing* (LSH), in detail. With LSH, one applies a hash function to every data point. The hash function must be chosen so that there is a high probability for points which are close to each other to be hashed into the same bin. To search for nearest neighbours, one then applies the hash function to the query and searches the bin into which the query was hashed. However, LSH relies on the distance measure being a metric.

We describe a method which combines *vantage indexing* with *tunneling*, an approach applicable to certain non-metric distance measures. Our specific distance measure is non-metric; however, we can decompose our search space into disjoint metric subspaces. Within the metric subspaces, the non-metric reduces to the l_1 norm. We use a new variant of the vantage indexing method which can be used for solving the exact r-near neighbour problem (i. e., without resorting to approximation) for the l_1 norm in moderately high dimensional spaces (no more than about 9 dimensions). It guarantees to retrieve exactly the items within the ball of interest, without any false positives or false negatives, with a time complexity of essentially $\mathrm{O}(\log n)$ (n=number of data points), but at the cost of using $\mathrm{O}(n2^{j-1})$ storage space (j=number of dimensions). It is possible to use less storage space, but at a cost: false positives can occur.

For certain locality sensitive hash functions, LSH is similar to vantage indexing. Andoni and Indyk propose hash functions for the l_1 and l_2 norms which involve calculating the distances to an arbitrary number of randomly chosen vantage objects and then putting all data points into the same bin whose distances to the vantage objects lie in a certain range. We, on the other hand, use exactly as many vantage objects as needed, placed in the space so that we are guaranteed to retrieve all data points we should retrieve, and no others. Also, we avoid the need to pick a bin size which might not fit very well with the actual search radius.

While our method for vantage indexing is specific to instances where the EMD reduces to metric subspaces with regular ball shapes, the second part of our combined method is more generally applicable to prametric spaces that can be decomposed into metric subspaces. By "tunneling", we can efficiently search between metric subspaces, still without introducing false negatives. False positives can occur, but their percentage among the retrieved items shrinks as the search radius grows, making them tolerable.

2 MEASURING MELODIC OR RHYTHMIC SIMILARITY WITH THE EMD

2.1 The Earth Mover's Distance

The distance measure discussed in this paper measures the distance between weighted point sets. Intuitively speaking,

[1] R. Typke gratefully acknowledges support by the Austrian Research Fund (FWF), Project Number M1027-N15.

a weighted point set a_i can be imagined as an array of piles of dirt each equal to w_i units, situated at position x_i. The role of the supplier is arbitrarily assigned to one array and that of the receiver to the other one, and the arrays of piles are made to look as similar as possible by shifting dirt from piles in the supplier array to piles in the receiver array. The Earth Mover's Distance (EMD) then measures the minimum amount of work needed to make two arrays of piles as similar as possible in this way. See [3] for a more detailed description of the EMD. We now define the EMD formally:

Definition. Fix a *ground distance* d on $\mathbb{R}^k$. The ground distance can, but need not be, a metric.

Let $A = \{a_1, a_2, \ldots, a_m\}$, $B = \{b_1, b_2, \ldots, b_n\}$ be *weighted point sets* such that $a_i = \{(x_i, w_i)\}, i = 1, \ldots, m$, $b_j = \{(y_j, v_j)\}, j = 1, \ldots, n$, where $x_i, y_j \in \mathbb{R}^k$ with $w_i, v_j \in \mathbb{R}^+ \cup \{0\}$ being the respective weights.

Let $W_A = \sum_{j=1}^{n} w_i$ be the total weight of set A; the total weight W_B of the set B is defined analogously.

Let $d_{ij} = d(x_i, y_j)$ denote the ground distance between individual coordinates in A and B, without regard to their weight.

A *flow matrix* $F = (f_{ij})$ between A and B is an $m \times n$ matrix of non-negative real numbers, such that for each $1 \leqslant i \leqslant m$, $\sum_{j=1}^{n} f_{ij} \leqslant w_i$, and for each $1 \leqslant j \leqslant n$, $\sum_{i=1}^{m} f_{ij} \leqslant v_j$. Furthermore, we require that $\sum_i \sum_j f_{ij} = \min(W_A, W_B)$. Denote by $\mathcal{F}$ the collection of all possible flow matrices between A and B.

The *Earth Mover's Distance*, $\mathrm{EMD}_d(A, B)$, between A and B is defined as

$$\mathrm{EMD}_d(A, B) = \frac{\min_{F \in \mathcal{F}} \sum_{i=1}^{m} \sum_{j=1}^{n} f_{ij} d_{ij}}{min(W_A, W_B)}.$$

For the remainder of this paper, we will assume that the ground distance for the EMD is the Euclidean metric l_2. With this assumption in mind, we drop the subscript referring to the ground distance from our formulas, writing $\mathrm{EMD}(A, B)$ instead of $\mathrm{EMD}_{l_2}(A, B)$.

The EMD is a useful measure for music similarity, as was demonstrated at the annual MIREX comparison of music retrieval algorithms in 2006. Useful properties of the EMD include its continuity, its ability to support partial matching, and its robustness against distortions of tempo and pitch when measuring melodic similarity for symbolically encoded music. For doing so, one can represent every note by a point in the two-dimensional space of onset time and pitch. The weights of points can be used to encode the importance of notes. See [6] for details.

The EMD has one major disadvantage which can cause technical difficulties: it is in general not a metric. Specifically, the triangle inequality does not hold, and while the EMD of a point to itself is 0, there can exist distinct points that are also EMD 0 from one another. While there is no universally accepted terminology in the mathematical literature for weak distance measures, there is some precedent for calling weak distance measures with properties like the EMD *symmetric prametrics*.

It should be emphasized that the EMD does behave as a metric if one restricts the domain of the EMD to point sets having a given weight, assuming that the ground distance is a metric [3]. One can take advantage of this property when working with an EMD which is a prametric by decomposing the space of possible point sets into subspaces each containing only point sets having a given weight. We will refer to such subspaces as *metric subspaces* of the EMD space.

2.2 Comparing symbolic rhythmic patterns: a nice special case

The instance of the EMD described in this subsection is of particular interest for searching rhythmic patterns. We call this EMD the *Manhattan EMD*, and note that it is well-behaved from a geometric point of view.

Rhythmic patterns can naturally be represented as sequences of onset times. See Figure 1 for an illustration. To render irrelevant a musical segment's tempo and location within a piece of music, we scale their numeric representations to a fixed duration (say, 60) and translate them so that they start at position 0.

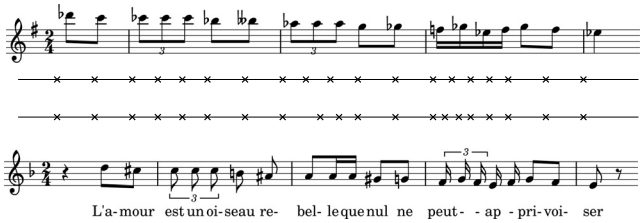

Figure 1. Comparing rhythms using sequences of onset times. *Top:* Ferruccio Busoni: Sonatina No.6: Chamber Fantasia on Bizet's Opera Carmen (3rd theme); *bottom:* Georges Bizet: Carmen: Habanera; *middle:* two sequences of onset times representing the incipits at the top and bottom. A possible numeric representation of the two series of onset times: **Busoni**: (0, 3, 6, 8, 10, 12, 15, 18, 20, 22, 24, 27, 30, 31.5, 33, 34.5, 36, 39, 42); **Bizet**: (0, 3, 6, 8, 10, 12, 15, 18, 21, 22.5, 24, 27, 30, 31, 32, 33, 34.5, 36, 39, 42).

When comparing normalized sequences of onsets containing the same number of onsets, the Manhattan EMD equals the sum of the absolute differences between corresponding onsets a_i and b_i in onset sequences $A = a_1 \ldots a_n$ and $B = b_1 \ldots b_n$, divided by the number of onsets: $\mathrm{EMD}(A, B) = \frac{\sum_{i=1}^{n} |a_i - b_i|}{n}$. Thus, if we restrict ourselves to point sets of a certain given length, the Manhattan EMD (with l_2 as ground distance) is a metric and is equal to the l_1 norm (also known as "Manhattan norm").

Since every normalized segment starts with 0 and ends with 60, we omit these two numbers and view n-onset segments as points in an $(n-2)$-dimensional space. All seg-

ments lie in the subset of $\mathbb{R}^{n-2}$ where the coordinates are strictly increasing.

3 VANTAGE INDEXING FOR MANHATTAN EMD

Vantage indexing (introduced by Vleugels and Veltkamp [7]) is an approach to the retrieval of objects from a large database which are similar to a given query. The search is restricted to items whose pre-calculated distances to a small set of pre-chosen *vantage objects* are similar to the query's distances to the same vantage objects.

3.1 Ball shapes

If one works with the l_1 norm, a ball (the set of all points whose distance lies within a certain radius around a point of interest) has the shape of a cross-polytope. A one-dimensional cross-polytope is a line segment, a two-dimensional cross-polytope is a square, for three dimentions, an octahedron, and so forth.

3.2 Creating a ball by intersecting "onion layers"

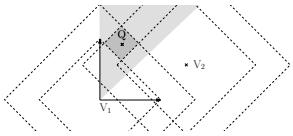

Figure 2. In this two-dimensional example, only the light gray area is inhabited by database objects since only there the second coordinate is greater than the first. The dark gray ball around Q with a given radius can be created by intersecting onion layers around V_1 and V_2 whose thickness equals the radius.

We call the area between the surfaces of two balls of differing radii around a vantage object an "onion layer".[1] An onion layer contains all items in the database whose distance from the vantage object lies within a certain range. Searching an onion layer can be done efficiently if one has built an index (for example, a range tree) for the database column that contains the distance to a vantage object. By building an index for multiple columns containing distances to different vantage objects (for example, a nested range tree), one can also search intersections of onion layers efficiently.

If one can cover the ball around the query object whose radius equals the search radius by using intersections of onion layers in a way that the intersection of onion layers equals exactly the ball of interest, neither false negatives nor false positives will occur. Since the intersection of onion layers can be searched efficiently, the ball of interest can as well. See Figure 2 for an illustration.

Certain metric spaces have the property that there exists a finite number of points $x_1, \ldots, x_n$ such that $\bigcap_i (B(x_i, R + r) \setminus B(x_i, R)) = B(y, r)$ for some point y (here we denote the ball with radius r around a point x with $B(x, r)$). In other terms, some spaces have the property that one needs only finitely many vantage objects so that the intersection of onion layers is exactly that of a ball. The l_1 space has this property, as do other spaces whose balls are regular objects with finitely many faces. An example of a metric space that does not have this property is Euclidean space.

[1] Inspiration has many sources, and onions have other good uses [5].

3.3 Optimum placement and number of vantage objects necessary for optimum indexing

In the case of one dimension, one vantage object is enough to always achieve the desired effect of perfectly covering only the interesting ball with onion layers. However, one has to place the vantage object correctly. See Figure 3 for good and bad examples.

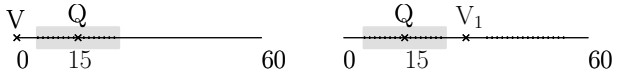

Figure 3. Of interest is the ball around Q (at 15) with radius 10, shown as a grey bar. *Left:* This ball is identical to the intersection of an onion layer around V with distance range of 5 to 25 and the inhabited part of the space. *Right:* If the vantage object is not put either to the left or to the right of inhabited space, corresponding onion layers can intersect with the inhabited space in two disjoint places (shown as dotted lines) – one would need to use another vantage object for the intersection of onion layers to be identical to the ball of interest.

For two dimensions, two vantage objects are necessary, and they are sufficient if they are placed, for example, like in Figure 2. Here, the onion layers intersect in two square-shaped regions, but only one of them (the one in the gray area) can ever contain data points. Therefore, no false positives can occur.

One can cover exactly the ball of interest and no other region (except for regions outside the inhabited space) by using 2^{j-1} vantage objects (j is the number of dimensions of the space) and placing them as follows:

- Let m be some number which is larger than any coordinate that can ever occur in the inhabited subspace (in the Bizet and Busoni example from Section 2.2, we could let $m = 60$).
- For one dimension, place one vantage object at the coordinate (0) (as illustrated in Fig. 3, left).
- For j dimensions, use twice as many vantage objects as for $j - 1$ dimensions. The coordinates of the vantage objects for j dimensions are based on those for $j - 1$ dimensions: for each vantage object position in $j - 1$ dimensions, create two new lists of coordinates for j dimensions by appending 0 or m, respectively,

to the list of coordinates for $j-1$ dimensions.

We need at least 2^{j-1} vantage objects because a cross-polytope has 2^j facets (a *facet* is a $(j-1)$-dimensional structure). For the intersection of onion layers to equal the ball of interest, every facet of the ball needs to coincide with the surface of an onion layer. One can use as few vantage objects as possible if one covers opposite facets of the cross-polytope with the surfaces of an onion layer around the same vantage object. This is exactly what happens if one uses 2^{j-1} vantage objects and places them as above.

4 PARTIAL MATCHING BY TUNNELING BETWEEN METRIC SUBSPACES

When searching sequences of onsets that were detected in audio files, there are two problems: the detection is not always completely reliable, producing both false negatives and false positives, and often there will be multiple voices with onsets, while a user might want to search for rhythmic patterns which occur only in one voice. Therefore, for successfully searching sequences of onsets from audio data, one should be able to ignore a certain number of onsets and still find matching subsequences or supersequences. In this section, we will show how one can use tunnels between metric subspaces to achieve partial matching while still benefiting from the optimum vantage indexing within the subspaces of sequences with equal numbers of onsets.

The EMD provides partial matching as described above for point sets whose weight sums are unequal. Unfortunately, the EMD does not obey the triangle inequality in such a case. This makes it impractical to directly apply vantage indexing since there would be no way of controlling the number of false negatives. Also, the locality sensitive hashing method, which also relies on the triangle inequality, becomes unusable.

Our approach is inspired by the case of the Manhattan EMD, which is rather easy to visualize. In particular, one can precisely visualize the neighbourhood, or "ball" of r-near neighbours in the n-dimensional metric subspace for an m-dimensional query point: assume that $(q_1, q_2, \ldots, q_m)$ is a query, where the initial 0 and final 60 are dropped, so that the query is an m-dimensional object.

First, we visualize the neighbourhoods in higher-dimensional metric subspaces, say of dimension $m+i$. In this case, the set of points which are of distance 0 from the query comprise $\binom{m+i}{i}$ many i-dimensional hyperplanes which are parallel to axes, and pass through the points that are the various possible completions of the query vector to $m+i$ dimensional space using i-many zeros. Then, the r-near neighbours of the query are found in the union of the r-balls, as calculated in the $m+i$-dimensional metric subspace, around each of the points of distance 0. The shape of these neighbourhoods is rather awkward, and is no longer a ball, but rather the cartesian product of a ball and a hyperplane. This shape is difficult to search efficiently. The reader may wish to think about the case of $m = 2$ and $i = 1$.

The neighbourhoods in lower-dimensional metric subspaces are a easier to describe. The points of distance 0 from the query in the $m-i$-dimensional metric subspace are simply the $\binom{m}{i}$ many points obtained by reducing the query to an $m-i$-dimensional point. The neighbourhood is then the union of the r-balls around these points.

Our intuition is that a given point can be somehow connected with "tunnels" to the points in other metric subspaces which are of distance 0, and then take advantage of the triangle inequality in the metric subspaces to use vantage indexing. However, the points in the database can possibly not include any such points of distance 0, so we have to modify this simplistic idea of a tunnel by instead linking to nearest neighbours. We precalculate, for every point set in all spaces except for that with the lowest dimensionality, links to its nearest neighbours in lower-dimensional spaces. If such links exist, one can efficiently retrieve not only the nearest neighbours in the same space as the query (e.g. as described in Section 3), but also the nearest neighbours in higher-dimensional and lower-dimensional spaces which are linked to the retrieved items from the space with the same dimensionality as the query. This yields almost the same result as an exhaustive linear search, but requires only logarithmic complexity.

One needs, however, to limit the number of connections per high-dimensional object by limiting the range of dimensionalities across which one wants to search. By introducing such a constant bound, one ensures that the number of connections grows only linearly with the number of items in the database. For the application of searching for rhythmic and melodic patterns, this means that one should limit the partial matching to a certain maximum number of missing or extra notes which may be ignored when the dissimilarity is calculated. Such a limit is probably desirable: without it, a short (low-dimensional) pattern could become very similar to very different long (high-dimensional) patterns.

4.1 Efficiently building tunnels

By projecting high-dimensional objects onto lower-dimensional subspaces and then using vantage indexing as described in Section 3 for finding the nearest neighbour of the projection, one can benefit from the logarithmic search complexity of vantage indexing for the purpose of building connections between nearest neighbours in metric subspaces of differing dimensionality.

Algorithm 1 can be used for building connections between objects of different dimensionality. Note that its runtime lies in $\mathrm{O}(n \log n)$ since we limit the number of dimensionalities to cross (the number of notes to ignore) to a constant t. The outermost loop is executed $\mathrm{O}(n)$ times. The

Algorithm 1 Build connections between nearest neighbours in subspaces of different dimensionality.

for all point sets p **do**
  for i := 1 to max. number of dimensions to cross **do**
    for all subspaces s with i dimensions less than p **do**
      project point set p onto subspace s
      use the vantage index of subspace s for finding p's nearest neighbours in s
      create a connection between p and its nearest neighbour in s (or all of them if they have equal distances).
    end for
  end for
end for

two inner loops each are executed only a constant number of times. Each point set p with j dimensions can be projected onto $(j-i)$-dimensional space in $\binom{j}{i}$ many ways, and this is bounded by a constant as long as the maximum possible number of dimensions and the number of subspaces to cross are bounded. Using the vantage index within the subspace onto which p is projected takes $\mathrm{O}(\log n)$ steps, which leads to an overall runtime of $\mathrm{O}(n \log n)$.

The space complexity can be expected to lie in $\mathrm{O}(n)$ since the number of connections to create should be bounded by $\binom{j}{i}$. There is one exception: if there are many nearest neighbours in the subspace which all have the same distance from the projected point, we might store more than $\binom{j}{i}$ connections for one projected point.

4.2 Using tunnels for retrieving items from higher dimensions than the query

To find rhythmic patterns which are similar to a given query, but contain up to t additional notes (which should be ignored when calculating dissimilarity), one can tunnel into higher dimensions by using the connections previously described. Given a query Q, search radius r, and a database with a vantage index for each metric subspace and connections between items and the nearest neighbours of their projections in subspaces with up to t fewer dimensions, one can retrieve Q's r-near neighbours with the same dimensionality as Q or up to t dimensions higher as follows:

Retrieve Q's nearest neighbours n within radius r with as many dimensions as Q, using the vantage index for this subspace.
for all r-near neighbours n **do**
  Retrieve every item which is connected to n and has a dimensionality higher than that of Q.
end for

This will retrieve exactly the desired objects from the subspace where the query Q resides, but for the higher-dimensional objects, both false positives and false negatives may occur.

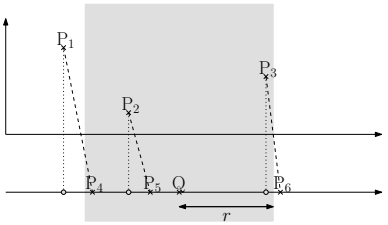

Figure 4. False negatives and false positives resulting from tunneling, and how to avoid them.

The database shown in Figure 4 contains 6 point sets $P_1, \ldots, P_6$. Three, $P_1, \ldots, P_3$, are two-dimensional, the others, one-dimensional. The query Q is one-dimensional. The area of interest within the search radius r around Q is marked grey.

False positives: It is conceivable that the projection of a higher-dimensional object onto Q's subspace lies just outside the search radius, but its nearest neighbour in Q's subspace happens to lie within the search radius. An example is P_1, whose projection onto the subspace (shown as a circle) has a nearest neighbour beyond the border of the grey area.

False negatives: It is also possible that while the projection of a higher-dimensional object onto Q's subspace lies inside the search radius, the closest object in Q's subspace lies outside the search radius. In this case, illustrated with P_3 and P_6, the higher-dimensional object will not be retrieved. In the extreme case that there is no single object inside the search radius in Q's subspace, no higher-dimensional objects whatsoever will be retrieved.

Controlling false negatives and false positives. To avoid all false negatives and limit the badness of false positives to a threshold e, one can add the projections as additional "ghost points" to the database if their nearest neighbour in the subspace is further away than $e/2$, and extend the search radius by $e/2$.

The distance of false positives to the query will be at most e higher than desired because in the worst case, the nearest neighbour of the projection will lie on the line connecting the projection with the query. The nearest neighbour can be up to $r + e/2$ away from the query, while the projection can be another $e/2$ away from the nearest neighbour, leading to a total maximum distance of $r + e$ for false positives.

Such additional ghost points would be added as part of the task of building the index, and so would not slow down the search process. It would also not increase the computational complexity of building the index – the only price is some extra space for storing ghost points wherever some point from higher dimensions gets projected into a sparsely populated area in a subspace. There is a tradeoff between the required additional space and the maximum possible distance error for false positives.

4.3 Using tunnels for retrieving items from lower dimensions than the query

As we have seen in Algorithm 1, subspaces can be searched efficiently for a query even without using the pre-computed connections. Since these connections are already there, they can be used instead of projecting the query onto each possible subspace. To avoid false negatives, one still needs to search the area around the nearest neighbour in the subspace with the search radius $r + a$, using the vantage index of the subspace, where a is the distance between the projection of Q onto the subspace and its nearest neighbour in that subspace. Whenever a is greater than zero, the possibility of false positives gets introduced. If such false positives cannot be tolerated, one can resort to not using the connections but instead projecting the query onto all possible subspaces.

5 EXPERIMENTAL EVALUATION OF VANTAGE INDEXING WITH TUNNELING

For an experimental evaluation, we used Dixon's onset detector from his "BeatRoot" system [4] for extracting onsets from various recordings with piano music (piano concertos and solo works by Beethoven, Liszt, and Schumann). We used piano music because piano onsets are relatively easy to detect correctly. The detected onsets were grouped into overlapping sequences of 5 to 8 onsets. $40,000$ of these sequences were put into a database and indexed for optimum retrieval within subspaces as described in Section 3. We also added connections between subspaces as described in Section 4, with connections spanning up to 3 dimensionalities, thus allowing for ignoring up to 3 extra or missing notes.

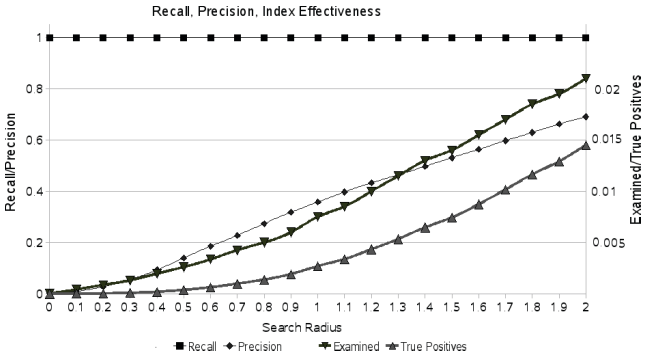

Figure 5. Recall ■, precision ♦ (left scale), ratio of retrieved items ▼, and ratio of true positives ▲ (right scale) for the tunneling method and different search radii, averaged over 20 queries.

For every search radius in a range from 0 to 2, we randomly picked 20 items from the database as queries and used the index to retrieve the most similar items according to the EMD. To calculate precision and recall, we also did exhaustive searches for each of these queries (by calculating the EMD between the query and each of the $40,000$ items in the database) and counted false positives and false negatives.

With this experiment, we verified that it is indeed possible to reduce the number of false negatives to zero and limit the distance error of false positives to e by adding the projections of high-dimensional points to the database in cases where their nearest lower-dimensional neighbour is further away than $e/2$, and by increasing the search radius by $e/2$. Figure 5 shows that our index works well: for example, with a search radius of 1.9, we need to retrieve 1.3% of the whole database (curve: ▲), but we actually retrieve 1.95% (▼ Examined $= 0.0195$), which still relieves us from looking at 98.05% of the database. For this experiment, the value of e was 1, i.e., we inserted ghost points only if the projection and its nearest neighbour were more than 0.5 apart. With this threshold, we needed about as many ghost points as real points. The nested range tree we used for the vantage table [1] has a space complexity of $O(n \log n)$.

The larger our search radius, and thus the more important it is that we do not get swamped with false positives, the lower their percentage in the search results. This is probably due to the fact that false positives can only occur near the surfaces of searched balls, and when the search radius is increased, one expects an ever larger percentage of objects inside the ball rather than near its surface. Although the precision is not very good for very low search radii, this does not matter much since for low search radii, very few items are retrieved at all from the database. As the number of retrieved database items grows with the search radius, the pain resulting from false positives is relieved by the fact that at the same time, the percentage of false positives among the retrieved items shrinks.

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AUTOMATIC MAPPING OF SCANNED SHEET MUSIC TO AUDIO RECORDINGS

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ABSTRACT

Significant digitization efforts have resulted in large multimodal music collections comprising visual (scanned sheet music) as well as acoustic material (audio recordings). In this paper, we present a novel procedure for mapping scanned pages of sheet music to a given collection of audio recordings by identifying musically corresponding audio clips. To this end, both the scanned images as well as the audio recordings are first transformed into a common feature representation using optical music recognition (OMR) and methods from digital signal processing, respectively. Based on this common representation, a direct comparison of the two different types of data is facilitated. This allows for a search of scan-based queries in the audio collection. We report on systematic experiments conducted on the corpus of Beethoven's piano sonatas showing that our mapping procedure works with high precision across the two types of music data in the case that there are no severe OMR errors. The proposed mapping procedure is relevant in a real-world application scenario at the Bavarian State Library for automatically identifying and annotating scanned sheet music by means of already available annotated audio material.

1 INTRODUCTION

The last years have seen increasing efforts in building up large digital music collections. These collections typically contain various types of data ranging from audio data such as CD recordings to image data such as scanned sheet music, thus concerning both the auditorial and the visual modalities. In view of multimodal searching, navigation, and browsing applications across the various types of data, one requires powerful tools that support the process of analyzing, correlating, and annotating the available material. In the case of digitized audio recordings, first services have been established to automate the annotation process by identifying each recording and assigning available metadata such as title, artist, or lyrics. Here, the metadata is drawn from specialized annotation databases provided by commercial services such as Gracenote [6] or DE-PARCON [9].

Opposed to acoustic music data, which is increasingly available in digital formats, most sheet music is still produced and sold in printed form. In the last years, digital music libraries have started to systematically digitize their holdings of sheet music resulting in a large number of scanned raster images. To make the raw image data available to content-based retrieval and browsing, methods for automatically extracting and annotating semantically meaningful entities contained in the scanned documents are needed. In this context, *optical music recognition* (OMR) [3] is a key task. Here, the goal is to convert scanned sheet music into a computer readable symbolic music format such as MIDI or MusicXML [13]. Even though significant progress has been made in the last years, current OMR algorithms are substantially error-prone, resulting in systematic errors that require subsequent correction [2]. Similarly, there is still a high demand for reliable solutions for the more general task of automatic sheet music annotation in the digital library community.

In this paper, we present a novel approach for automatically annotating scanned pages of sheet music with metadata. Our approach is based on a new procedure for *mapping* the scanned sheet music pages to an existing collection of annotated audio recordings. The mapping allows for identifying and subsequently annotating the scans based on the metadata and annotations that are already available for the audio recordings. In particular, as it is the case in the specific application scenario at the Bavarian State Library, we assume the existence of an audio collection containing annotated digitized audio recordings for all pieces to be considered in the sheet music digitization process. The conversion of both the audio recordings (by employing filtering methods) and the scanned images (by employing OMR) to a common feature representation allows for a direct comparison of the two different types of data. Using the feature sequence obtained from a few consecutive staves or an entire page of the scanned sheet music as query, we compute the top match within the documents of the audio database. The top match typically lies within a musically corresponding audio recording, which then allows for identifying the scanned page and for transferring all available audio anno-

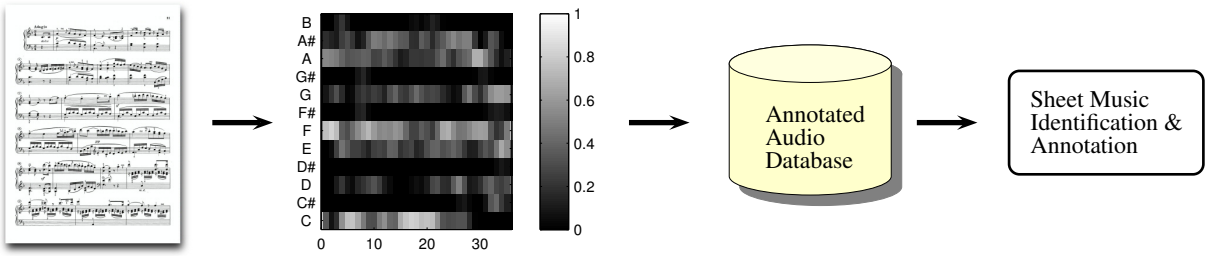

Figure 1. Overview of the mapping procedure for automatic identification and annotation of scanned sheet music using an annotated audio database. The first page of the second movement of Beethoven's piano sonata Op. 2 No. 1 and the resulting scan chromagram are shown.

tations to the scanned sheet music domain. This procedure is described in Sect. 2 and illustrated by Fig. 1. We have tested and analyzed our mapping procedure by means of a real-world application scenario using the corpus of the 32 piano sonatas by Ludwig van Beethoven. In Sect. 3, we discuss the outcome of our experiments showing that the mapping across the two music domains is robust even in the presence of local OMR errors, but suffers in the presence of severe global OMR errors. We also describe a postprocessing procedure that allows for detecting most of the misclassifications and automatically reveals most of the passages within the scanned pages where the severe OMR errors occurred. In Sect. 4, we conclude this paper with prospects on future work and indicate how to improve the identification rate by correcting and compensating for severe OMR errors prior to the mapping stage.

2 MAPPING PROCEDURE

One key strategy of our mapping procedure is to reduce the two different types of music data, the audio recordings as well as the scanned sheet music, to the same type of feature representation, which then allows for a *direct* comparison *across* the two domains. In this context, chroma-based features have turned out to be a powerful mid-level music representation [1, 7, 12]. Here, the *chroma* correspond to the twelve traditional pitch classes of the equal-tempered scale and are commonly indicated by the twelve pitch spelling attributes C, C$^\sharp$, D, . . .,B as used in Western music notation. In the case of audio recordings, normalized chroma-based features indicate the short-time energy distribution among the twelve chroma and closely correlate to the harmonic progression of the underlying piece. Based on signal processing techniques, the transformation of an audio recording into a chroma representation (or chromagram) may be performed either by using short-time Fourier transforms in combination with binning strategies [1] or by employing suitable multirate filter banks [12]. In our implementation, we use a quantized and smoothed version of chroma features, referred to as CENS features, see [12]. The transformation of scanned sheet music into a corresponding chromagram requires several steps, see [11]. First, each scanned page is analyzed using optical music recognition (OMR) [2, 3]. In our system, we use the commercially available SharpEye software [8] to extract musical note parameters (onset times, pitches, durations) along with 2D position parameters as well as bar line information from the scanned image. Assuming a fixed tempo of 100 BPM, the explicit pitch and timing information can be used to derive a chromagram essentially by identifying pitches that belong to the same chroma class. A similar approach has been proposed in [7] for transforming MIDI data into a chroma representation. Fig. 1 shows a resulting scan chromagram obtained for the first page of the second movement of Beethoven's piano sonata Op. 2 No. 1. Note that the tempo assumption (of always choosing 100 BPM) is not a severe restriction since the mapping algorithm to be described next can handle local and global tempo variations anyway.

For the actual scan-audio mapping, we use a matching procedure similar to the one described in [10]. First, in a preprocessing step, all recordings of the given audio database are converted into sequences of CENS vectors. (In our implementation, we use a feature sampling rate of 1 Hz.) While keeping book on document boundaries, all these CENS sequences are concatenated into a single audio feature sequence. Then, each scanned page of sheet music to be identified is also converted into a sequence of CENS features. The resulting scan feature sequence is then compared to the audio feature sequence using subsequence dynamic time warping (DTW). For a detailed account on this variant of DTW we refer to [12]. In our experiments, it turned out that the DTW step sizes $(2, 1), (1, 2), (1, 1)$ (instead of the classical step sizes $(1, 0), (0, 1), (1, 1)$) lead to more robust matching results, and are hence used in the remainder of this paper. As a result of the DTW computation, one obtains a *matching curve*. The ith position of the matching curve contains the costs for matching the scan feature sequence to the most similar subsequence of the audio feature sequence ending at position i. Therefore, the curve's local minima close to zero correspond to audio feature subsequences similar to the scan feature sequence. These subsequences are referred to as *matches*. Because of the book-keeping, doc-

ument numbers and positions of matches within each audio document can be recovered easily. Note that DTW can compensate for possible temporal differences between scan feature sequences and corresponding audio feature subsequences thus also relativizing the above tempo assumption.

3 EXPERIMENTS AND EVALUATION

The basic identification and annotation procedure of a given scanned page of sheet music can be summarized as follows. First, map a given scanned page against the audio database and derive the top match of lowest cost. Then determine the audio recording that contains the top match and transfer all annotations available for the audio recording to the image domain. Note that in the ideal case the top match not only identifies the scanned page but also indicates the time position within the audio recording where the music notated on the page is actually played.

We now show to which extent this approach also works in practice by discussing a real-world application scenario using the musically relevant corpus of Beethoven's piano sonatas. Our test database and some technical details are described in Sect. 3.1. In Sect. 3.2, we discuss a baseline experiment using MIDI versions instead of extracted OMR data. This experiment indicates which identification rates one may expect in the optimal (but unrealistic) case where no OMR extraction errors have occurred. Then, in Sect. 3.3, we describe several experiments performed on the actual OMR data. Here, we also discuss various types of OMR errors that significantly degrade the mapping quality. Finally, in Sect. 3.4, we describe a postprocessing strategy that automatically reveals most of the misclassifications.

3.1 Test Database

Our experiments have been conducted on the basis of the 32 piano sonatas by Ludwig van Beethoven, which play a key role in the evolution of the sonata form and are considered as one of the greatest treasures in the music literature. Because of its outstanding musical significance and the large number of available digitized audio recordings, the automated analysis and organization of the corpus of Beethoven's piano sonatas is highly relevant to musicologists and librarians.

Our audio database consists of a complete recording of the 32 piano sonatas conducted by Daniel Barenboim, comprising 101 audio documents (basically corresponding to the movements) and 11 hours of audio material. Furthermore, we have a scanned version of the corresponding sheet music (Volume 1 & 2, G. Henle Verlag) at our disposal amounting to a total number of 604 digitized pages (3693 two-stave systems). In the following, dealing with piano music, the term *line* is used to denote a two-stave system consisting of a stave for the right and a stave for the left hand. The scanned pages, which are available as 600dpi b/w images in the TIFF format, have been processed by the OMR Engine of SharpEye 2.68 and saved in the MusicXML file format as one file per page. Finally, each of the 604 MusicXML files was transformed into a sequence of CENS vectors (one feature per second) assuming a fixed tempo of 100 BPM. Subsequently, using the extracted OMR information on the notated systems, the CENS sequences were segmented into 3693 subsequences corresponding to the lines.

3.2 Baseline Experiment: MIDI-Audio Mapping

In a baseline experiment, we investigated what identification rates one may expect in the case that there are no severe OMR extraction errors. To this end, we used a complete set of MIDI files for the 32 Beethoven sonatas and randomly generated a large number of MIDI fragments of various lengths, which were used instead of the OMR extraction results. Then, for each of these MIDI fragments we computed a matching curve with respect to the audio database and determined the topmost audio match. Recall that in the identification scenario the objective is to determine the piece of music underlying the respective MIDI fragment by using the audio recordings of the database as an identifier. Therefore, we consider a match as *correct* if it lies within the audio document that corresponds to the same movement as the MIDI document from which the respective query is taken. Otherwise the match is considered as *incorrect*.

In particular, we investigated the dependence of the number of correct audio matches subject to the length L (given in seconds) of the MIDI query. To this end, we randomly generated 1000 MIDI queries for each of the seven parameters $L \in \{10, 20, 30, \ldots, 70\}$. Each of the queries lies within a single MIDI file and therefore has a unique correct assignment to one of the 101 movements. The second column of Table 1 shows the number of correct matches. As an example, consider the case $L = 10$, where 823 of the 1000 matches were correct. Note that the number of correct matches increases significantly with the query length. For example, for $L = 40$ only 3 of the 1000 queries were misclassified. To give a more detailed picture of the matching quality, Table 1 additionally provides various cost and confidence values. The third, fourth, and fifth column show the average cost values, the standard deviations, and the maximal cost values for the correct top matches. For example, in the case $L = 10$, the average cost value (standard deviation/maximal cost value) for the 823 correct matches is 0.059 (0.024/0.223). The latter cost values are with respect to a range from 0 (no costs) to 1 (maximum costs). Increasing L leads to slightly higher cost values stabilizing around the value 0.07 even for long queries.

Similarly, the sixth, seventh, and eighth columns of Table 1 show the corresponding values for the incorrect top matches. For example, in the case $L = 10$, the average cost of the 177 incorrect top matches is 0.084 with a standard de-

Length	#(Cor.)	Cost (correct)			Cost (incorrect)			Gap
(in sec.)	(in %)	av.	std.	max.	av.	std.	max.	av.
10	82.3	0.059	0.024	0.223	0.084	0.034	0.207	0.044
20	96.7	0.068	0.026	0.206	0.102	0.031	0.196	0.070
30	99.2	0.070	0.024	0.189	0.139	0.040	0.214	0.093
40	99.7	0.071	0.024	0.218	0.177	0.027	0.198	0.106
50	99.9	0.072	0.023	0.204	0.117	0.000	0.117	0.118
60	99.9	0.071	0.021	0.193	0.159	0.000	0.159	0.128
70	99.9	0.071	0.022	0.196	0.229	0.000	0.229	0.135

Table 1. Results for the baseline experiment of mapping MIDI fragments of various lengths $L \in \{10, 20, \ldots, 70\}$ (given in seconds) to the audio database. Each line shows the length L, the percentage of correct matches for the 1000 MIDI fragments of the respective length, the average values (av.), the standard deviations (std.), and the maximum values (max.) of the correct matches and incorrect matches, and the average confidence gap.

Length	#(Cor.)	Cost (correct)			Cost (incorrect)			Gap
(in sec.)	(in %)	av.	std.	max.	av.	std.	max.	av.
10	48.4	0.080	0.033	0.198	0.104	0.040	0.247	0.034
20	67.9	0.103	0.039	0.261	0.147	0.051	0.285	0.050
30	78.4	0.114	0.044	0.292	0.173	0.049	0.317	0.062
40	84.9	0.120	0.043	0.356	0.192	0.051	0.340	0.072
50	87.1	0.132	0.043	0.305	0.208	0.051	0.367	0.080
60	87.0	0.143	0.050	0.304	0.232	0.044	0.356	0.080
70	87.1	0.153	0.052	0.316	0.247	0.049	0.373	0.078

Table 2. Experimental results mapping OMR fragments of various lengths (given in seconds) to the audio database. For each length parameter $L \in \{10, 20, \ldots, 70\}$ we randomly generated 1000 OMR chroma subsequences, each corresponding to a subpart of exactly one of the scanned pages. The table has the same interpretation as Table 1.

viation of 0.034. Note that in the case of incorrect matches, when increasing the query length, the average cost increases at a much higher rate than in the case of correct matches.

We also investigated how well the correct matches were separated by successive matches that do not lie in the respective correct audio document. To this end, we computed for each query the minimal cost value of a restricted matching curve, where the correct audio document had been removed. Then, for all correctly identified queries, we computed the difference of this minimal value and the cost of the correct top match. This difference value, which we refer to as *confidence gap*, indicates the identification reliability based on the top match. The average confidence value is shown in the last column of Table 1. For example, in the case $L = 10$ the average confidence gap amounts to the value 0.044. Increasing L leads to a significant increase of the confidence gap up to the value of 0.135 for $L = 70$. In conclusion, one may say that one obtains very good identification rates (with an error rate of less than 1%) for MIDI fragments of at least 30 seconds of duration.

3.3 OMR-Audio Mapping

Next, we describe a similar experiment, now using the (potentially flawed) OMR extraction results instead of the "clean" MIDI data. For each of the 604 scanned pages, we computed a CENS feature sequence as explained in Sect. 2. Then, from these sequences, we randomly generated 1000 subsequences of length L for each of the length parameters $L \in \{10, 20, \ldots, 70\}$. Table 2 summarizes the OMR-audio mapping results. Obviously, the identification rate drops significantly compared to the pure MIDI case. For example, in the case $L = 10$ only 484 out of the 1000 OMR query fragments appear as top match in the correct audio document (opposed to the 823 correct matches in the MIDI case). The identification rate increases to roughly 87% for OMR feature sequences that correspond to a duration of 50 seconds and above. A comparison with Table 1 shows that, in the OMR case, the average costs of the correct matches are much higher than the ones in the MIDI case. Furthermore, the confidence gap is much smaller.

All these numbers indicate that the OMR-audio mapping procedure significantly suffers from the artifacts that are mainly caused by OMR extraction errors. In particular, a manual investigation of samples of the OMR extraction results revealed that there are two prominent types of OMR errors that significantly degrade the quality of the CENS feature sequences. First, for roughly 7% of the lines (two-stave systems) the key signature was extracted incorrectly. In particular, one or even more accidentals notated at the beginning of each stave were missing. Such an error generally distorts the CENS subsequence for an entire line, since a missing accidental causes all notes of a specific pitch class to be shifted upwards or downwards by one semitone, which may significantly corrupt the chroma distribution. Second, in almost 5% of the measures there were some note or beam extraction errors that resulted in inconsistencies with respect to the notated time signature. In such cases, the conversion tool of our OMR software, which transforms the OMR extraction parameters into a MusicXML file, simply discards all voices within those measures that reveal such inconsistencies. This also results in a significant corruption of the chroma distribution. Obviously, the automated detection and correction of such OMR extraction errors would overcome these problems resulting in significantly improved identification rates. These issues are left for future work and are further discussed in Sect. 4.

We continue the analysis of our OMR-audio mapping procedure based on the raw OMR material. Instead of using randomly chosen OMR fragments of a specific duration, we now investigate the mapping quality based on musical units such as pages or lines. Using entire pages in the OMR-audio mapping leads to an identification rate of roughly 82.5%. The average length of the corresponding CENS sequences

Lines	Length (in sec.)	$k=1$ (in %)	$k=2$ (in %)	$k=5$ (in %)	$k=10$ (in %)	$k=20$ (in %)	$k=50$ (in %)
1	9.133	44.57	52.97	65.77	76.20	84.59	92.26
3	27.099	71.30	76.66	83.62	88.06	92.45	96.13
5	45.053	77.04	81.23	86.41	90.12	93.37	96.86
7	62.995	77.74	81.83	86.84	90.85	93.83	96.89

Table 3. Identification rates depending on the number of lines used in the OMR-audio mapping. The columns indicate the recall percentage (out of 3693 mappings, respectively) of the correct audio document within the top k matches.

amounts to 55 seconds yielding robust mappings if there are no severe OMR errors. Another problem that often leads to misclassifications is that a single scanned page may refer to more than one pieces of music. In particular for our Beethoven corpus, a single page may contain both the end and the beginning of two consecutive movements. To overcome this problem, one may use single lines in the mapping process instead of entire pages. This also yields the advantage of having several identifiers per page. On the downside, the average length of the CENS sequences corresponding to the lines lies below a duration of 10 seconds yielding an identification rate of only 44.57%, see Table 3. To improve the identification rate of the line-based mapping strategy, we query each line in the context of ℓ preceding and ℓ subsequent lines. In other words, instead of using a single line we use a block of $2\ell+1$ subsequent lines with the reference line positioned in the middle. Here, we assume that all pages belonging to one movement are in the correct order, hence allowing us to consider blocks of lines ranging across two consecutive pages. To systematically investigate the identification rate depending on the number of lines used in the OMR-audio mapping, for each of the 3693 lines of our scanned Beethoven material, we generated CENS query sequences corresponding to 1, 3, 5, and 7 lines. Table 3 shows both the resulting identification rates based on the top match ($k=1$) and the recall values for the correct audio document for the top k matches with $k \in \{1,2,5,10,20,50\}$. For example, using three lines, the top match ($k=1$) was correct in 71.30% of the 3693 OMR-audio mappings. Considering the top 5 matches ($k=5$), at least one of these matches was correct in 83.62% of the mappings.

3.4 Postprocessing

We now show how the additional information of considering the k top matches (instead of considering only the top match) can be used to detect most of the incorrect identifications. The only assumption we use is that the scanned pages that correspond to a specific movement are given as a sequence of consecutive pages, i. e., pages of different movements are not interleaved. We explain our postprocessing procedure by means of our Beethoven scenario using $Q = 3693$ OMR queries each consisting of 7 subsequent lines and considering the $k=5$ top matches. Recall that the objective is to map each of the queries to one of the $P=101$ audio documents (representing the pieces or movements). We construct a $P \times Q$ *mapping matrix* M, where the rows correspond to the pieces and the columns to the queries. Then an entry $M(p,q)$, $1 \leq p \leq P$, $1 \leq q \leq Q$, is non-zero if and only if the p^{th} audio document appears among the top k matches for the q^{th} query. In this case $M(p,q)$ is set to $1-c$, where $c \in [0,1]$ denotes the cost of the corresponding match. In case there are several matches for the entry (p,q) among the top k matches, we define c to be the minimal cost value over these matches. Note that $M(p,q)$ expresses a kind of confidence that the q^{th} query belongs the p^{th} piece. Furthermore, M indicates the kind of confusion that occurred in the identification procedure. Fig. 2 shows the mapping matrix for the Beethoven scenario.

For our Beethoven corpus, both the audio recordings and the scanned pages are sorted with respect to increasing opus and movement numbers. Therefore, a correct mapping of all queries corresponds to a diagonal staircase-like structure in M. In the following, we do not assume that the scanned pages are given in the same order (on the piece and movement level) as the audio recordings, since this assumption is often violated in real-world digitization applications. For example, many music books contain a more or less unsorted mixture of various pieces and movements. Therefore, we only make the assumption that the pages that correspond to a specific audio document (referring to a specific movement) are given in the correct order. Then, in case of a correct identification of the OMR queries, the matrix M reveals a structure of horizontal line segments, where each such segment corresponds to an audio document.

In the following, a tuple (p,q) is referred to as *positive* if the entry $M(p,q)$ is non-zero. Furthermore, a positive tuple (p,q) is referred to as *true positive* if the q^{th} query semantically corresponds to the p^{th} audio document, otherwise (p,q) is called *false positive*. Now, the idea is that positive tuples included in long horizontal line segments within M are likely to be true, whereas isolated positive tuples are likely to be false. Intuitively, our procedure classifies the positive tuples by looking for groups of tuples included in long horizontal line segments (these tuples are classified as true) and discards isolated positives tuples (these tuples are classified as false). Due to space limitations, we do not give technical details and refer to Fig. 2 for an illustration.

We have applied this postprocessing procedure to the Beethoven scenario using $Q = 3693$ queries each consisting of 7 subsequent lines and considering the top match only. As a result, 78.42% of the queries were mapped correctly and 17.17% of the queries were not mapped (by discarding false positives). The remaining 4.41% are incorrect mappings. Note that the result of this type of postprocessing is the detection rather than the correction of incorrect identifi-

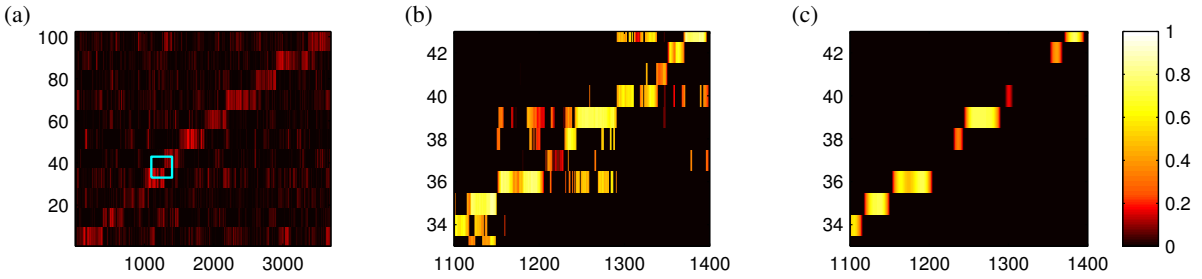

Figure 2. **(a)** Mapping matrix M for the Beethoven scenario. The rows correspond to the audio documents ($P = 101$) and the columns to the OMR queries ($Q = 3693$). **(b)** Enlargement of the marked region of M. **(c)** The same region after applying the postprocessing procedure.

cations. Having identified incorrect mappings allows to both further improve the identification process and to automatically reveal passages within the sheet music where severe OMR errors have occurred.

Rather than identifying incorrect mappings, one may also increase the number of correct identifications. For this, certain tuples are specified as true positives by "filling" small gaps within horizontal line segments. Thus, OMR queries are assigned to a specific audio document if neighboring OMR queries are consistently assigned to the same audio document. Using $k = 3$ in our example increases the number of correct identifications to 86.70% (instead of 77.74% without postprocessing). Note that there is a natural trade-off between eliminating the incorrect identifications and boosting the correct identifications.

4 CONCLUSIONS

In this paper, we have introduced the problem of mapping sheet music to audio recordings. Based on an automated mapping procedure, we have presented a novel approach for automatically identifying scanned pages of sheet music by means of a given audio collection. Such a procedure, which constitutes an important component in the digitization and annotation process of multimodal music material, is needed for building up the Probado music repository [4] currently set up at Bavarian State Library in Munich, Germany. This music repository, which contains digitized sheet music and audio data for a large collection of classical and romantic piano sonatas (Haydn, Mozart, Beethoven, Schubert, Schumann, Chopin, Liszt, Brahms) as well as German 19th centuries piano songs, is continuously expanded requiring automated procedures for music processing and annotation.

As our experiments show, the proposed procedure for mapping scanned sheet music and audio material works well in the case that there are no severe OMR extraction errors. Our postprocessing procedure allows for automatically revealing most of the critical passages containing these OMR errors. In the future, we will use various heuristics to correct typical OMR errors prior to the mapping step. For example, in the case of piano music, different key signatures for the left and right hand staves can be assumed to be invalid and easily corrected by considering neighboring stave lines. Furthermore, similar to the strategy suggested in [2], one can simultaneously employ various OMR extraction results obtained from different OMR software packages to stabilize the mapping result. Based on these strategies, we expect to achieve a significant improvement of the identification rates reaching the ones reported in our MIDI baseline experiment.

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DEVELOPMENT OF A MUSIC ORGANIZER FOR CHILDREN

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ABSTRACT

Software development for children is challenging; children have their own needs, which often are not met by 'grown up' software. We focus on software for playing songs and managing a music collection—tasks that children take great interest in, but for which they have few or inappropriate tools. We address this situation with the design of a new music management system, created with children as design partners: the *Kids Music Box*.

1. INTRODUCTION

Digital documents such as music, movies, and pictures have become widely available; it is not uncommon for an individual's hard drive to contain hundreds of songs and videos, and thousands of photos. However, both the interface and usability of digital document organizers are aimed for adults, and far less attention has been focused on the development of applications that are suitable for primary school children (six to ten years old). Children are as information thirsty as adults; they want to access their favourite songs and movies as much as adults do, and prize the ability to independently select and enjoy their entertainment 'just like a big person'. But at present children are forced to use computer software that requires complex interaction, good spelling, and reading skills beyond their current abilities.

A number of music-related systems aimed at children are described in the research literature. They primarily focus on supporting children in creating music (eg, [4]), often in the context of formal music education (eg, [6]). MIR research attention has not yet turned to supporting children in playing, browsing, searching, and organizing collections of *existing* music. This paper describes the design and prototyping of such an organizer, the *Kids Music Box* (KMB). The development of KMB was grounded in previous research on creating software for children (Section 2), and follows the Participatory Design [10] model for software development (Section 3). Structured techniques for matching functionality and interface/interaction design to the target users were also employed (eg, expert review and usability testing). The major contribution of this paper is to demonstrate this principled approach to developing music-interaction software for children.

2. DESIGNING SOFTWARE FOR CHILDREN

Children are not miniature adults [4]; they have different needs, capabilities and expectations of computer technologies. Making cosmetic changes to adult software does not adequately adapt it for children [3]. To guide development of KMB, we first reviewed existing literature regarding software design for children.

2.1. Cognitive Development

Chiasson et al. [3] found that most adult user-interfaces assume that users are proficient readers with fairly extensive vocabularies; most children, however, have not reached this level of proficiency. Older children may not fully understand text-based instructions, while young children may not even know the alphabet. Children are creative spellers and it is hard for an interface to recognise their text input [6]. Given these difficulties in interpreting on-screen text, it is particularly important that icons be meaningful and intuitive. Where possible, the icons should represent familiar, real-world objects [5].

Children expect to see the results of their actions. If nothing happens after an action has been performed, it is very likely that children will repeat their action until there is some response from the application. Although constant visual or audio feedback might be annoying for adult users, children often expect and enjoy it [9].

Young children have difficulty with abstract concepts, and may not be able to navigating complex interfaces. Their most common method for learning to use new software is trail-and-error. Once they have found a method that works, it is very likely that they will keep using it, instead of searching for a more efficient method.

To summarise:

- Interfaces should be strongly visual, avoiding text where possible [4]
- Content specific metaphors are useful in helping children navigate interfaces [4].
- Where possible, allow children to select input from a list rather than directly enter text [6].
- Children are impatient and want instant feedback on their actions [9].
- Icons should be visually meaningful [5].

- Interfaces should take into account the fact that children may not yet understand abstract concepts [4].
- Interfaces should not make use of menus and sub-menus as children may not have the ability to categorize so as to navigate efficiently [4].
- Rollover audio, animation, and highlighting should be used to indicate where to find functionality [4].

2.2. Physical Development

Since children's motor control skills develop over time, until fully developed, it is difficult to perform tasks such as controlling the mouse and targeting small areas on the screen. For example, tasks requiring them to hold down mouse for extended period are tiring and difficult [5]. Typing on a standard keyboard is also an issue for children as their strategy is "hunt-and-peck".

Children find it challenging to accurately drag an item on screen [11]. They may also lack the fine-motor control needed to target small items on screen. It is important to have icons big enough for children to identify and at the same time, icons should be spaced to minimise the chance that children accidentally press the wrong button [4].

To summarise:

- Mousing should be as simple as possible [4].
- Screen items should be big enough and distanced from each other to compensate inaccuracy in targeting [4].
- Dragging is difficult for children [5].

3. KMB DESIGN AND DEVELOPMENT

We base our work on the *Participatory Design model* , the essence of which is to respect users as partners in the design process, by giving them a significant role in the design phase [10]. However, it is unreasonable to expect that that children can have the same level of explicit input as an adult. Since children are usually unable to clearly verbally articulate their feelings and reactions to software designs, the onus is on designers to identify these reactions from the child partners' emotional displays and physical actions during participatory design exercises [5]. To that end, we videotaped design exercises.

That said, it is also central to the Participatory Design model to empower the users to negotiate team decisions with the designers. This can be a significant issue when working with children, as the children must learn to trust that adults will listen to their contributions, and adults must learn to elaborate on children's ideas rather than dismissing them [1].

3.1. Evaluation of suitability of existing organizers

Our initial step was to evaluate existing digital music organizers for their suitability with our target age group (6 to 10 years old). We identified one organizer intended specifically for children (*KidsPlayer*, developed by SororSoft); for contrast, we also examined the suitability of two organizers intended for adults (*Windows Media Player v10* and *iTunes v7.01*).

These three organizers were investigated in a focus group study involving eight participants aged between six and ten years old. Three of the participants were female and five were male; the average age was 7.8 years. All participants had regularly used a computer. For the study, the participants used all three organizers to: import songs; manage a playlist by deleting songs, assigning genre and ratings, and so forth; loading a saved playlist; and selecting and playing one or more songs. These tasks include the minimum requirements of a digital music organizer. The study included a debriefing, and 'homework' in the form of an opportunity to draw a design for an ideal music organizer.

This evaluation study uncovered significant usability problems in all three organizers. The most significant barriers were in importing music and creating a new playlist, primarily because these require the user to understand the Windows file system. All participants required assistance from the researcher to locate specific folders for songs—a difficulty corroborated by the children in the debriefing. When asked the question, "*Do you look after the songs on your computer yourself?*", all participants responded that their parents or other family members help them look after both their physical (CDs and tapes) and digital music.

The children liked KidsPlayer's big and colorful buttons, corroborating suggestions in the literature review that children prefer vibrant colors and large controls [6]; the iTunes interface was dismissed as '*boring*' with its small buttons and grayscale windows. However, KidsPlayer uses a 'bubble wrapping' effect on buttons that causes them to look like part of the background graphics, and the children had difficulties in identifying its clickable objects. An attractive interface must still adhere to basic usability principles—and in particular, the children had difficulty with small fonts, small buttons and 'handles', right clicks, drag-and-drop, and double-clicking.

A major difficulty with the products designed for adults (*Media Player* and *iTunes*) is that they provide too much information and functionality for children. Children simply do not know what to focus on when too many details are presented to them—and so, for example, the presentation of many columns of metadata turned the task of locating a single song by its title into a frustrating chore. The children also responded positively to the 'clear and simple' layout of *Media Player*, which more cleanly groups related features than do the other two organizers.

The children were particularly enthusiastic about two functionalities: the ability to rate songs in *iTunes* ("*So I can give five stars to my favourite songs!*"), and the *Media Player* visualization that synchronizes with the music ("*I*

like to see those colourful and moving graphics when I play my song").

3.2. Prototyping and Expert Review

Based on the background research into effective interface/interaction designs for children (Section 2) and the evaluation of existing music organizers (Section 3.1), a list of essential design requirements emerged: clear navigation paths; access to visualizations and games while listening to music; features grouped by functionality, and clear guidance through multi-step processes; use of point-and-click rather than double-clicking, right-clicking, or drag-and-drop; and a colorful design with large clickable areas. These principles guided the creation of a C# based horizontal prototype (ie, a largely non-functional demonstration of the design) [7].

With adult users, at this point a 'walkthrough' of Use Cases is staged with a group of potential users to refine the interface and interaction design. Children, however, find it overwhelmingly difficult to provide feedback from a non-functional prototype, particularly when the most engaging design aspects (eg, playing songs, viewing visualizations) are not supported. For this reason the initial design was instead the subject of an expert evaluation [7] by two design experts with significant experience in creating interfaces for children. The expert review identified many minor improvements to the interface design (for example, in the choice of colors and shape of the frames distinguishing spaces within a window). The most significant changes were to further simplify the interface and clarify the interaction flow, and to support a variety of 'skins' for KMB (discussed in Section 3.4).

3.3. Design Overview and Implementation

Kids Music Box was developed with *C#, MySQL* and *Macromedia Flash*. *C#* was selected for its support of rapid development. *MySQL* database was used to store all the account and song information and to provide responses to user queries. *Macromedia Flash* was used to construct all the interactive features, such as the buttons and the 'playground' of games and visualizations.

The 'box' was chosen as the primary visual interface metaphor for KMB. The real world analogy is to children and their toys – children are taught early that toys cannot be scattered all over the house; they needed to be sorted in a box or a drawer, so the toys can be retrieved when needed and the house is not cluttered. Digital organizers in many aspects are the counterpart to storage boxes and drawers that hold children's toys.

KMB has a multi-coloured and non-rectangular interface; children from the focus group study have agreed that colours and shapes were more interesting. The main display area has multiple functionalities; it is able to display both pictures and text, display visualisations and it is also an interactive playground. The main reason why MediaPlayer's visualisation was so well received in the initial user study (Section 3.1) is that it offered children "something to do" while listening to music [7]; children are not interested in sitting quietly and listening to songs. One real-world example occurs in music classes for kids, which often involve several activities (dancing, clapping, games) while singing or listening.

Children enjoy making and receiving presents: every parent's refrigerator is covered with art given proudly by their child, school art classes help children create cards and gifts for every occasion, and children (particularly girls) exchange handmade tokens of friendship. Music–related gifts hold a special position for adult, who may give gifts of songs and CDs or create elaborate and emotionally meaningful playlists or compilation CDs [4]. To support this behavior, users can create a 'gift' music box that can be opened by another KMB.

Participants in the study of existing organizers (Section 3.1) liked the idea of organizing their songs as playlists or groups of 'favorites', but found it difficult to manipulate files. KMB takes an approach to managing playlists that is similar to that of iTunes (the playlist manager that the children found to be most usable): users do not have to know where the playlists are stored, since both saving and loading is handled by the organizer.

The final KMB interface relies heavily on images—each button or functional area has an associated icon. Children aged 6 – 10 are often still learning to read, and find it easier to process information from pictures rather than text. Tool tips and brief onscreen explanations clarify the functions attached to icons, and support the children in associating images with functionality. The Comic Sans MS font in 14 point is used in KMB, as it is the font that is most attractive and easiest/fasted to read for children [2].

3.4. Expert and Participant Design Review

A second expert review was conducted on an 'alpha' prototype of KMB, uncovering a few minor issues with functionality (eg, the lack of an exit button) and icons (the rating system was changed from stars to an image more familiar to children: the smiley face). Appropriate modifications were made, to create a 'beta' prototype.

All participants in the initial design evaluation study (Section 3.1) took part in a formal review of the KMB beta. The goals were: to uncover remaining usability problems; to investigate how well the KMB design addresses the issues and desires raised by the children in the initial study; and to examine the 'learnability' of KMB (the extent to which it can be sued without formal training) [7]. Learnability is particularly important for entertainment-oriented software—struggling through a tutorial detracts from the user experience.

Again, minor usability problems were uncovered. The most significant insights gained were in the use of skins, the suitability of the icons, the song metadata displayed, and the box metaphor. One of the older participants (10 years old), dismissed all of the skins provided as being

"*too kiddie*" and "*not cool*"; the boys thought that some skins were "*too girlie*". A child's interests and tastes change dramatically from six to ten, as the child begins to aspire to a more mature appearance and activities. A full version of KMB should allow the user many more possibilities for tailoring the interface to support their emerging image of themselves.

For each of the icons, the participants were asked to speculate on the functionality represented. All of the participants were able to identify the function represented by most of the icons, and the children felt that any remaining uncertainty could be easily cleared up by reading the tooltips text.

Another concern was the minimal song metadata displayed by KDM (song title, artist, and user-provided rating). The expert reviewers speculated that children might wish to view additional descriptors. The participants, however, were emphatic that they found the larger amount of metadata to be confusing.

Participants did not find the 'box' metaphor to be compelling; given that users do not physically drag items into a box, the analogy is indeed weak. Two of the children did, however, suggest reinforcing the visual link to the metaphor by always including an image within box outlines—a suggestion that we adopted.

4. THE KMB INTERFACE

4.1. Login Window and Parental Settings

The *Login window* allows multiple users to create their own account for the organizer—each user will have their own music collections, and can organize music the way they like. Before a user can access KBM by logging into in their account, the parental settings must be configured (Figure 1). Only songs stored within the folders specified in the *Music settings* section folders are imported to the organizer window automatically; this aids children in importing music into their organizer while giving parents control over the songs that KMB contains. The *Picture setting* similarly restricts the images used in KMB (for to identify playlists and users). Parents can further restrict the import of songs whose metadata contain terms specified in the *Filter settings*. *Gift setting* identifies the default directory where KMB 'gift boxes' are to be stored. The grayscale, standard 'adult software' appearance of the Parental Settings window was deliberately chosen to make the adults feel more comfortable using that part of the software, and to make it less appealing to children.

4.2. The Organizer Window

Before the primary organizer window displays, KMB loads all the songs from the music folders that were added from the Parental Settings window into the Music Library. The Organizer window comprises the *music box* (Playlists) manager, the *music library* display, *visualisation* and *global control* (Figure 2).

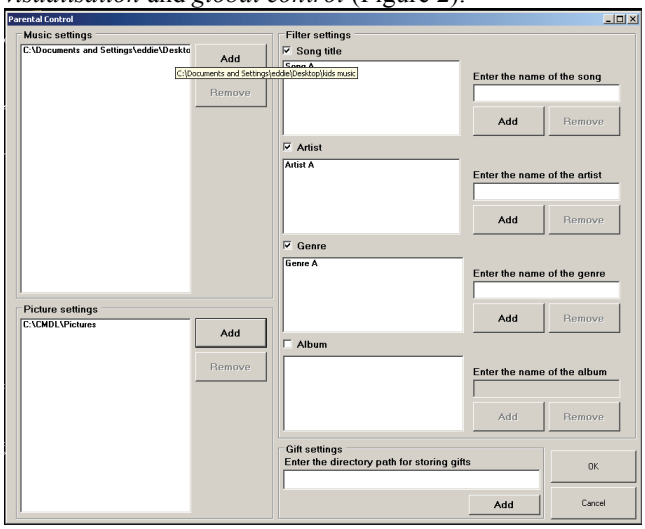

Figure 1. Parental Settings Window

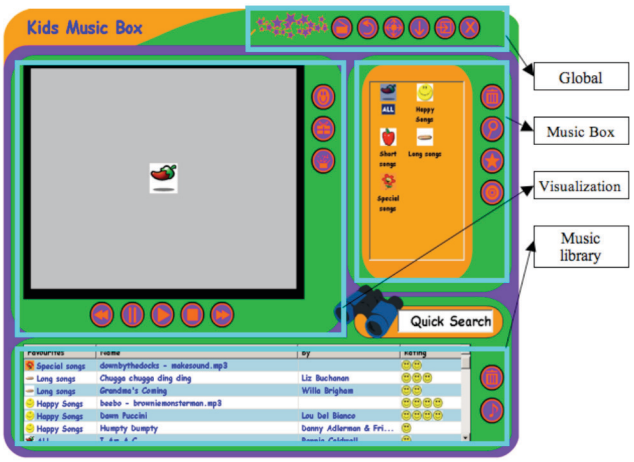

Figure 2. The Organizer window

The main function of the *Music Box* area is to display all the music boxes (playlists) that the user has created. The music boxes are displayed by a picture along with the name for the box. There will always be one box that is selected. All the songs that belong to this box will be displayed in the *Music Library* area. To change to another box, the user simply clicks on the box's icon.

The four vertical buttons in the Music Box area are, from top to bottom, the *remove music box*, *search for songs*, *make a new music box* and *get songs from CDs*.

4.3. Searching for songs

Users can search for songs by the title, artist, playlist ("music box"), and song rating (Figure 3). Search fields that are not expected to include very large numbers of values (artist, music box, and rating) prompt the user with a pull-down list of values. Metadata for songs matching the query are listed immediately below the query.

Figure 3. Searching for songs by The Tomato Collection Friends

4.4. The Music Library area

The music library displays all the songs that are contained in the currently selected music box. The main functionalities for the *Music Library* are: *sort songs within a music box, move songs to different music boxes*, *assign ratings to songs*, *performing the quick search function*, *import songs* and *remove songs* (Figure 4a)

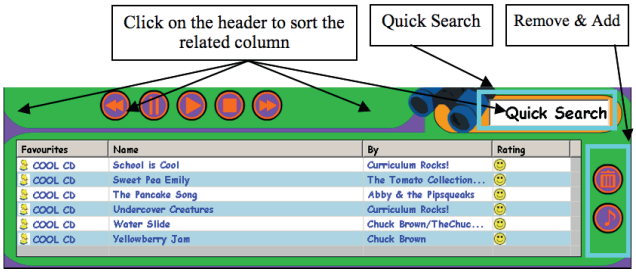

Figure 4a. The Music Library area

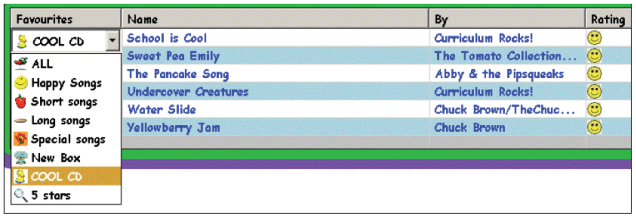

Figure 4b. Move a song to a different music box

Song metadata is limited to *song title*, *artist* and the *song rating*. Songs can be sorted by all these attributes. To move a song to a different music box, the user clicks on the Favourites label for the song and selects a new music box from the drop down box (Figure 4b). Users can similarly assign ratings (smiley faces) to individual songs. To remove a song, the user clicks on the song to select it and then clicks the Remove button. A song is added by clicking the Add button, and then following through the import dialog that appears.

4.5. The Visualisation area

The *Visualisation area* occupies largest screen area of the interface. This area has multiple functionalities: *image viewer*, a *playground with a number of Macromedia Flash games* and *music visualisations* by using the embedded *Window's Media Player* (Figure 5). The games and visualisations in the current KMB are essentially samples – in a non-prototype version, it would be more appropriate for users to manually insert their own games. The user can switch between different image albums, games and visualizations by clicking on the Switch Display button or using the large arrows to scroll between possibilities. The music playing buttons (play, pause, go to beginnning/end) are included in this area to reinforce the connection between the song and the playground activity. Note that we use the conventional music/video player icons; while these are not intuitive, it seems more reasonable to support children in learning standard icons than to require them to learn controls idiosyncratic to KMB.

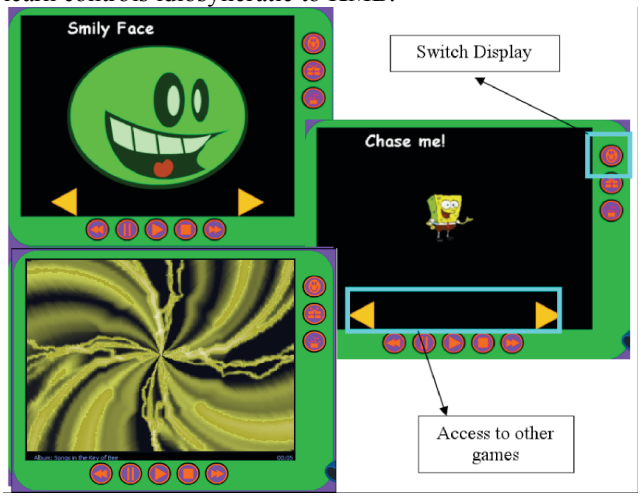

Figure 5. A range of games, image albums, and visualizations

4.6. Making a Gift

Clicking on the Make a Gift button (middle vertical button in the Music Library window) invokes a wizard that guides the user through selecting the songs, cover image, item for the visualization area, and a message to the recipient (Figure 6). The gift is packaged as a single file that is transferred to the recipient's computer. When opened, the gift is stored in KMB as a new music box within the recipient's music box library.

5. CONCLUSIONS

This paper has described the design, and the design process, for a music organizer for children. The most significant design features to adapt the organizer concept

for children are: a 'playground' to interact with while listening to songs; a set of themed interface skins to allow children to tailor the interface to their own tastes; a parental assistance window to support the child's caregiver in ensuring that appropriate images/music are included in KMB; the movement of difficult file management tasks from the user to the system; and an appealing and usable interface and interaction design.

Figure 6. An opened gift

However, the true test of a design lies in its use (or lack of use) over time [7]. To further evaluate the acceptability of KMB to its target users, it was installed on three of the participants' personal computers in their homes for a month, together with a logging application. The logging software created a timestamped log entry for each user action, and also created a screenshot every 20 so that the researchers could manually clarify ambiguous log states. This version of KDM also included popups triggered by starting up and exiting the software; the popups contained brief questions about why the child was using the system and the quality of the user experience.

The three participants engaged in a total of 85 sessions, with a median session length of about 15 minutes. The users' first few sessions were with their parents, but they began using KMB independently as they became more familiar with it. This pattern was expected, as parental assistance is necessary to set up the songs and images that the children can use. Tooltips were used extensively in earlier sessions, but then their display rapidly tapered off—indicating that the icons are indeed well linked to functionality, and that the system learnability is acceptable. For over 80% of the usage sessions, the participants reported that KMB was fun and easy to use.

The 'playground' was well received; for 33 of the 85 sessions the main reason for using KMB was to play with the games, rather than to organize or listen to music. In retrospect, this should not have been surprising—adults frequently listen to music as they play computer games or browse the Web [8]. This does point to the need for a variety of games and visualizations—and perhaps to make it easier to run KMB in the background to other games or physical activities.

None of the participants used the 'make a gift' feature, likely because a KMB gift can only be opened by another KMB. Evaluation of this feature has to wait until a significant user base develops—or the gift feature is redesigned to be compatible with conventional organizers.

Participant Design has enabled us to create a music organizer tailored to the eight children who worked with us in this study. The next step is to test our design with a larger pool of potential users. To this end, we plan to make KMB available as Open Source, perhaps as a One Laptop Per Child resource (www.olpc.org).

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RHYTHM COMPLEXITY MEASURES: A COMPARISON OF MATHEMATICAL MODELS OF HUMAN PERCEPTION AND PERFORMANCE

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ABSTRACT

Thirty two measures of rhythm complexity are compared using three widely different rhythm data sets. Twenty-two of these measures have been investigated in a limited context in the past, and ten new measures are explored here. Some of these measures are mathematically inspired, some were designed to measure syncopation, some were intended to predict various measures of human performance, some are based on constructs from music theory, such as Pressing's cognitive complexity, and others are direct measures of different aspects of human performance, such as perceptual complexity, meter complexity, and performance complexity. In each data set the rhythms are ranked either according to increasing complexity using the judgements of human subjects, or using calculations with the computational models. Spearman rank correlation coefficients are computed between all pairs of rhythm rankings. Then phylogenetic trees are used to visualize and cluster the correlation coefficients. Among the many conclusions evident from the results, there are several observations common to all three data sets that are worthy of note. The syncopation measures form a tight cluster far from other clusters. The human performance measures fall in the same cluster as the syncopation measures. The complexity measures based on statistical properties of the inter-onset-interval histograms are poor predictors of syncopation or human performance complexity. Finally, this research suggests several open problems.

1 INTRODUCTION

Many music researchers consider rhythm to be the most important characteristic of music. Furthermore, one of the main features of rhythm is its complexity. Threfore measures of the complexity of a rhythm constitute key features useful for music pattern recognition and music information retrieval, as well as ethnomusicological analyses of world music [17, 18]. Since the notion of complexity is flexible, it is not surprising that in the past a variety of different measures of complexity has appeared in the literature. Areas where such measures have been applied range from psychology, enginneering, computer science and mathematics, to music theory. Given such a wide range of applicable fields, different techniques for measuring complexity have been developed. For example, one can analyze a rhythm's binary sequence representation, ask listeners to rate a rhythm's complexity, or ask musicians to perform a rhythm. Therefore, in our work, we include measures of information and coding complexity, performance complexity, and cognitive complexity. Furthermore, there are traditional concepts in music such as *syncopation* [10] which may also be considered as measures of rhythm complexity [7, 8].

With the exception of [7, 8], previous research on rhythm complexity has been limited to determining how good a feature it is for music pattern recognition, or how well it models human judgements of complexity [17, 18]. Moreover, for such studies researchers have used data (families of rhythms) that were generated artificially and randomly with some constraints. Here, we not only use a large group comprised of 32 measures of complexity that employ a wide variety of measurement techniques, but we also validate these measures against human judgements of perceptual, meter, and performance complexity using three diverse data sets.

2 COMPLEXITY MEASURES

One can broadly categorize the complexity measures used in this study into two distinct categories: human performance measures directly obtained from psychological experiments, and measures obtained from mathematical models of rhythm complexity. The human performance measures can be subdivided into three types: *perceptual* complexity, *meter* complexity, and *performance* complexity. Perceptual complexity is obtained by asking human subjects to judge complexity as they listen to rhythms. Meter complexity is obtained by measuring how well the human subjects are able to track the underlying *metric beat* of a rhythm. It is worth noting that some researchers, for example in music psychology [4], refer to the metric beat as the *pulse*.

Here we reserve the word *pulse* for the largest duration interval that evenly divides into all the inter-onset onsets (IOI) present in a family of rhythms. This is common terminology in ethnomusicology and music technology. Performance complexity measures pertain to how well the subjects can reproduce (execute, play-back) the rhythms, usually by tapping. The mathematical models can be subdivided into two main categories: those that are designed to measure *syncopation*, and those that are designed to measure *irregularity*. The irregularity measures can be divided into *statistical* and *minimum-weight-assignment* measures.

Due to lack of space, we cannot provide a detailed description of all the complexity measures tested. Thus we list the complexity measures with each corresponding essential reference in the literature for further information, along with a label in parentheses pertaining to the phylogentic tree labels used in Figures 1, 2, and 3. Measures of syncopation are listed first. The Longuet-Higgins and Lee measure (*lhl*) [4, 14], along with Smith and Honing's version (*smith*) [19], take advantage of a metric hierarchy of weights [13] to calculate syncopation. A variation of Toussaint's metrical complexity (*metrical*) [21] and Keith's measure (*keith*) [10] also use this hierarchy to judge syncopation. The Weighted Note-to-Beat Distance (*wnbd, wnbd2, wnbd4, wnbd8*) [7] uses the distance from onsets to *metric beats* to gauge syncopation.

Second, we list the measures regarding mathematical irregularity. IOI histogram measures for entropy (*ioi-g-h, ioi-l-h*), standard deviation (*ioi-g-sd, ioi-l-sd*), and maximum bin height (*ioi-g-mm, ioi-l-mm*) were used to determine the complexity of both global (full) IOIs [24] and local (relative, adjacent) IOIs [18]. Also, pertaining to entropy calculations are the Coded Element Processing System (*ceps*) [26], H(*k*-span) complexity (*hk*) [25], and the H(run-span) complexity (*hrun*) [25], which all measure the uncertainty [5] of obtaining sub-patterns in a rhythm. The directed swap distance (*dswap, dswap2, dswap4, dswap8*) [1] computes the minimum weight of a linear assignment between onsets of a rhythm and a meter with an onset at every second, fourth, or eigth pulse, and also the average over each meter. Two other measures, Rhythmic Oddity (*oddity*) [22] and Off-Beatness (*off-beatness*) [22] take a geometric approach.

Third, those measures which do not easily fall into a category are listed. These include the Lempel-Ziv compression measure (*lz*) [12], Tanguiane's [20] complexity measure, which looks at sub-patterns at each metrical beat level, and Pressing's Cognitive Complexity measure (*pressing*) designed on the basis of music theory principles, which generates rhythmic patterns at each metrical beat, assigning appropriate weights to special patterns [16]. Furthermore, Tanguiane's measure uses the max (*tmmax*) and average (*tmavg*) complexities over different metrical beat levels. In addition, derivatives (*tmuavg, tmumax*) without the restriction of subpatterns starting with an onset, were tested.

3 EXPERIMENTAL DATA

The measures of complexity in § 2, were compared using three rhythm data sets. Each data set had been compiled to test human judgements regarding the perceptual, meter, and performance complexities of the rhythms. The first data set shown in Table 1 was synthesized by Povel and Essens in 1985 [15] and then later studied by Shmulevich and Povel in 2000 [17]. The second data set shown in Table 2 was created by Essens in 1995 [2]. The third data set shown in Table 3 was generated by Fitch and Rosenfeld in 2007 [4]. In addition to the rhythms themselves, the results of several human performance complexity measures used in this work are contained in Tables 1, 2, and 3. In the following we describe the methodologies of Povel and Essens [15], Shmulevich and Povel [17], Essens [2], and Fitch and Rosenfeld [4], used to obtain the human judgements of complexity.

3.1 Povel and Essens 1985

Previous work by Povel and Essens [15] studied the reproduction quality of temporal patterns. The rhythms, shown in Table 1, were presented to the participants in random order. For each presentation, the participant was asked to listen to the pattern, and then reproduce the pattern by tapping [15]. Once the participant had felt they could reproduce the rhythm, they stopped the audio presentation and proceeded to then tap the pattern they just heard, repeating the pattern 4 times. Afterwards, they could choose to move to the next rhythm or repeat the one they had just heard [15]. From this experiment, we derive an empirical measure for the reproduction difficulty of temporal patterns; i.e., rhythm performance complexity. This was based on Povel and Essens' *mean deviation percentage* which calculates the amount of adjacent IOI error upon reproduction [15]. See column 3 of Table 1.

3.2 Shmulevich and Povel 2000

Shmulevich and Povel [17] studied the perceptual complexity of rhythms using the same data as Povel and Essens [15]. All participants were musicians with musical experience averaging 9.2 years [17]. A pattern was repeated four times before the next was randomly presented. The resulting perceptual complexity in column 4 of Table 1 represents the average complexity of each rhythm across all participants.

3.3 Essens 1995

A study of rhythm performance complexity was conducted by Essens [2]. The rhythms used for that study are shown in Table 2. The procedure Essens used to test the reproduction accuracy of rhythms was very similar to that of Povel and Essens [15]. We use the mean deviations of Essens to rank the rhythms by increasing complexity, as seen in column 3

of Table 2. Essens also studied the perceptual complexity of rhythms [2]. Participants were asked to judge the complexity of each rhythm in Table 2 on a 1 to 5 scale where 1 means very simple and 5 means very complex [2]. Note that some participants had been musically trained for at least 5 years. The order of the patterns was random. The perceptual complexity in column 4 of Table 2 is the average complexity over the judgements from each subject.

3.4 Fitch and Rosenfeld 2007

Most recently, Fitch and Rosenfeld [4] conducted an experimental study of metric beat-tracking or, in their terminology, pulse-tracking (i.e., rhythmic *meter complexity*) and rhythm reproduction (i.e., *performance complexity*). The rhythms used in the experiments are shown in Table 3. These rhythms were generated in such a way as to vary the amount of syncopation among the rhythms, as measured by the Longuet-Higgins and Lee syncopation measure [14].

The metric beat-tracking experiment yielded two measures of meter complexity [3]. The first pertained to how well participants could tap a steady beat (*beat tapping error adjusted* for tempo) when different rhythms were played. The second counted the number of times (*number of resets*) the participant tapped the metric beat exactly in between the points where the metric beat should be [4]. The values are shown in columns 3 and 4 of Table 3. The second experiment for rhythm reproduction accuracy was interleaved with the metric beat-tracking experiment. Hence the subject now taps the rhythm just heard from experiment 1 while the computer provides the metric beat [4]. The adjacent IOI error of the target and reproduced rhythms gives a performance complexity shown in column 5 of Table 3.

4 RESULTS

We adhered to the following procedure to validate the complexity measures in § 2 using the three rhythm data sets. The complexity scores were obtained using the rhythms as input for each measure. The Spearman rank correlation coefficients [11] between all pairs of rankings of the rhythms according to the computational and empirical measures for each rhythm data set were calculated.

Phylogenetic trees were used to visualize the relationships among the correlation coefficients. This technique has proved to be a powerful analytical tool in the computational music domain [1, 8, 21, 22, 23]. The program SplitsTree [9] was used to generate the phylogenetic trees using the BioNJ algorithm [6]. Figures 1, 2, and 3, picture the phylogenetic trees where the distance matrix values are the correlation coefficients subtracted from one. Each tree yields a fitness value greater than or equal to 94.0 on a 100.0 scale. The least-squares fitness is the ratio of A/B where A is the sum of the squared differences between the geodesic distances

No.	Rhythm	Performance Complexity Povel and Essens	Perceptual Complexity Shmulevich and Povel
1	x x x x x . . x x . x . x . . .	5	1.56
2	x x x . x . x x x . . x x . . .	1	2.12
3	x . x x x . x x x . . x x . . .	0	2.08
4	x . x . x x x x x . . x x . . .	2	1.88
5	x . . x x . x . x x x x x . . .	3	1.80
6	x x x . x x x . x x . . x . . .	9	2.44
7	x . x x x x . x x . . x x . . .	7	2.20
8	x x . . x x x x x . x . x . . .	4	2.56
9	x x . . x . x x x . x x x . . .	14	3.00
10	x . x x x . x x x x . . x . . .	18	2.04
11	x x x . x x . . x x . x x . . .	19	2.76
12	x x . x x x x . x . . x x . . .	15	2.72
13	x x . x x . x x x x . . x . . .	13	3.00
14	x x . . x x . x x . x x x . . .	27	3.16
15	x . . x x x . x x x . x x . . .	10	2.04
16	x x . x x x x . x x . . x . . .	11	2.88
17	x x . x x x . x x x . . x . . .	17	2.60
18	x x . x x x . . x x . x x . . .	22	2.60
19	x x . . x x . x x x x . x . . .	21	2.64
20	x x . . x x . x x x . x x . . .	25	3.24
21	x x x x x . x x . x . . x . . .	29	3.08
22	x x x x . x . . x x x . x . . .	20	3.04
23	x x x . . x x . x x x . x . . .	16	3.04
24	x . x x x . . x . x x x x . . .	6	2.56
25	x . x . . x x x x . x x x . . .	8	2.56
26	x x x x . x . x . . x x x . . .	26	2.84
27	x x . x x x . x . . x x x . . .	23	3.60
28	x x . x . . x x x . x x x . . .	32	2.68
29	x . x x x x . x . . x x x . . .	28	3.28
30	x . . x x x x . x x . x . . .	21	3.08
31	x x x x . x x x . . x . x . . .	30	3.52
32	x x x x . . x x . x x . x . . .	31	3.60
33	x x . x x x x . . x x . x . . .	24	3.04
34	x x . x . . x x x x x . x . . .	33	2.88
35	x . x . . x x x . x x x x . . .	12	3.08

Table 1. Thirty-five rhythms from Povel and Essens with the Performance Complexity and Perceptual Complexity.

No.	Rhythm	Performance Complexity Essens	Perceptual Complexity Essens
1	x x x . x x x . x x x . x x . .	0	2.2
2	x x x . x . x x x . x x x x . .	8	3.1
3	x . x x x . x x x . . x x x . .	4	3.2
4	x . x x x . . x x . x x x x . .	19	2.9
5	x x x . x x x . x x . x x . . .	2	2.2
6	x x x . x . . x x . x . x x . .	7	3.1
7	x x x x x x x . x x x . x x x .	10	2.6
8	x x x . x x x x x . . x x x . .	5	4.2
9	x x x x x x . x x . x x x . . .	13	2.9
10	x . x . x . x x x . x x . .	6	2.8
11	x x x x x x x . x x x . x . x .	1	3.1
12	x x x . x x . . x . x . x . . .	3	2.5
13	x . . x x x x . x x . . x x . .	20	3.5
14	x . x x x x . x x x . x x x . .	12	2.5
15	x . . x x x . x x x . x x x . .	14	2.4
16	x . . x x x . x x x x . x x . .	11	3.0
17	x x . x x x . x x x x . x . . .	17	3.0
18	x . . x x x x x x x . x x x . .	18	3.1
19	x . x . x x . x x x . x x x . .	22	2.4
20	x x . x x x x . x x . x x . . .	16	3.2
21	x x . x x x . x x x x x x . . .	15	2.4
22	x x . . x x . x x x x x x . . .	11	2.9
23	x . x . x x . x x x x x x x . .	21	2.7
24	x x . x x x x x x . x . x x . .	9	3.8

Table 2. Twenty-four rhythms from Essens with Performance Complexity and Perceptual Complexity.

between pairs of leaves in the tree, and their corresponding distances in the distance matrix, and B is the sum of the squared distances in the distance matrix. Then this value is subtracted from 1 and multiplied by 100 [27]. Note that the phylogenetic tree is used here as a visualization tool, and not in order to obtain a phylogeny of complexity measures.

5 DISCUSSION AND CONCLUSION

There are several noteworthy observations common to all three data sets. The syncopation measures form a tight cluster far from the other clusters. The human performance mea-

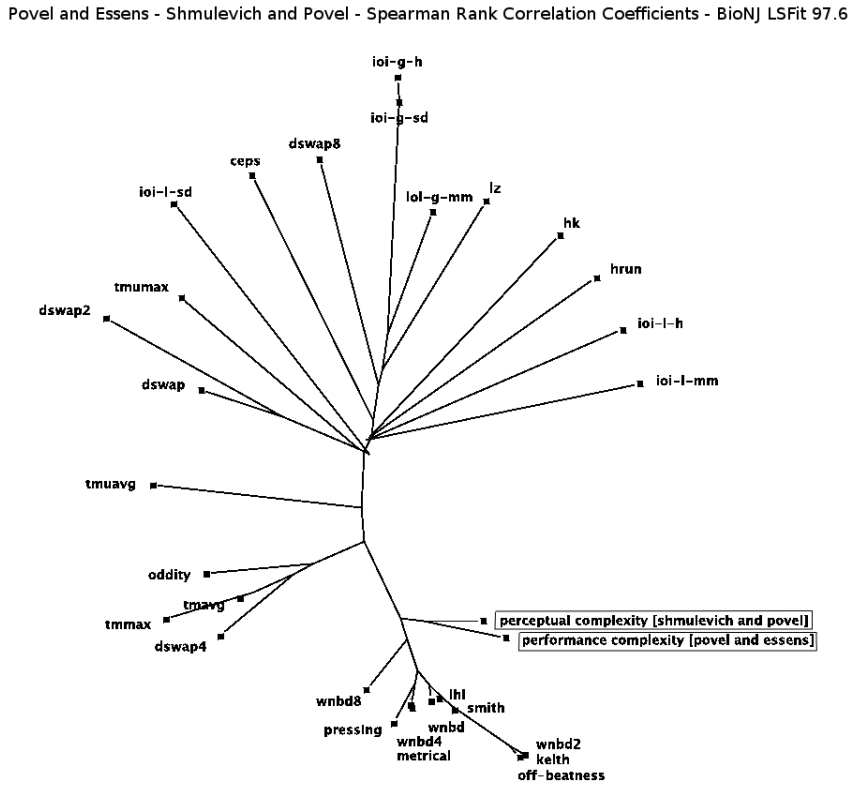

Figure 1. BioNJ of measures compared to Shmulevich and Povel and Povel and Essens human judgements.

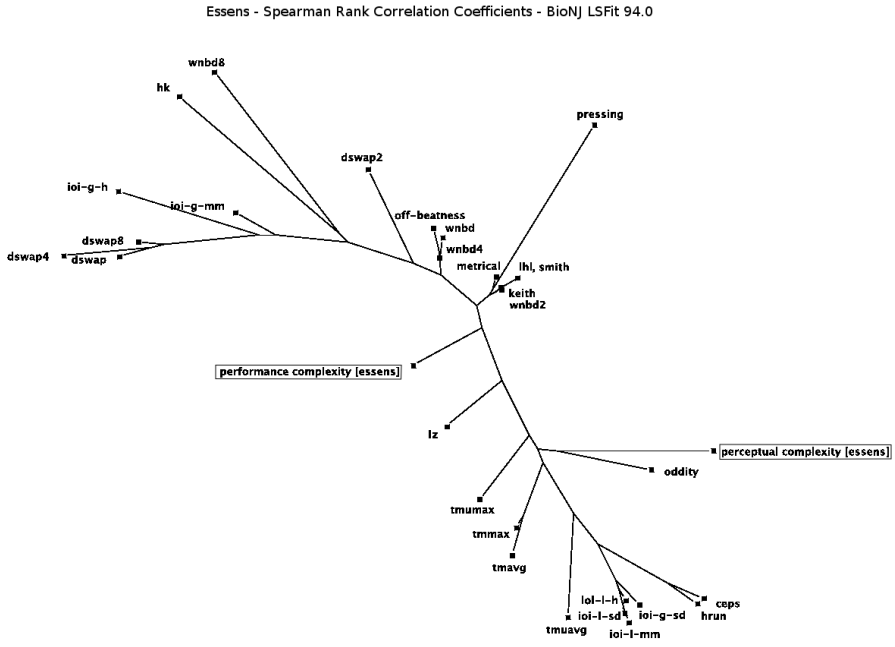

Figure 2. BioNJ of measures compared to Essens' human judgements.

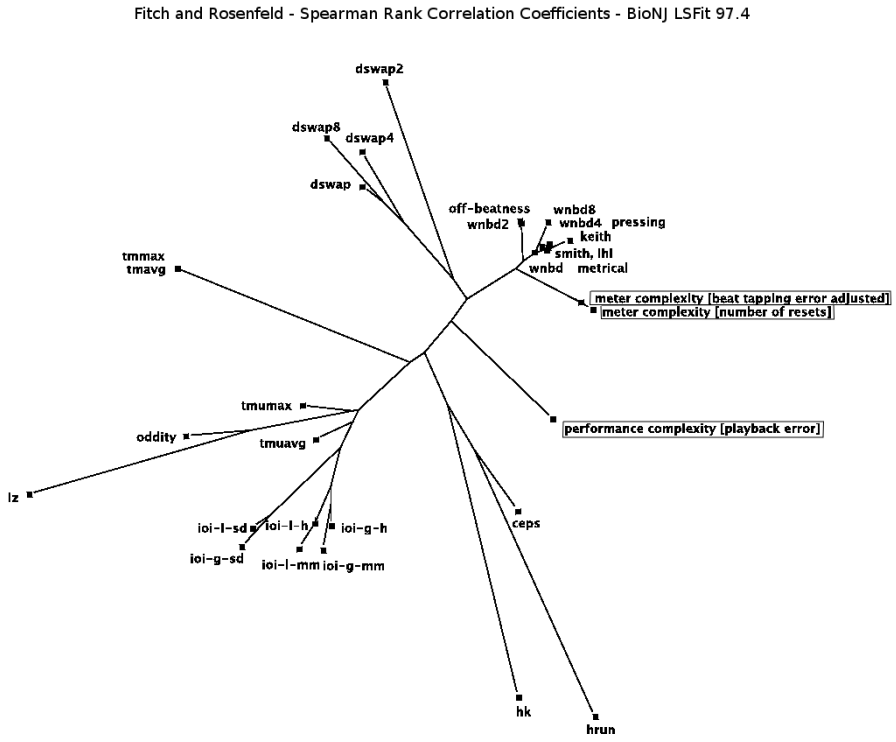

Figure 3. BioNJ of measures compared to Fitch and Rosenfeld's human judgements.

No.	Rhythm	Meter Complexity Beat Tapping Adjusted	Meter Complexity Number of Resets	Performance Complexity Play-back Error
1	x . . . . . x . x . . . . . x .	0.075	2.500	0.138
2	x . . . x . . . x . . . x . x .	0.082	2.250	0.145
3	x . x . x . . . . x . x .	0.075	2.313	0.153
4	x . . . x x x . . . . . x . . .	0.119	8.750	0.257
5	. . x . . . x . x . x . x . . .	0.103	5.500	0.133
6	. . x . . . x . x . . . x . x .	0.082	3.063	0.235
7	x . . . . . x . . x . . . . x .	0.112	6.000	0.215
8	x . . x . . . x . . x . x . . .	0.110	5.188	0.208
9	x . . . . x . . . x . . . x x .	0.141	6.938	0.250
10	x . . . . x . . . x . . . x . .	0.144	10.375	0.171
11	. . x . . . x . . . x x . x . .	0.130	6.875	0.220
12	. x . . . x . . . x . . x . x .	0.124	6.438	0.226
13	. . x . . . x x . . . x . x . .	0.130	6.965	0.387
14	. . x . . x . . . . . x . . . x	0.159	11.688	0.239
15	. . . x . x . . . . x x . . . x	0.172	13.688	0.485
16	x . . . x . x x x	0.085	2.625	0.173
17	x . x . x . . . . . x . x . . .	0.077	2.313	0.179
18	. . x . x . . . x . x . x . . .	0.077	2.438	0.182
19	x . x . . . x . x . . x	0.074	1.938	0.252
20	. . x . . . x . x . . . x . . .	0.098	3.375	0.142
21	x . . . . x . x . . x . x . . .	0.161	11.063	0.305
22	. . x . . . x . x . x . . . x .	0.129	8.500	0.321
23	x x . . . x . . . . x . . . x .	0.145	7.375	0.320
24	. x . . . x . . . x x . x . . .	0.134	7.188	0.265
25	. . x . . . x . . . x . x . .	0.146	8.625	0.176
26	. . x . . . x . . . x x . . . x	0.118	6.500	0.326
27	. . x . . x . . . . x . . x . x	0.117	6.188	0.368
28	. . x x . . . x . . . x . x . .	0.154	10.813	0.344
29	. x . x . x . . . x	0.191	15.750	0.185
30	. x . x . . . x . . . x . . . x	0.164	11.938	0.158

Table 3. Thirty rhythms from Fitch and Rosenfeld with Meter Complexity and Performance Complexity.

sures fall in the same cluster as the syncopation measures. The complexity measures based on statistical properties of the inter-onset-interval histograms appear to be poor predictors of syncopation or of human performance complexity.

There are also some important differences between the three figures. The overall appearance of clusters is much stronger in Figure 3 than in the other two. This is perhaps due to the fact that the rhythms used in Figure 3 are much more realistic and sparser than the rhythms used in Figures 1 and 2. Similarly, the six IOI (inter-onset-interval) measures are scattered in Figures 1 and 2, but are in one cluster in Figure 3. The cognitive complexity measure of Pressing, designed on the basis of principles of music perception falls squarely in the group of syncopation measures in Figures 1 and 3. However, in Figure 2, although it falls into the syncopation cluster, it is quite distant from the other measures, probably because of the great density of the rhythms in this data set. Also worthy of note is a comparison of the human meter complexity measures with the human performance (playback) measure. In Figure 3 we see that the meter complexity is considerably closer to the syncopation measures than the play-back performance measure. This suggests that the mathematical syncopation measures are better predictors of human meter complexity than performance complexity.

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CONTENT-BASED MUSICAL SIMILARITY COMPUTATION USING THE HIERARCHICAL DIRICHLET PROCESS

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ABSTRACT

We develop a method for discovering the latent structure in MFCC feature data using the Hierarchical Dirichlet Process (HDP). Based on this structure, we compute timbral similarity between recorded songs. The HDP is a nonparametric Bayesian model. Like the Gaussian Mixture Model (GMM), it represents each song as a mixture of some number of multivariate Gaussian distributions However, the number of mixture components is not fixed in the HDP, but is determined as part of the posterior inference process. Moreover, in the HDP the same set of Gaussians is used to model all songs, with only the mixture weights varying from song to song. We compute the similarity of songs based on these weights, which is faster than previous approaches that compare single Gaussian distributions directly. Experimental results on a genre-based retrieval task illustrate that our HDP-based method is both faster and produces better retrieval quality than such previous approaches.

1 INTRODUCTION

We develop a new method for estimating the timbral similarity between recorded songs. Our technique is based on the hierarchical Dirichlet process, a flexible Bayesian model for uncovering latent structure in high-dimensional data.

One approach to computing the timbral similarity of two songs is to train a single Gaussian or a Gaussian Mixture Model (GMM) on the Mel-Frequency Cepstral Coefficient (MFCC) feature vectors for each song and compute (for the single Gaussian) or approximate (for the GMM) the Kullback-Leibler (K-L) divergence between the two models [1]. The basic single Gaussian approach with full covariance matrix ("G1" [2]) has been successful, forming the core of the top-ranked entries to the MIREX similarity evaluation task two years running [3, 4].

Although MFCC data are not normally distributed within songs, using a richer model such as the GMM to more accurately represent their true distribution provides little or no improvement in numerous studies [2, 5, 1]. This suggests that a "glass ceiling" has been reached for this type of representation. Moreover, the computational cost of the Monte Carlo estimation procedure involved in comparing two GMMs is orders of magnitude more than that incurred by computing the K-L divergence between two single Gaussians exactly. This is a very significant issue if we want to compute similarity matrices for large sets of songs, since the number of comparisons between models that must be done grows quadratically with the number of songs.

Another approach [6] produced results statistically indistinguishable from the other top algorithms in MIREX 2007 by using a mid-level semantic feature representation to compute similarity. Using painstakingly human-labeled data, Barrington et al. trained GMMs to estimate the posterior likelihood that a song was best characterized by each of 146 words. These models then produced a vector for each test song defining a multinomial distribution over the 146 semantic concepts. To compute the dissimilarity of two songs, the K-L divergence between these multinomial distributions for the songs was computed.

The success of this method suggests that alternative statistical representations of songs are worth exploring. Rather than take a supervised approach requiring expensive hand-labeled data, we make use of the Hierarchical Dirichlet Process (HDP), which automatically discovers latent structure within and across groups of data (songs, in our case). This latent structure generates a compact alternative representation of each song, and the model provides a natural and efficient way of comparing songs using K-L divergence.

2 HDP-BASED SIMILARITY USING LATENT FEATURES

The hierarchical Dirichlet process (HDP) is an extension of the Dirichlet process (DP), a nonparametric Bayesian model of mixtures of an unknown number of simple densities. We first outline the DP and then describe how we model songs with an HDP.

2.1 Dirichlet Process Mixture Models

The Gaussian Mixture Model (GMM) is a generative process that assumes that each of our feature vectors was generated by one of K multivariate Gaussian distributions. To

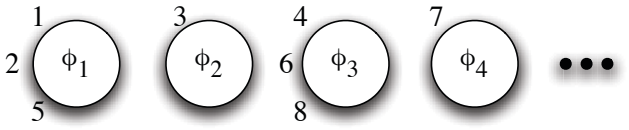

Figure 1. Four tables and eight customers in a Chinese Restaurant Process (CRP). In this example, the 1st, 3rd, 4th, and 7th customers all sat at an empty table, whereas the 2nd, 5th, 6th, and 8th sat at existing tables. The 9th customer will sit at table 1, 2, 3, or 4 with probabilities $\frac{3}{8+\alpha}$, $\frac{1}{8+\alpha}$, $\frac{3}{8+\alpha}$, and $\frac{1}{8+\alpha}$ respectively, or will sit at a new table with probability $\frac{\alpha}{8+\alpha}$

draw a new vector y_t, the process first chooses a mixture component index $z_t \in 1...K$ with probability π_{z_t} (where π is a vector of mixture probabilities summing to one), then draws the vector from the z_tth Gaussian distribution. Given K and a set of vectors assumed to have been generated by a GMM, algorithms such as Expectation-Maximization (EM) can find a maximum-likelihood estimate of the mixture probabilities $\pi_{1...K}$, the parameters defining the K Gaussians $\mu_{1...K}$ and $\Sigma_{1...K}$, and which mixture component z_t generated each vector y_t.

A nagging issue in mixture modeling is model selection, i.e., choosing the number of components K with which to explain the data. Recent work in nonparametric Bayesian statistics has produced models such as the Dirichlet Process Mixture Model (DPMM) that sidestep this issue. Where the GMM assumes the existence of K mixture components, the DPMM [7] assumes the existence of a countably infinite set of mixture components, only a finite subset of which are used to explain the observations.

A traditional metaphor for the way a DP generates data is called the Chinese Restaurant Process (CRP). In the CRP, we imagine a Chinese restaurant with an infinite number of communal tables and a positive scalar hyperparameter α. The restaurant is initially empty. The first customer sits at the first table and orders a dish. The second customer enters and decides either to sit at the first table with probability $\frac{1}{1+\alpha}$ or a new table with probability $\frac{\alpha}{1+\alpha}$. When sitting at a new table the customer orders a new dish. This process continues for each new customer, with the tth customer choosing either to sit at a new table with probability $\frac{\alpha}{\alpha+t-1}$ or at the kth existing table with probability $\frac{n_k}{\alpha+t-1}$, where n_k is the number of other customers already sitting at table k. Notice that popular tables become more popular, and that as more customers come in they become less and less likely to sit down at a new table.

We obtain a DPMM from a CRP as follows. The "dishes" in the CRP correspond to probability density functions, and the process of "ordering" a dish k corresponds to drawing the parameters ϕ_k to a PDF from a prior distribution H over those parameters. (For example, each dish ϕ_k can be a Gaussian with parameters $\{\mu_k, \Sigma_k\} = \phi_k \sim H$.) The process of a customer t choosing a table z_t corresponds to choosing a distribution ϕ_{z_t} from which to draw an observation y_t (in our case, a feature vector). Since customers in the CRP tend to sit at tables with many other customers, the DPMM tends to draw points from the same mixture components again and again even though it has an infinite number of mixture components to choose from.

Analysis under a DPMM involves inferring the posterior distribution over its latent parameters conditioned on the data. This provides a partition of the data (feature vectors) into an unknown number of clusters (the number of tables) and the identities of the parameters ϕ (the means and covariances of the Gaussian mixture components). The posterior distribution $P(\phi, z_{1...T}|y_{1...T})$ of the set of mixture component parameters ϕ and the cluster labels for each feature vector $z_{1...T}$ to a DPMM conditioned on the data $y_{1...T}$ can be inferred using Markov Chain Monte Carlo (MCMC) methods such as Gibbs sampling [7]. For simple data, there will be relatively few unique cluster labels in z, but more clusters will be necessary to explain more complex data.

2.2 The Hierarchical Dirichlet Process

The Hierarchical Dirichlet Process (HDP) [8] is a model of *grouped data*, which is more appropriate than the DPMM for modeling songs represented as a collection of MFCCs. Rather than associate each song with a single table in the restaurant, each song is represented as a group of features which sit at a song-specific "local" restaurant. The dishes for this restaurant, however, are drawn from a "global" set of dishes. Thus, each song is represented as a distribution over latent components, but the population of latent components is shared across songs. Similarity between songs can be defined according to the similarity between their corresponding distributions over components.

The generative process underlying the HDP can be understood with the Chinese Restaurant Franchise (CRF). The CRF takes two hyperparameters α and γ. Each song j has its own CRP, and each feature vector $y_{j,t}$ chooses a table from CRP(α). If it sits down at a new table, then it chooses a dish for that table from a global CRP (with hyperparameter γ) shared by all songs – that is, it either chooses a dish that is already being served at some number of other tables m with probability proportional to m, or it chooses a new dish with probability proportional to γ.

Although we have defined the CRP as a sequential process, in fact data under a CRP are exchangeable – the probability of a seating plan under the CRP is the same regardless of the order in which the customers sat down. This allows us to think of the CRP as defining an implicit prior on infinite multinomial distributions over mixture components. In the DPMM, the infinite-dimensional vector of probabilities $\bar{\pi}$ defining such an infinite multinomial distribution is analo-

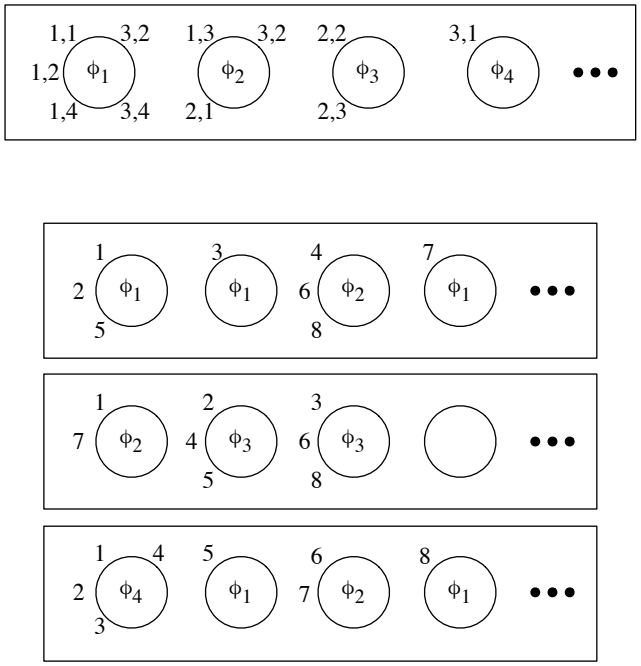

Figure 2. Chinese Restaurant Franchise (CRF) for three songs with eight observations. Below are three CRPs (corresponding to the three songs), and above is the global CRP from which the CRPs get their dishes. Each customer j, i sitting at a table in the global CRP corresponds to table i in restaurant j, and customer j, i's table membership in the global CRP determines the dish that is served at table i in restaurant j. If a new customer coming into a restaurant j sits down at a new table, then the dish for that table will be ϕ_1, ϕ_2, ϕ_3, or ϕ_4 with probability $\frac{5}{\gamma+11}$, $\frac{3}{\gamma+11}$, $\frac{2}{\gamma+11}$, or $\frac{1}{\gamma+11}$ respectively, or a new dish with probability $\frac{\gamma}{\gamma+11}$.

gous to the K-dimensional vector π in the GMM. The HDP has J such vectors $\bar{\pi}_{1...J}$, each of which defines a different distribution over the same mixture components.

We use Gibbs sampling to approximate the posterior distribution over the latent variables conditioned on observed data. The distribution is over the cluster partition assigning feature vectors to clusters and a truncated vector π_j defining the mixture proportions for each song over the finite subset of K mixture components that are actually assigned to observations. We let $\pi_{j,1...K} = \bar{\pi}_{j,1...K}$, and $\pi_{j,K+1} = 1 - \sum_{k=1}^{K} \bar{\pi}_{j,k}$, where $\pi_{j,K+1}$ is the probability of drawing an observation from a mixture component that has not been used to explain any feature vector in any song.

For a complete exposition of the HDP, including how to infer the posteriors for its parameters conditioned on data, see [8].

2.3 Representing Songs Using the HDP

The mixture components parameterized by $\phi_{1...K}$ capture the latent structure in the feature data, and the mixture proportion vectors $\pi_{1...J}$ express the feature data for songs $1...J$ in terms of that latent structure. ϕ and π_j together can describe the empirical distribution of feature vectors for a song j as richly as a GMM can, but the HDP does not require that we choose a fixed value of K, and represents the songs in a more compact way.

To compute the distance between two songs i and j, we can compute the symmetrized KL divergence between the posterior distributions $P(\pi_i|\beta, m)$ and $P(\pi_j|\beta, m)$ which are of the form

$$P(\pi_j|\beta, m) = \text{Dir}(\beta_1 + m_{j,1}, ..., \beta_K + m_{j,K}, \beta_{K+1}) \quad (1)$$

where $m_{j,k}$ is the number of tables in restaurant j serving dish k, β_k is the likelihood of choosing a dish k from the global CRP, and β_{K+1} is $1 - \sum_{k=1}^{K} \beta_k$, the likelihood of choosing a previously unseen dish in the global CRP.

This allows us to compare two songs in terms of the latent structure of their feature data, rather than directly comparing their distributions over the low-level features as the G1 algorithm and GMM-based algorithms do. The KL divergence between these two posteriors can be efficiently computed. The KL divergence between two Dirichlet distributions with parameters v and w each of length K is [9]:

$$D(\text{Dir}(v)||\text{Dir}(w)) = \log \frac{\Gamma(\sum v)}{\Gamma(\sum w)} + \sum_{s=1}^{K} \frac{\log(\Gamma(w_s))}{\log(\Gamma(v_s))} + \sum_{s=1}^{K} ((v_s - w_s)(\Psi(v_s) - \Psi(\sum v))$$

where $\Gamma(x)$ is the gamma function, $\Psi(x)$ is the digamma function (the first derivative of the log gamma function), and $\sum v$ and $\sum w$ denote the sum of the K elements of v and w respectively.

This is less expensive to compute than the KL divergence between two high-dimensional multivariate Gaussian densities. It can be sped up further by computing the gamma and digamma terms offline for each song.

2.4 Generalizing to New Songs

It is important that our approach be scalable to new songs not seen during training. Once we have inferred the global dish likelihoods β and the mixture component parameters $\phi_{1...K}$, we can infer the posterior distribution over the mixture proportions π_{J+1} for a new song $J + 1$ conditioned on β, ϕ, and the new data y_{J+1} using the same Gibbs sampling techniques originally used to train the model, holding all other parameters constant.

3 EVALUATION

In this section we describe the experiments we performed to evaluate our approach against G1, GK (the analogous algorithm for K-component GMMs), and an approach based on Vector Quantization (VQ).

3.1 South by Southwest Dataset

We test our approach on a dataset that we compiled from the South by Southwest (SXSW) 2007 and 2008 festivals' freely distributed "artist showcase" mp3s [10]. We selected a set of up to twenty mp3s (all by different artists to avoid biasing the results) for seven genres: country, electronic, hip-hop, jazz, metal, punk, and rock. Songs that we felt were unrepresentative of their genre were removed or replaced prior to any quantitative evaluations. There were fewer than 20 usable songs available for country (12), jazz (14), and metal (15), so those genres are slightly underrepresented. There are a total of 121 songs in the dataset.

3.2 Features

All models were trained on the same sets of feature vectors, which for each frame consisted of 13 MFCCs (extracted using jAudio [11]) combined with 26 delta features computed by subtracting the MFCCs for frame t from those at frame $t-1$ and $t-2$, for a total of 39 dimensions. Each frame was approximately 23 ms long, or 512 samples at the files' sampling rate of 22050 Hz, with a hop size of 512 samples (no overlap). 1000 feature vectors were extracted from the middle of each song.

3.3 Models Evaluated

3.3.1 G1

As described above, G1 models each song's distribution over feature vectors with a single multivariate Gaussian distribution with full covariance matrix. Models are compared using the symmetrized KL divergence.

3.3.2 K-component GMMs

We train K-component GMMs for each song using the E-M algorithm. The symmetrized KL divergence between models is approximated by drawing 1000 synthetic feature vectors from the trained models and evaluating their log likelihoods under both models [1]. This approach is evaluated for $K = 5$, 10, 20, and 30.

3.3.3 VQ Codebook

This algorithm is meant to be a simple approximation to the HDP method we outlined above. First, we cluster all of the feature vectors for all songs into K groups using the k-means algorithm, renormalizing the data so that all dimensions have unit standard deviation. This defines a codebook of K cluster centers that identifies every feature vector with the cluster center to which it is closest in Euclidean space. For each song j, we compute the vector $\pi_{j,1\ldots K}$ of the relative frequencies of each cluster label. Each $\pi_{j,1\ldots K}$ defines a multinomial distribution over clusters, and we compute the distance between songs as the symmetrized KL divergence between these multinomial distributions (smoothed by a factor of 10^{-5} to prevent numerical issues).

This algorithm, like our HDP-based method, represents each song as a multinomial distribution over latent cluster identities discovered using an unsupervised algorithm, and lets us see how a much simpler algorithm that uses similar ideas performs compared with the HDP.

3.3.4 HDP

We train an HDP on all of the data using the direct assignment method [8], inferring the posterior distributions over the π_j's for each song j and computing the distance between two songs i and j as the KL divergence between the posteriors over π_i and π_j. We place vague gamma priors on α and γ [8]:

$$\alpha \sim \text{gamma}(1, 0.1), \quad \gamma \sim \text{gamma}(1, 0.1) \qquad (2)$$

and learn them during inference. For the prior H over ϕ, we use the normal-inverse-Wishart distribution [12] with parameters $\kappa_0 = 2$, $\nu_0 = 41$ (the number of dimensions plus two), and $\mu_0 = \bar{y}$ (the mean of all feature vectors across songs). The normal-inverse-Wishart matrix parameter Λ_0 was chosen by averaging the covariance matrices from 100 clusters of feature vectors, each of which was obtained by choosing a feature vector at random and choosing the 24,200 feature vectors closest to it under a Euclidean distance metric. (The number 24,200 was chosen because it was $1/5$ of the total number of points.) The goal of this process is to choose a matrix Λ_0 that resembles the covariance matrix of fairly large cluster of points, encouraging the model to find similarly shaped clusters. Using smaller (larger) clusters to choose Λ_0 would result in the model creating more (fewer) latent topics to explain the data.

3.4 Experiments

Since human-labeled ground truth similarity data is inherently expensive and difficult to acquire, we follow previous researchers [1, 2] in using genre as a proxy for similarity. We assume that all songs labeled with the same genre are "similar," which allows us to use evaluation metrics from the information retrieval literature. We first compute a full 121x121 distance matrix between all songs using each algorithm. For each query song s_q, each other song s_i is

G1	G5	G10	G20	G30	VQ5	VQ10	VQ30	VQ50	VQ100	HDP
13.24	829	1487	2786	4072	0.58	0.59	0.63	0.686	0.85	**0.25**

Table 1. Time in seconds required to compute a 121x121 distance matrix for G1, GMM-based ($K = 5, 10, 20, 30$), VQ-based ($K = 5, 10, 30, 50, 100$), and HDP-based algorithms.

	G1	G5	G10	G20	G30	VQ5	VQ10	VQ30	VQ50	VQ100	HDP
RP	0.3254	0.3190	0.3287	0.3144	0.3146	0.2659	0.2997	0.3191	0.340	0.3313	**0.3495**
AP	0.3850	0.3761	0.3746	0.3721	0.3706	0.3171	0.3546	0.3850	0.3989	0.3910	**0.3995**
AUC	0.6723	0.6712	0.6687	0.6679	0.6661	0.6513	0.6675	0.6846	0.6893	0.6758	**0.7002**

Table 2. Three measures of retrieval quality: mean R-Precision (RP), mean Average Precision (AP), and mean Area Under ROC Curve (AUC) for G1, GMM-based ($K = 5, 10, 20, 30$), VQ-based ($K = 5, 10, 30, 50, 100$), and HDP-based algorithms on the large SXSW dataset.

	G1	HDP
RP	0.5486	**0.6000**
AP	0.6807	**0.7154**
AUC	0.8419	**0.8983**

Table 3. Mean R-Precision (RP), mean Average Precision (AP), and mean Area Under ROC Curve (AUC) for G1 and our HDP-based algorithm on the smaller dataset.

given a rank $r_{q,i}$ based on its similarity to s_q. The quality of this ranking, i.e. how well it does at ranking songs of the same genre as s_q more similar than songs of different genres, is summarized using R-Precision (RP), Average Precision (AP), and the Area Under the ROC Curve (AUC), which are standard metrics from the information retrieval literature [13]. All experiments were conducted on a MacBook Pro with a 2.0 GHz Intel Core Duo processor and 2 GB of RAM. All models were implemented in MATLAB.

3.4.1 Testing on Additional Data

To test our HDP-based method's ability to generalize to unseen data using the method in section 2.4, we use the HDP trained on the large SXSW set to compute a similarity matrix on a smaller set consisting of 5 artist-filtered songs per genre (35 in all) by artists not in the training set. The electronic, punk, rap, and rock songs came from the SXSW artist showcase collection, and the country, jazz, and metal songs came from a dataset previously used by George Tzanetakis [14]. We also compute a similarity matrix on this dataset using G1, and compare the RP, AP, and AUC metrics for retrieval quality obtained using both algorithms.

4 RESULTS

Tables 1, 2, and 3 summarize the results of our experiments. The best results in each row are in bold.

The amount of time required to compute the distance matrices for the GMMs was, as expected, enormous by comparison to the other models. The cost of computing the KL divergence for the VQ-based and HDP-based models was more than an order of magnitude lower even than the cost of computing the KL divergence between single Gaussians.

The HDP performed better than the other models for all three standard information retrieval metrics, although the VQ model with $K = 50$ was a very close second. None of the GMMs outperformed G1.

The results in table 3 show that the HDP-based approach does generalize well to new songs, showing that the algorithm can be scaled up efficiently to databases of many songs.

4.1 SIMILARITY HUBS

The G1 and GK approaches are known to produce "hubs" [1] – an undesirable phenomenon where certain songs are found to be similar to many other songs. The hub phenomenon is a potentially serious concern, since it can result in very bad matches being selected as similar to a query song.

Our HDP-based approach does not suffer from this problem. Figure 3 shows how often each song is ranked in the top five of another song's similarity list for similarity matrices obtained from G1, the HDP, and choosing distances at random. The randomly generated histogram shows the sort of distribution of hubs one would expect to see due to chance in a dataset of this size. The HDP's histogram closely resembles the random one, indicating an absence of abnormal hubs. G1's histogram, by contrast, shows more severe and more numerous hubs than the other two histograms.

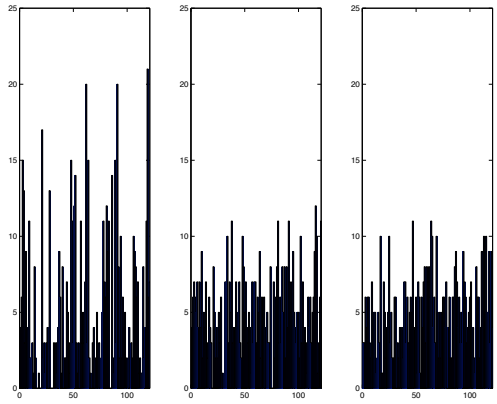

Figure 3. Histograms of how often each song is ranked in the top five of another song's similarity list for similarity matrices obtained using G1 (left), the HDP (center), and by choosing distances at random (right).

5 CONCLUSION

We developed a new method for assessing the similarity between songs. Our HDP-based approach outperformed the G1 algorithm, can compute large distance matrices efficiently, and does not suffer from the "hub" problem where some songs are found to be similar to all other songs. Since our approach does not have access to any information about temporal structure beyond that provided by the MFCC deltas (about 69 ms in total), we expect that combining the distances it provides with fluctuation patterns or some similar feature set would provide an improvement in similarity performance, as it does for the G1C algorithm [2].

One area of future work involves relaxing the bag-of-feature-vectors assumption. For example, we might learn distributions over texture patches of feature vectors instead of individual feature vectors. Hidden Markov models can also be fit into the HDP framework [8], and may yield improved results.

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HIT SONG SCIENCE IS NOT YET A SCIENCE

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ABSTRACT

We describe a large-scale experiment aiming at validating the hypothesis that the popularity of music titles can be predicted from global acoustic or human features. We use a 32.000 title database with 632 manually-entered labels per title including 3 related to the popularity of the title. Our experiment uses two audio feature sets, as well as the set of all the manually-entered labels but the popularity ones. The experiment shows that some subjective labels may indeed be reasonably well-learned by these techniques, but not popularity. This contradicts recent and sustained claims made in the MIR community and in the media about the existence of "Hit Song Science".

1. INTRODUCTION

Claims have recently been formulated about the possibility of a "Hit Science" that aims at predicting whether a given cultural item, e.g. a song or a movie, will be a hit, prior to its distribution. Such claims have been made in the domains of music [4] as well as movie [7], and are the basis of hit counseling businesses [9], [17].

More precisely, the claim is that cultural items would have specific, technical features that make them preferred by a majority of people, explaining the non uniform distribution of preferences [6]. These features could be extracted by algorithms to entirely automate the prediction process from a given, arbitrary new item (a song or a movie scenario).

A study showed the inherent unpredictability of cultural markets [19]. The unpredictability was shown to stem from a cumulative advantage or rich-get-richer effect. The study did not conclude, however, that there was no objective substrate to user preferences, but demonstrated the existence of a preference bias introduced when users are exposed to judgments of their pairs.

This study assesses to which extent this claim is scientifically grounded in the domain of music, i.e. can we extract automatically features accounting for song popularity, regardless of the cultural biases evidenced by [19].

In particular, [4] describe an experiment in which a system is trained to learn a mapping between various musical features extracted from the acoustic signal and from the lyrics, and the popularity of the song. They conclude from this experiment that their system learns something about popularity, and so that Hit Song Science is indeed possible.

However, the idea that popularity can be inferred from such technical features contradicts the natural intuitions of any musically-trained composer.

In this paper, we describe a larger-scale and more complete experiment designed to further validate this claim. We use a 32.000 song database of popular music titles, associated with fine-grained human metadata, in the spirit of the Pandora effort [16]. To ensure that the experiment is not biased, we use three sets of different features. We describe the various experiments conducted and conclude that popularity is basically not learned by any of these feature sets.

2. EXTRACTING GLOBAL DESCRIPTORS

The most widely used approach to extract global information from acoustic signals is to identify feature sets supposed to be representative of musical information contained in the signal, and to train classifiers such as SVMs (Support Vector Machines) on manually annotated data (*Train* set). These classifiers are then tested, typically on other data sets (the *Test* set), and their performance is evaluated. If the experiment is performed without biases, a good performance of the classifier means that the feature set considered does carry some information pertaining to the classification problem at hand.

In this paper we describe an experiment similar in sprit to that of [4] on a 32,000 song database. We use three different feature sets to train our classifiers: a *generic* acoustic set *à la* MPEG-7, a *specific* acoustic set using proprietary algorithms, and a set of high-level metadata produced by humans. These feature sets are described in the next sections.

2.1. Generic Audio Features

The first feature set we consider is related to the so-called *bag-of-frame* (BOF) approach. The BOF approach owns

his success to its simplicity and generality, as it can be, and has been, used for virtually all possible global descriptor problems. The BOF approach consists in modelling the audio signal as the statistical distribution of audio features computed on individual, short segments. Technically, the signal is segmented into successive, possibly overlapping frames, from which a feature vector is computed. The features are then aggregated together using various statistical methods, varying from computing the means/variance of the features across all frames to more complex modelling such as Gaussian Mixture Models (GMMs). In a supervised classification context, these aggregated features are used to train a classifier. The BOF approach can be parameterized in many ways: frame length and overlap, choice of features and feature vector dimension, choice of statistical reduction methods (statistical moments or Gaussian Mixture Models), and choice of the classifier (Decision Trees, Support Vector Machines, GMM classifiers, etc.). Many papers in the MIR literature report experiments with variations on BOF parameters on varied audio classification problems [1], [5], [12], [15]. Although perfect results are rarely reported, these works demonstrate that the BOF approach is relevant for extracting a wide range of global music descriptors.

The *generic* feature set we consider here consists of 49 audio features taken mostly from the MPEG-7 audio standard [11]. This set includes spectral characteristics (Spectral Centroid, Kurtosis and Skewness, HFC, Mel Frequency Cepstrum Coefficients), temporal (ZCR, Inter-Quartile-Range), and harmonic (Chroma). These features are intentionally chosen for their generality, i.e. they do not contain specific musical information nor musically *ad hoc* algorithms.

Various experiments [14] were performed to yield the optimal BOF parameters for this feature set: localization and duration of the signal, statistical aggregation operators used to reduce dimensionality, frame size and overlap. The best trade-off between accuracy and computation time is achieved with the following parameters: 2048 sample frames (at 44,100 Hz) with a 50% overlap computed on a 2-minute signal extracted from the middle part of the title, the features are the two first statistical moments of this distribution, i.e. the mean and variance, are considered, yielding a total feature vector of dimension 98 (49 means + 49 variances).

2.2. Specific Audio Features

The *specific* approach consists in training the same (SVM) classifier with a set of "black-box" acoustic features developed especially for popular music analysis tasks by Sony Corporation. These proprietary features have been used in commercial applications such as hard disk based Hi-Fi systems. Altogether, the specific feature set also yields a feature vector of dimension 98, which guaranties a fair comparison with the generic feature set. As opposed to the generic set, the specific set does not use the *BOF* approach: each feature is computed on the whole signal, possibly integrating specific musical information. For instance, one feature describes the proportion of perfect cadences (i.e. resolutions in the main tonality) in the whole title. Another one represents the proportion of percussive sounds to harmonic sounds. We cannot provide here a detailed description of these features as we are mostly interested in comparing the performances of acoustic classifiers on two reasonable, but different feature sets.

2.3. Human Features

Lastly, we trained a classifier with human-generated features. We use the 632 Boolean labels provided by our manually annotated database (see following section) to train the classifiers. This is not directly comparable to the 98 audio features as these labels are Boolean (and not float values). However, as we will see, these features are good candidate for carrying high-level and precise musical information that are typically not well learnt from features extracted from the acoustic signal.

3. THE HIFIND DATABASE

3.1. A Controlled Categorization Process

Several databases of annotated music have been proposed in the MIR community, such as the RWC database [8], the various databases created for the MIREX tests [3]. However, none of them has the scale and number of labels needed to test our hypothesis. For this study we have used a music and metadata database provided by the HiFind Company [10]. This database is a part of an effort to create and maintain a large repository of fine-grained musical metadata to be used in various music distribution systems, such as playlist generation, recommendation, advanced music browsing, etc. The HiFind labels are binary (0/1 valued) for each song. They are grouped in 16 categories, representing a specific dimension of music: *Style*, *Genre*, *Musical setup*, *Main instruments*, *Variant*, *Dynamics*, *Tempo*, *Era/Epoch*, *Metric*, *Country*, *Situation*, *Mood*, *Character*, *Language*, *Rhythm* and *Popularity*. Labels describe a large range of musical information: objective information such as the "presence of acoustic guitar", or the "tempo range" of the song, as well as more subjective characteristics such as "style", "character" or "mood" of the song. The *Popularity* category contains three (Boolean) labels, *low*, *medium* and *high*. It represents the popularity of the title, as observed e.g. from hit charts and records of music history. These three labels are, in principle, mutually exclusive.

The categorization process at work at HiFind is highly controlled. Each title is listened to entirely by one

categorizer. Labels to be set to *true* are selected using an *ad'hoc* categorization software. Label categories are considered in some specific order. Within a category, some rules may apply that prevent some combinations of labels to be selected. The time taken, for a trained categorizer, to categorize a single title is about 6 minutes. The categorized titles are then considered by a categorization supervisor, who checks, among other things, aspects such as consistency and coherence to ensure that the description ontologies are well-understood and utilized consistently across the categorization team. Although errors and inconsistencies can be made during this process, it nevertheless guaranties a relative good "quality" and consistency of the metadata, as opposed for instance to collaborative tagging approaches in which there is no supervision. Additionally the metadata produced is extremely precise (up to 948 labels can be considered per title), a precision which is difficult to achieve with collaborative tagging approaches.

There is no systematic way to ensure that the categorization produces absolutely correct and consistent information, so we had to consider the database as it was provided as ground truth. Some minor "clean up" was performed before use, by discarding titles with metadata of obviously of low quality. For instance, we discarded songs having much less labels set to "true" than the average (37). Additionally, we kept only those labels for which we had a significant amount of titles (above 20) with the *true* and *false* values, to build training and testing sets of sufficient size. As a result of this cleanup, the total number of titles considered in this study is 32978, and the number of labels 632. (Note that those labels correspond to the 632 human features for the experiment described in Section 2.3) Acoustic signals were given in the form of a *wma* file at 128 kbps. This database was used both for training our classifiers and for testing them, as described in Section 4.1.

3.2. Database Redundancy

The HiFind database is *sparse*: the mean number of labels set to true per song (occupation factor) is 5.8% (i.e. 37 on a total of 632). Sparseness suggests the dominant role of the true-valued labels compared to false-valued labels for a given song. It is also *redundant*. For instance, labels 'Country Greece' and 'Language Greek' are well correlated. This redundancy has been analyzed and exploited for performing statistical inference, e.g. to infer unknown attributes from a partial description of a music title, or for suggesting modifications [18].

3.3. Assessing Classifiers

To avoid the problems inherent to the sole use of precision or recall, the traditional approach is to use *F-Measure* to assess the performance of classifiers. For a given label, the *recall R* is the proportion of positive examples (i.e. the titles that are *true* for this label) that were correctly predicted. The *precision P* is the proportion of the predicted positive examples that were correct. When the proportion of positive examples is high compared to that of negative examples, the precision will usually be artificially very high and the recall very low, regardless of the actual quality of the classifier. The F-measure addresses this issue and is defined as:

$$F = 2 \times {R \times P}/{R + P}$$

However, in our case, we have to cope with a particularly unbalanced 2-class (*True* and *False*) database. So the mean value of the F-measure for each class (*True* and *False*) can still be artificially good. To avoid this bias, we assess the performance of our classifiers with the more demanding *min F-measure*, defined as the minimum value of the F-measure for the positive and negative cases. A *min-F-measure* near 1 for a given label really means that the two classes (*True* and *False*) are well predicted.

4. EXPERIMENT

4.1. Experiment Design

We first split the *HiFind* database in two "balanced" parts *Train* and *Test*, so that *Train* contains approximately the same proportion of examples and counter-examples for each labels as *Test*. We obtained this state by performing repeated random splits until a balanced partition was observed. We trained three classifiers, one for each feature set (*generic*, *specific* and *human*). These classifiers all used a Support Vector Machine (SVM) algorithm with a Radial-Basis Function (RBF) kernel, and were trained and tested using *Train* and *Test*. More precisely, each classifier, for a given label, is trained on a maximally "balanced" subset of *Train*, i.e. the largest subset of *Train* with the same number of "True" and "False" titles for this label (popularity *Low*, *Medium* and *High*). In practice the size of these individual train databases varies from 20 to 16320. This train database size somehow represents the "grounding" of the corresponding label. The classifiers are then tested on the whole *Test* base. Note that the *Test* base is usually not balanced with regards to a particular label, which justifies the use of the *min-F-measure* to assess the performance of each classifier.

4.2. Random Oracles

To assess the performance of our classifiers, we compare them to that of *random oracles* defined as follows: given an label with *p* positive examples (and therefore *N-p* negative ones, with *N* the size of the test set), this oracle returns *true* with a probability *p/N*.

By definition, the min-F-measure of a random oracle only depends on the proportion of positive and negative examples in the test database.

For instance, for a label with balanced positive and negative examples, the random oracle defined as above has a *min-F-measure* of 50%. A label with 200 positive examples (and therefore around 16,000 negative examples) leads to a random oracle with a min-F-measure of 2.3%. So the performance of the random oracle is a good indicator of the size of the train set, and can therefore be used for comparing classifiers as we will see below.

4.3. Evaluation of the Performance of Acoustic Classifiers

4.3.1. Comparison with random oracles

The comparison of the performance of acoustic classifiers with random oracles shows that the classifiers do indeed learn something about many of the HiFind labels. More than 450, out of 632, are better learned with the acoustic classifiers than with our random oracle. Table 1 indicates, for each feature set, the distribution of the relative performances of acoustic classifiers with regards to random oracles.

Improvement	Specific	Generic
50	8	0
40	12	15
30	43	20
20	111	79
10	330	360
0	128	158

Table 1. Number of labels for which an acoustic classifier improves over a random classifier by a certain amount. Column "Improvement" reads as follows: there are 111 labels for which a specific acoustic classifier outperforms a random classifier by +20 (in min-F-measure).

Table 1 also shows that around 130 to 150 labels lead to low-performance classifiers, i.e. acoustic classifiers that do not perform significantly better than a random oracle (the last row of the table); approximately half of the labels lead to classifiers that improve over the performance of a random classifier by less than 10; the rest (top rows) clearly outperform a random oracle, i.e. they are well-modeled by acoustic classifiers.

4.3.2. Distribution of performances for acoustic classifiers

At this point, it is interesting to look at the distribution of the performances of these acoustic classifiers. These performances vary from 0% for both feature sets to 74% for the generic features and 76% for the specific ones. The statistical distribution of the performances is close to a power law distribution, as illustrated by the log-log graph of **Figure 1**.

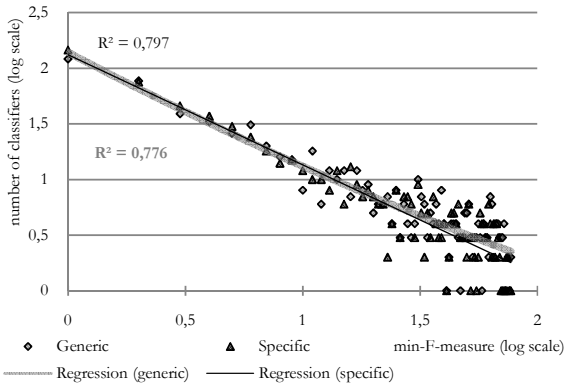

Figure 1. Log-log graph of the distribution of the performance of acoustic classifiers for both feature sets. This distribution of the performance of classifiers is close to a power law.

These power laws suggest that a natural organization process is taking place in the representation of human musical knowledge, and that the process of automatic audio classification maintains this organization.

4.3.3. Specific features slightly outperform generic features

Not surprisingly, we can see that specific features perform always better than the generic ones. This is illustrated by **Figure 2**. Since the classifiers are both based on the same SVM/kernel, the difference can only come from 1) the actual features extracted or 2) the aggregation method. For the generic features, the aggregation is based on means and averages over all the segments of the song. For the specific features, the aggregation is *ad hoc*.

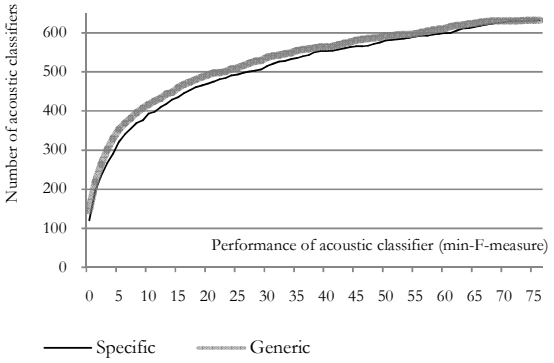

Figure 2. Cumulated distribution of the performance of acoustic classifiers for the generic and specific feature sets. There are more classifiers with low performance for the generic feature sets (leftmost side of the graph).

4.3.4. Acoustic classifiers perform better for large training sets

Lastly, we can observe the relationship between the performance and the size of the training set. The trend lines in Figure 3 show that the performances of acoustic classifiers increase with the training dataset size, regardless of the feature set. This is consistent with the acknowledged fact that machine-learning algorithms require large numbers of training samples, especially for high-dimensional feature sets.

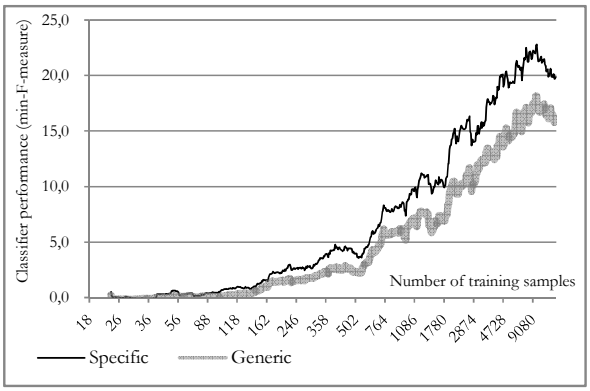

Figure 3. The relative performances of the 632 acoustic classifiers (i.e. the difference between the min-F-measures of the classifier and of the corresponding random oracle) for specific and generic features, as a function of the training database size. The performance of the acoustic classifiers increases with the size of the training database.

These experiments show that acoustic classifiers definitely learn some musical information, with varying degrees of performance. It also shows that the subjective nature of the label do not seem to influence their capacity to be learned by audio features. For instance, the label "Mood nostalgic" is learnt with the performances of 48% (specific features), and 43% (generic features), to be compared to the 6% of the random oracle. Similarly, label "Situation evening mood" is learnt with 62% and 56% respectively, against 36% for random. So popularity is, *a priori*, a possible candidate for this task.

4.4. Inference from Human Data

This double feature experiment is complemented by another experiment in which we train a classifier using all the HiFind labels but the Popularity ones. This is justified by the low entropy of the database as discussed in Section 3.2. Contrarily to the acoustic classifiers, we do not present here the performances of the classifiers for all HiFind labels. Indeed, some pairs of HiFind labels are perfectly well correlated so this scheme works perfectly for those, but this result is not necessarily meaningful (e.g. to infer the country from the language). The same *Train / Test* procedure described above applied with the 629 non-popularity labels as input yields the following result (*min-F-measure*): 41% (*Popularity-Low*), 37% (*Popularity-Medium*) and 3% (*Popularity-High*).

4.5. Summary of Results for Inferring Popularity

The results concerning the Popularity labels are summarized in Table 2.

Popularity label	Generic features	Specific features	Corrected specific	Human features	Dumb features	Random oracle
Low	36	35	31	41	32	27
Medium	36	34	38	37	28	22
High	4	3	3	3	3	3

Table 2. The performances (min-F-measures) of the various classifiers for the three Popularity labels. No significant improvement on the random oracle is observed.

These results show clearly that the Popularity category is not well-modeled by acoustic classifiers: its mean performance is ranked fourth out of 16 categories considered, but with the second lowest maximum value among categories.

Although these performances appear to be not so bad at least for the "Low" label, the comparison with the associated random classifiers shows that popularity is in fact practically not learnt. Incidentally, these performances are not improved with the *correction scheme*, a method that exploits inter-relations between labels to correct the results [14], in the spirit of the contextual approach described in [2].

Interestingly, the use of human features (all HiFind labels) does not show either any significant performance.

Lastly, we also considered *a priori* irrelevant information to train our classifiers: the letters of the song title, i.e. a feature vector of size 26, containing the number of occurrences of each letter in the song title. The performances of the corresponding classifiers are respectively 32% 28% and 3% (for the low, medium and high popularity labels, see Table 2). This shows that even dumb classifiers can slightly improve the performances of random classifiers (by 5% in this case for the medium and low popularity labels), but that this information does not teach us anything about the nature of hits.

These results suggest that there are no significant statistical patterns concerning popularity using these features sets.

5. CONCLUSION

We have shown that the popularity of a song cannot be learnt by using state-of-the-art machine learning

techniques with two sets of reasonable audio features. This result is confirmed when using supposedly higher-level human metadata. This large-scale evaluation, using the best machine-learning techniques available to our knowledge, contradicts the claims of "Hit Song Science", i.e. that the popularity of a music title can be learned effectively from known features of music titles, either acoustic or human. We think that these claims are either based on spurious data or on biased experiments. This experiment is all the more convincing that some other subjective labels can indeed be learnt reasonably well using the features sets described here (e.g. the "mood nostalgic" label).

This experiment does not mean, however, that popularity cannot be learnt from the analysis of a music signal or from other features. It rather suggests that the features used commonly for music analysis are not informative enough to grasp anything related to such subjective aesthetic judgments. Current works are in progress to determine "good features" using feature generation techniques [13], which have been shown to outperform manually designed features for specific analysis tasks. However, more work remains to be done to understand the features of subjectivity for even simpler musical objects such as sounds or monophonic melodies. Hit song science is not yet a science, but a wide open field.

6. ACKNOWLEDGEMENT

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MELODY EXPECTATION METHOD BASED ON GTTM AND TPS

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ABSTRACT

A method that predicts the next notes is described for assisting musical novices to play improvisations. Melody prediction is one of the most difficult problems in musical information retrieval because composers and players may or may not create melodies that conform to our expectation. The development of a melody expectation method is thus important for building a system that supports musical novices because melody expectation is one of the most basic skills for a musician. Unlike most previous prediction methods, which use statistical learning, our method evaluates the appropriateness of each candidate note from the view point of musical theory. In particular, it uses the concept of melody stability based on the generative theory of tonal music (GTTM) and the tonal pitch space (TPS) to evaluate the appropriateness of the melody. It can thus predict the candidate next notes not only from the surface structure of the melody but also from the deeper structure of the melody acquired by GTTM and TPS analysis. Experimental results showed that the method can evaluate the appropriateness of the melody sufficiently well.

1. INTRODUCTION

We have developed a method for predicting the next notes in a melody that uses the generative theory of tonal music (GTTM) [1] and the tonal pitch space (TPS) [2]. Our melody prediction method helps a novice construct a melody or play an improvisation by displaying candidates for the next notes. This method is designed to be used with a "prediction piano" (Figure 1), which was developed to assist musical novices play improvisations. On the lid of this piano, there is a 32 × 25 full-color LED matrix that displays a piano roll view that scrolls down in time with the music.

We identified two key requirements for our melody expectation method to make it useful to musical novices playing an improvisation on the expectation piano.

1) Candidate notes are predicted and output even if the input melody is novel.
2) The output is appropriate from a musical point of view.

Two approaches were considered when developing this method: statistical learning and music theory. With the statistical learning approach, the predictions depend on the characteristics of the data used for learning: composer, genre, period, country, etc. [3, 4]. Moreover, predicting candidate notes for a novel melody is problematic because the system may not be able to find a similar melody in the learning data and thus may be unable to evaluate whether the notes are appropriate or not. With the music theory approach, the predictions do not depend on the characteristics of the data used for learning. It can thus be applied to novel melodies.

Although many music theories have been proposed [5–8], GTTM is the most suitable for predicting notes in a melody because it can be used to represent the various aspects of music in a single framework. Furthermore, it includes a concept of stability, meaning that it can be used to evaluate the appropriateness of the predicted notes.

This paper is organized as follows. Section 2 briefly explains GTTM and the analyzers we constructed. Section 3 explains our melody prediction method. In Section 4, we describe the expectation piano, and, in Section 5, we present some experimental results. We conclude in Section 6 with a summary of the main points and mention of future work.

Figure 1. Expectation Piano.

2. GTTM AND ANALYZERS

The GTTM is composed of four modules, each of which assigns a separate structural description to a listener's understanding of a piece of music. These four modules output a grouping structure, a metrical structure, a time-span tree, and a prolongational tree, respectively (Figure 2). The time-span tree is a binary tree with a hierarchical structure that describes the relative structural importance of the notes that differentiate the essential parts of the melody from the ornamentation.

There are two types of rules in GTTM, i.e., "wellformedness rules" and "preference rules". Wellformedness rules are the necessary conditions for assigning structures and restrictions on the structures. When more than one structure satisfies the well-formedness rules, the preference rules indicate the superiority of one structure over another.

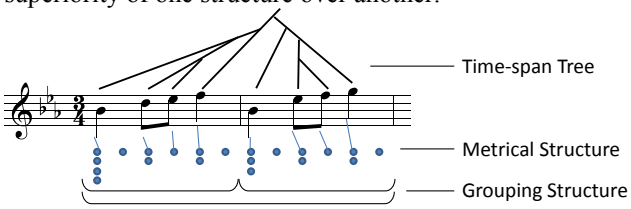

Figure 2. Time-span tree, metrical structure, and grouping structure.

2.1. Problems in implementing GTTM

In this section, we specify the problems with the GTTM rules in terms of computer implementation.

2.1.1. Ambiguous concepts defining preference rules

GTTM uses some undefined words that can cause ambiguities in the analysis. For example, GTTM has rules for selecting structures for discovering similar melodies (called parallelism), but does not have the definition of similarity itself. To solve this problem, we attempted to formalize the criteria for deciding whether each rule is applicable or not.

2.1.2. Conflict between preference rules

Conflict between rules often occurs and results in ambiguities in the analysis because there is no strict order for applying the preference rules. Figure 3 shows a simple example of a conflict between grouping preference rules (GPR). GPR3a (register) is applied between notes 3 and 4 and GPR6 (parallelism) is applied between notes 4 and 5. A boundary cannot be perceived at both 3-4 and 4-5, because GPR1 (alternative form) strongly prefers that note 4, by itself, cannot form a group.

Figure 3. Simple example of conflict between rules.

To solve this problem, we introduced adjustable parameters that enable us to control the strength of each rule.

2.1.3. Lack of working algorithm

Knowledge represented in the rule form is in general declarative, which is advantageous in the sense that a knowledge programmer does not need to take into account an algorithm for reasoning. A system is required to perform automatic reasoning on the declaratively described knowledge.

Unfortunately, GTTM has few descriptions of the reasoning and working algorithms needed to compute analysis results.

2.2. exGTTM

In our previous work [9–11], we extended the GTTM theory through full externalization and parameterization and devised a machine-executable extension of GTTM, exGTTM. The externalization includes introducing an algorithm for generating a hierarchical structure of the time-span tree in a mixed top-down/bottom-up manner. Such an algorithm has not previously been represented for GTTM. The parameterization includes a parameter for controlling the priorities of rules to avoid conflicts among them as well as parameters for controlling the shape of the hierarchical time-span tree. Although it has been suggested that such parameters are required in GTTM, they were not explicitly presented.

Here, we distinguish two kinds of ambiguity in music analysis: one involves the musical understanding by humans, and the other concerns the representation of a music theory. The former kind of ambiguity derives from the ambiguity of the music itself. For the latter type of ambiguity, related to GTTM, either no concept for mechanization has been presented, or it has only been presented in an implicit way. Therefore, due to the former kind of ambiguity, we assume there is more than one correct result. We avoid the latter kind of ambiguity as much as possible by performing full externalization and parameterization.

2.2.1. Full externalization and parameterization

The significance of full externalization and parameterization is twofold: precise controllability and coverage of the manual results. Whenever we find a correct result that exGTTM cannot generate, we introduce new parameters and give them appropriate values so that it can generate the correct result. In this way, we repeatedly externalize and introduce new parameters until we can obtain all of the results that people consider correct. In total, we introduced 15 parameters for grouping-structure analysis, 18 for metrical-structure analysis, and 13 for

	Parameters	Description
Grouping structure	$S_{GPR\,R}$ ($0 \leqq S_{GPR\,R} \leqq 1$)	Strength of each grouping preference rule. The larger the value is, the stronger the rule acts. $R \in$ { 2a, 2b, 3a, 3b, 3c, 3d, 4, 5, and 6 }
	σ ($0 \leqq \sigma \leqq 0.1$)	Standard deviation of a Gaussian distribution, the average of which is the boundary by GPR5. The larger the value is, the wider its skirt becomes.
	W_m ($0 \leqq W_m \leqq 1$)	Balance between temporal similarity of attack points and that of pitch difference in GPR6. The larger the value is, the more the system estimates the pitch difference.
	W_l ($0 \leqq W_l \leqq 1$)	Weight for the length of parallel phrases. The larger the values is, the more the length of parallel phrases is prioritized in GPR6.
	W_s ($0 \leqq W_s \leqq 1$)	Balance determining whether the note *i* becomes the ending note of a group or the beginning note of the following group in GPR6. The larger the value is, the more the note tends to be the ending note.
	T_{GPR4} ($0 \leqq T_{GPR4} \leqq 1$)	Threshold at which the effects of GPR2,3 are considered to be salient in GPR4. The smaller the value is, the more probably GPR4 is applied.
	T^{low} ($0 \leqq T^{low} \leqq 1$)	Threshold in the lower-level boundary. The smaller the value is, the more salient the boundary becomes.
Metrical structure	$S_{MPR\,R}$($0 \leqq S_{MPR\,R} \leqq 1$)	Strength of each metrical preference rule. The larger the value is, the stronger the rule acts. $R \in$ {1,2,3,4,5a, 5b, 5c, 5d, 5e, and 10)
	W_m ($0 \leqq W_m \leqq 1$)	Balance between temporal similarity of attack points and that of pitch difference in MPR1. The larger the value is, the more the system estimates the pitch difference.
	W_l ($0 \leqq W_l \leqq 1$)	Weight for the length of parallel phrases. The larger the value is, the more the length of parallel phrases is prioritized in MPR1.
	W_s ($0 \leqq W_s \leqq 1$)	Balance determining whether the note *i* becomes the ending note of a group or the beginning note of the following group in MPR1. The larger the value is, the more the note tends to be the ending note.
	$T_{MPR\,R}$($0 \leqq T_{MPR\,R} \leqq 1$)	Value of the threshold that decides whether each rule is applicable. $R \in$ { 4, 5a, 5b, 5c }
Time-span tree	$S_{TSRPR\,R}$($0 \leqq S_{TSRPR\,R} \leqq 1$)	Strength of each rule. The larger the value is, the stronger the rule acts. $R \in$ {1, 2, 3, 4, 5a, 5b,5c, 5d, 5e, and 10}
	W_m ($0 \leqq W_m \leqq 1$)	The balance between the temporal similarity of attack points and that of the pitch difference in TSRPR4. The larger the value is, the more the system estimates the pitch difference.
	W_l ($0 \leqq W_l \leqq 1$)	The weight for the length of parallel phrases. The larger the values is, the more the length of parallel phrases is estimated in TSRPR4.
	W_s ($0 \leqq W_s \leqq 1$)	The balance determines whether the note i becomes the ending note of a group or the beginning note of the following group in TSRPR4. The larger the value is, the more the note tends to be the ending note.

Table 1. Adjustable parameters of the exGTTM and ATTA.

time-span reduction(Table 1).

We appropriately supply lacking parameters and make implicit parameters explicit[1]. The parameters introduced by exGTTM are categorized into identified, implied, and unaware.

A parameter in the first category is identified in GTTM but it is not assigned concrete values. Hence, we valuate such a parameter. For example, since the resulting value of the GPR2a application, D_{GPR2a}, is binary, if the rule holds, D_{GPR2a} makes 1, and 0 if it does not hold. On the other hand, since GPR6 holds indefinitely, the resulting value of GPR6, D_{GPR6}, varies continuously between 0 and 1.

A parameter of the second category is implied in GTTM. Hence, we make it explicit. For example, to resolve the preference rule conflict, we introduce parameters to express the priority for each preference rule ($S^{\mathrm{GPR\,R}}$, $S^{\mathrm{MPR\,R}}$, and $S^{\mathrm{TSRPR\,R}}$ in Table 1). Since each preference rule has its own priority, all of the priority patterns are realized. This is an example of full-parameterization.

For the third category, we need to complement parameters that are not recognized in the original theory, since some of them may nearly lack any musicological meaning. For example, GPR6 in exGTTM needs to add parameters for controlling the properties of parallel segments, including the weights for pitch-oriented matching or timing-oriented matching.

We add a comment to the domain of intermediate variables, denoted as D and B. The domain of all the intermediate variables is constrained within the range of 0 to 1, and for this purpose, these variables are normalized at every computing stage. Thanks to this property, exGTTM can flexibly combine any intermediate variables (and possibly parameters) and cascade as many weighted-mean calculations as needed. This facilitates precise controllability.

2.2.2. Algorithm for acquiring hierarchy

Among the issues that require working algorithms, the problems for acquiring hierarchical structures in the grouping- and metrical-structure analyses and the time-span tree reduction can be all regarded as constraint satisfaction problems (CSP). This is because only the properties to be satisfied for the hierarchical structures are represented in the form of a rule; that is, neither constraint nor order of generating hierarchical structures is determined in advance.

[1] In the paper, the word ``parameter'' is used not only for parameters used in controlling a system externally but also for internal variables (intermediated variables) connecting submodules.

The constraints stipulated by the GTTM rules are divided into two categories: local and global. The former includes GPR2 (proximity) and TSRPR1 (strong metrical position), and the latter GPR5 (symmetry) and MPR1 (parallelism). We need to handle global constraints carefully when generating hierarchical structures. For the example of GPR5 in Figure 4, given a group at Layer 1, an inner boundary likely occurs around the center of the group, that is, either between Notes 1 and 2 or between Notes 2 and 3. Here, we can consider two cases. In Case 1, the boundary between Notes 1 and 2 is selected, taking into account the effects of some other rules. Then in each subgroup in Layer 2, the inner boundary of the subgroup may occur on the left-hand side of a center note. On the other hand, in Case 2, the boundary between Notes 2 and 3 is selected. Therefore, the inner boundary may occur on the right-hand side of a center note. Consequently, in computing GPR5, the determined boundary position influences the identifications of remote boundaries in lower layers, and we have to take into account up-to-date global information every time. That is, a global constraint is inevitably dynamic.

In light of the above considerations, we are developing algorithms for generating hierarchical structures for exGTTM so that nodes are generated either from the bottom-most nodes or the top-most node incrementally and so that every time the nodes at a layer are calculated, global information is re-calculated before moving onto an adjacent layer.

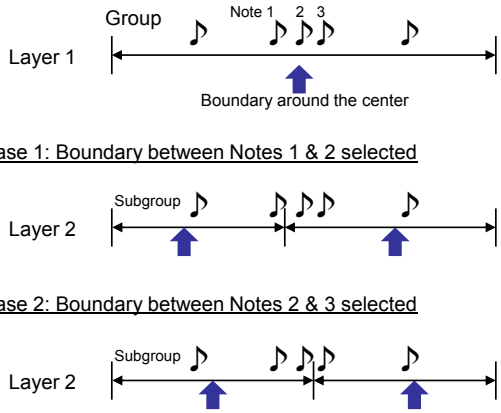

Figure 4. Simple example of conflict between rules.

2.3. FATTA: Fully Automatic Time-span Tree Analyzer

We implemented a time-span tree analyzer, called automatic time-span analyzer (ATTA), based on exGTTM. Although ATTA can automatically acquire a time-span tree, because the parameters are manually controlled, it takes too much time to find a set of optimal parameters. Therefore, we developed a method for automatically estimating the optimal parameters [12].

Two rules in GTTM [1] are not implemented in ATTA: GPR7 and TSRPR5.

> GPR7 (time-span and prolongational stability): prefer a grouping structure that results in a more stable time-span and/or prolongational reductions.
>
> TSRPR5 (metrical stability) In choosing the head of time-span *T*, prefer a choice that results in a more stable choice of metrical structure.

These rules require that information from later processes, such as time-span/prolongational reductions, be sent back to earlier processes.

To estimate the optimal parameters automatically, we evaluate the structural stability of the analysis results derived by ATTA. We use GPR7 and TSRPR5 to calculate the level of stability. Figure 5 shows the process flow of the FATTA, which consist of the ATTA and a loop by the GPR7 and TSRPR5.

2.3.1. Implementation of GPR7 with Tonal Pitch Space

GPR7 is applied to the loop between the time-span/prolongational reduction and grouping structure analysis. This rule leads to a preference for a grouping structure that results in a more stable time-span and/or prolongational reductions. The holding level of GPR7, which varies continuously between 0 and 1, is defined as

$$D_{GPR7} = \frac{\sum_i \text{distance}(p(i), s(i)) \times \text{size}(i)^2}{\sum_i \text{size}(i)^2}, \quad (1)$$

where i indicates the head of the time-span, which has primary and secondary branches, denoted by $p(i)$ and $s(i)$, respectively. Distance (x, y) indicates the distance between notes x and y in the tonality of the piece, which are defined using Lerdahl's tonal pitch space [2]. We normalized the distance from 0 to 1. The size(i) indicates the length of the time-span with head i. When calculating D_{GPR7}, we use the square of size(i) for the weighting for empirical reasons.

In the tonal pitch space, the distance between chord x = C_1/**R_1** and chord y = C_2/**R_2** is defined as follows:

$$\delta(x \to y) = i + j + k, \quad (2)$$

where i is region distance, j is chord distance, and k is basic space difference. The region distance is the smallest number of steps along the regional circle of fifth between **R_1** and **R_2**. The chord distance is the smallest number of steps along the chordal circle of fifth between the roots of C_1 and C_2 within each region. The basic space distance is a specially weighted to define each chord and region. Note that pitch class only has a meaning in terms the elements of the sets that define chords and regions, and chords are always understood as functioning within some region.

2.3.2. Implementation of TSRPR5

TSRPR5 is applied to the loop between the time-span reduction and metrical structure analyzer and results in a more stable metrical structure when choosing the head of a time-span. The holding level of TSRPR5, which varies continuously between 0 and 1, is defined as

$$D_{TSRPR5} = \frac{1}{\sum_i \text{size}(i)^2} \times \sum_i \begin{cases} \text{size}(i)^2 & dot(p(i)) \geq dot(s(i)) \\ 0 & dot(p(i)) < dot(s(i)) \end{cases}, \quad (2)$$

where $dot(x)$ indicates the number of metrical dots of note x.

2.3.3. Optimization of adjustable parameters

The set of optimal ATTA parameters is obtained by maximizing the average of D_{GPR7} ($0 \leq D_{GPR7} \leq 1$) and D_{TSRPR5} ($0 \leq D_{TSRPR5} \leq 1$). The parameters and default values are $S^{rules} = 0.5$, $T^{rules} = 0.5$, $Ws = 0.5$, $Wr = 0.5$, $Wl = 0.5$, and $\sigma = 0.05$. Because there are 46 parameters, a great amount of time is needed to calculate all parameter combinations. To minimize the calculation time, we constructed an algorithm that does the following:

(1) Maximize average of D_{GPR7} and D_{TSRPR5} by changing a parameter from its minimum to its maximum value.
(2) Repeat (1) for all parameters.
(3) Iterate (1) and (2) as long as the average of D_{GPR7} and D_{TSRPR5} is higher than after the previous iteration.

3. MELODY EXPECTATION METHOD

Our melody expectation method predicts candidate notes by using the level of stability of the time-span tree defined in FATTA. Our position is that we cannot always specify a single expected, following tone; and thus, we developed an expectation piano that simply suggests multiple candidates among stable pitch events with higher stability. The functions of FATTA are restricted as follows; FATTA only treats monophonic western tonal music. Thus, our expectation method can predict only monophonic musical structures of western tonal music.

3.1. Expectation Method based on FATTA

The main advantage of our melody expectation method is that, the stability of a melody is calculated by analyzing the whole melody from the beginning note to the expected note, not from only the local melody(a few notes previous to a relevant note) ; previous melody expectation methods based on music theory (eg. Steve Larson's theory of musical forces [8]) derive the expected note from the local melody. Music tends to be more interesting when it does not match the listener's expectations, such as a delayed note, and this may result in tension and relaxation. A composer can deliberately construct such music, which can make it difficult to predict the next notes in the melody with accuracy. For example, an

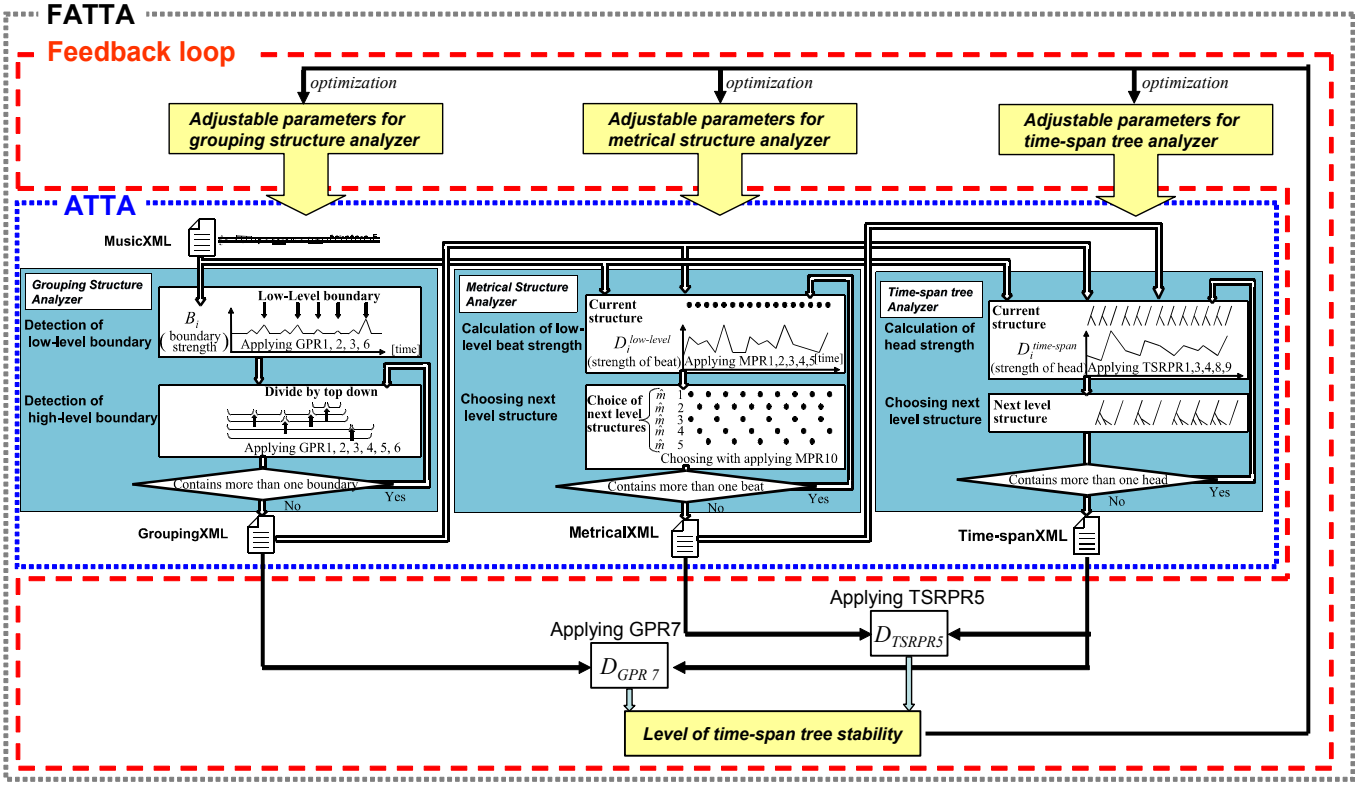

Figure 5. Processing flow of fully automatic time-span tree analyzer

ornamentation note is often inserted before the expected note. In such cases, our method can predict candidate notes fairly well because FATTA can evaluate the stability of the entire structure of the time-span trees, which includes branches connected to essential notes and leaves connected to ornament notes.

3.2. Real-time Extension for FATTA

To be able to predict notes on the basis of GTTM, FATTA must run in real time. However, FATTA needs several minutes to finish the analysis, so running in real time is difficult. Therefore, we extended the algorithm to enable real-time operation. First, to speed up the iteration described in 2.2, we use the set of optimal parameter values for the last melody, which is one note shorter than the present melody, as the initial parameter set. Second, to reduce the time used by ATTA, we introduce a window for analysis. The size of the window is the longest group length within 16 measures of the present position. This length can be acquired through preprocessing using the grouping structure analyzer in ATTA. If there is no grouping boundary in 16 measures from the present position, we use 16 measures as the window size.

3.3. Calculation Level of Stability of Melodies by FATTA

Our method evaluates the appropriateness of each candidate melody to occur after the present one by calculating its stability. We use the average of D_{GPR7} and D_{TSRPR5} as the level of stability acquired by FATTA.

Figure 6 shows the calculated level of stability from the primary note to the present note. The level of stability can be calculated after the third note because GTTM analysis needs at least four notes.

In the score, the primary note is a tonic of the region, and the tail note is also a tonic of the region. This means that the levels of stability indicate a large value at the tail of the melody and a smaller value in the middle of the melody. Therefore, the higher the level of stability, the relatively closer the tonic of the region. The region of the melody and chord progression were estimated in the implementation of GPR7 by applying the tonal pitch space.

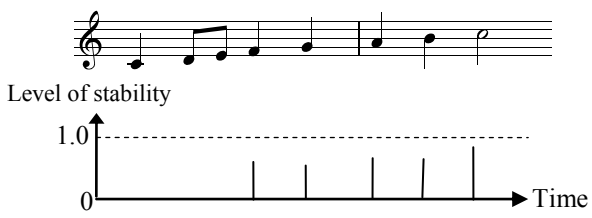

Figure 6. Level of stability over time.

4. EXPECTATION PIANO

Our expectation piano assists novices with musical improvisation by displaying the predicted notes on the piano lid. When the novice finds it hard to continue playing the melody, she/he can continue the improvisation by playing a note displayed on the lid, without impairing tonality.

4.1. Overview

The processing flow of the prediction piano is as follows (Figure 7). First, the MIDI signals for the music played on the piano are sent to a computer. The signals are quantized, and a MusicXML version of the performed melody is created. A learning-based quantization method [13] is used to eliminate the deviation in onset times, which are then aligned to the normalized positions. The predicted notes are acquired by inputting the MusicXML into FATTA. Finally, the predicted notes are displayed on the piano lid.

4.2. LED Scrolling Piano Roll

The predicted notes are displayed in piano roll format within the range of view of the keyboard. The roll scrolls down at a constant speed. Below the piano lid, which is made of semitransparent acrylic resin, there is a 32 × 25 full-color LED matrix for displaying the scrolling piano roll. The 32 represents two measures when the resolution is a sixteenth note, and 25 is the number of keys on the keyboard. The color of each LED in the matrix is determined under the assumption that the onset of the next note will start at the corresponding position on the piano roll and by calculating the level of stability. When the level of stability is high, the LEDs show yellow, when it is low, they show black, and when it is neither, they show red. There is also a 32 × 20 blue LED matrix that displays the bar lines of the piano roll.

4.3. Construction

The piano is 953 mm long and 710 mm wide and resembles a grand piano. It contains a MIDI keyboard, the LED display, a microcomputer, a power supply, and four speakers. The LED display is 630 mm long and

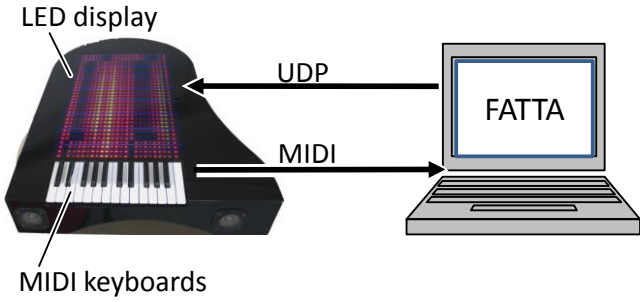

Figure 7. Overview of expectation piano.

LEARNING A METRIC FOR MUSIC SIMILARITY

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ABSTRACT

This paper describe five different principled ways to embed songs into a Euclidean metric space. In particular, we learn embeddings so that the pairwise Euclidean distance between two songs reflects semantic dissimilarity. This allows distance-based analysis, such as for example straightforward nearest-neighbor classification, to detect and potentially suggest similar songs within a collection. Each of the six approaches (baseline, whitening, LDA, NCA, LMNN and RCA) rotate and scale the raw feature space with a linear transform. We tune the parameters of these models using a song-classification task with content-based features.

1 INTRODUCTION

Measuring the similarity of two musical pieces is difficult. Most importantly, two songs that are similar to two lovers of jazz, might be very different to somebody that does not listen to jazz. It is inherently an ill-posed problem.

Still, the task is important. Listeners want to find songs that are related to a song that they like. Music programmers want to find a sequence of songs that minimizes jarring discontinuities. A system based on measurements from hundreds of thousands of users is perhaps the ultimate solution [8], but there is still a need to find new songs, before an item-to-item system has enough data.

It is difficult to construct a distance calculation based on arbitrary features. A simple approach places the feature values into a vector and then calculates an Euclidean distance between points. Such a calculation implies two things about the features: their independence and their scale. Most importantly, a Euclidean metric assumes that features are (nearly) orthogonal so the distance along different axis can be summed. A Euclidean metric also assumes that each feature is equally important. Thus a distance of 1 unit in the X direction is perceptually identical to one unit in the Y direction. This is unlikely to be true, and this paper describes principled means of finding the appropriate weighting.

Much work on music similarity and search calculates a feature vector that describes the acoustics of the song, and then computes a distance between these features. In this work we describe six means of assigning weights to the dimensions and compare their performance. The purpose of this paper is not to determine the best similarity measure—after all evaluation of a personal decision such as similarity is difficult—but instead to test and compare several quantitative approaches that MIR practitioners can use to create their own similarity metric. West's recent paper provides a good overview of the problem and describes successful approaches for music similarity [11]. We hope to improve the performance of future systems by describing techniques for embedding features in a metric space.

We measure the performance of our system by testing identification ability with a k-nearest neighbor (kNN) classifier. A kNN classifier is based on distances between the query point and labeled training examples. If our metric space is "good" then similar songs will be close together and kNN classification will produce the right identification. In our case, we try to identify the album, artist or blog associated with each song.

A kNN classifier has several advantages for our task. A kNN classifer is simple to implement, and with large amounts of data they can be shown to give an error rate that is no worse than twice the optimal recognizer [2]. Simple classifiers have often been shown to produce surprisingly good results [5]. The nearest-neighbor formulation is interesting in our application because we are more interested in finding similar songs, than we are in measuring the distance between distant songs or conventional classification. Thus kNN classification is a good metric for measuring our ability to place songs into a (linear) similarity space.

2 DATA

Our data comes from the top 1000 most-popular mp3 blogs on the Web, as defined by music blog aggregator, The Hype Machine's "TOP MUSIC BLOGS On The Net" [1]. We analyzed each new mp3 track that was posted on these mp3 blogs during the first three weeks of March 2008.

In the ongoing quest to discover new music, music blogs provide an engaging and highly useful resource. Their cre-

[1] http://hypem.com/toplist

ators are passionate about music, given that they're blogging about it all the time. And unlike a traditional playlist or set of recommended tracks, music blogs also provide personal commentary which gives the blog consumer a social context for the music.

We're interested in comparing the similarity across music posted on the same blog to the similarity between different tracks by the same artist or from the same album.

One of the difficulties in gathering this type of data is the large amount of uncertainty and noise that exists within the metadata that describes mp3s. Our general experience analyzing Web mp3s has been that less than a third of the tracks we encounter have reliable (if any) metadata in their ID3 tags. Therefore the metadata we used for our artists and album calculations is limited to what we were able to parse out of valid ID3 tags or information we could infer from the filename, or the HTML used to reference the track.

In these 1000 blogs, we found 5689 different songs from 2394 albums and 3366 artists. Just counting blogs for which we found labeled and unique songs, we were left with 586 different blogs in our dataset. After removing IDs for which we did not have enough data (less than 5 instances) we were left with 74 distinct albums, 164 different artists, and 319 blogs. The style of music represented in this collection differs from blog to blog. Many mp3 blogs could be broadly classified as "Indie" or "Indie Rock", but music shared on a specific blog is more representative of the blogger's personal taste than any particular genre.

3 FEATURES

We characterized each song using acoustic analysis provided via a public web API provided by The Echo Nest [4]. We send a song to their system, they analyze the acoustics and provide 18 features to characterize global properties of the songs. Although we did not test it, we expect that features from a system such as Marsyas [9] will give similar results.

The Echo Nest Analyze API splits the song into segments, each a section of audio with similar acoustic qualities. These segments are from 80ms to multiple seconds in length. For each segment they calculate the loudness, attack time and the other measures of the variation in the segment. There are also global properties such as tempo and time signature. The features we used are as follows [4]:

- segmentDurationMean: mean segment duration (sec.).
- segmentDurationVariance: variance of the segment duration (sec.^{2})—smaller variances indicate more regular segment durations.
- timeLoudnessMaxMean: mean time to the segment maximum, or attack duration (sec.).
- loudnessMaxMean: mean of segments' maximum loudness (dB).
- loudnessMaxVariance: variance of the segments' maximum loudness (dB^2). Larger variances mean larger dynamic range in the song.
- loudnessBeginMean: average loudness at the start of segments (dB).
- loudnessBeginVariance: variance of the loudness at the start of segments (dB^2). Correlated with loudnessMaxVariance
- loudnessDynamicsMean: average of overall dynamic range in the segments (dB).
- loudnessDynamicsVariance: segment dynamic range variance (dB^2). Higher variances suggest more dynamics in each segment.
- loudness: overall loudness estimate of the track (dB).
- tempo: overall track tempo estimate (in beat per minute, BPM). Doubling and halving errors are possible.
- tempoConfidence: a measure of the confidence of the tempo estimate (beween 0 and 1).
- beatVariance: a measure of the regularity of the beat (secs.^{2}).
- tatum: estimated overall tatum duration (in seconds). Tatums are subdivisions of the beat.
- tatumConfidence: a measure of the confidence of the tatum estimate (beween 0 and 1).
- numTatumsPerBeat: number of tatums per beat
- timeSignature: estimated time signature (number of beats per measure). This is perceptual measures, not what the composer might have written on the score. The description goes as follows: 0=None, 1=Unknown (perhaps too many variations), 2=2/4, 3=3/4 (eg waltz), 4=4/4 (typical of pop music), 5=5/4, 6=6/4. 7=7/4 etc.
- timeSignatureStability: a rough estimate of the stability of the time signature throughout the track

4 ALGORITHMS

We create a feature vector by concatenating the individual feature-analysis results (we used the order described in Section 3, but the order is irrelevant). Let us denote all input features as the matrix f, which is an mxn array of n m-dimensional feature vectors, one vector for each song's analysis results. Further let f_i be the i^{th} feature (column-)vector in f. To measure the distances between different feature vectors, we use learned Mahalanobis metrics [6].

A Mahalanobis (pseudo-)metric is defined as

$$d(f_i, f_j) = (f_i - f_j)^\top \mathbf{M}(f_i - f_j), \qquad (1)$$

where $\mathbf{M}$ is any well-defined positive semi-definite matrix. From Eq. (1) it should be clear that the Euclidean distance is a special case of the Mahalanobis metric with $\mathbf{M} = \mathbf{I}$, the identity matrix. We considered five different algorithms from the research literature to learn a Mahalanobis matrix to convert the raw features into a well-behaved metric space. Each of the algorithms either learns a positive semi-definite

matrix $\mathbf{M}$ or a matrix A, such that $\mathbf{M} = A^\top A$. We can uniquely decompose any positive semi-definite matrix as $\mathbf{M} = A^\top A$, for some real-valued matrix A (up to rotation). This reduces Eq. (1) to

$$d(f_i, f_j) = \|A(f_i - f_j)\|_2, \quad (2)$$

the Euclidean metric after the transformation $f_i \rightarrow Af_i$.

One of the approaches—whitening—is unsupervised, i.e. the algorithm does not require any side-information in addition to the pure feature vectors f. The other four use labels to tune the Mahalanobis matrix A so that similar songs are likely to be close to each other in the metric space. In this study we use album, artist and blog labels for each song as a measure of similarity. We evaluate our algorithm by testing the performance of a nearest-neighbor classifier in these new spaces.

The output of our algorithms is $F = Af$. The learned matrix A is of size $m \times m$ in this paper, but also can be $m' \times m$, where $m' < m$, in which case it reduces the dimensionality of the output space. The result matrix F has n points arrayed so similar points are close together. We partition the algorithms that we discuss into two groups: algorithms based on second-order statistics, and algorithms based on optimization. We will discuss each in turn.

4.1 Algorithms based on second-order statistics

The first three algorithms learn a linear transformation of the input space based on second-order statistics of the feature vectors. These methods rely heavily on the spread of information as captured by an outer product in the covariance calculation

$$cov(f) = \frac{1}{n}\sum_i (f_i - \bar{f}_i)(f_i - \bar{f}_i)^\top \quad (3)$$

where $\bar{f}_i$ is the mean of the feature vector over all songs. This equation is used in two different ways. The within-class covariance function is calculated from all vectors within one class and the between-class covariance is calculated from the means of all class clusters.

Whitening

The easiest way to massage the data is to normalize each dimension of the feature vector so that they all have the same energy. A more sophisticated approach adds rotations so that the covariance matrix of the whitened data is diagonal. We do this by computing

$$A_w = [cov(f)]^{-1/2} \quad (4)$$

where $cov(\cdot)$ is the covariance of a matrix. The covariance of fA_w is the identity matrix. This approach is completely unsupervised because whitening does not take into account what is known about the songs and their neighbors. Whitening is important as it removes any arbitrary scale that the various features might have. To use whitening as a pre-processing for distance computation was originally proposed by Mahalanobis [6] and is the original formulation of the Mahalanobis metric. A potential draw-back of whitening is that it scales all input features equally, irrespective of whether they carry any discriminative signal or not.

LDA

Linear discriminant analysis (LDA) is a common means to find the optimal dimensions to project data and classify it. It is often used as a means of rotating the data and projecting it into a lower-dimensional space for dimensionality reduction.

LDA assumes that the data is labeled with (normally) two classes. It further assumes that the data within each class is distributed with a Gaussian distribution and further assumes that each class of data shares the same covariance matrix. This is likely not true in our case since some artists or albums are more diverse that others. In this work we use a multi-class formulation for LDA proposed by Duchene [3].

LDA optimizes class distinctions, maximizing the between-class spread while minimizing the within-class spread. This procedure is based on the assumption that each class is independently sampled from a single uni-modal distribution so the distribution is characterized by a single mean and variance, which may not apply in many more complicated real world scenarios.

RCA

Relevant component analysis (RCA) [1] is related to whitening and LDA as it is entirely based on second-order statistics of the input data. One can view RCA as local within-class whitening. Different from LDA, it does not maximize the between-class spread and therefore makes no uni-modal assumption on the data.

4.2 Algorithms based on optimization

The next two algorithms explicitly learn a matrix A by minimizing a carefully constructed objective function that mimics the kNN leave-one-out classification error.

The problem with optimizing a nearest-neighbor classifier is that the objective function is highly non-continuous and non-differentiable. Many changes to the solution, the A matrix in this case, make no change to a point's nearest neighbors and thus no change to the objective function. Then an infinitesimally small change to A will shift the nearest neighbors and the objective function will make a large jump. The two algorithms we consider next introduce two different surrogate loss functions whose minimization

loosely translates into the minimization of the kNN leave-one-out classification error.

NCA

In neighborhood component analysis (NCA) the hard decisions incumbent in identifying nearest neighbors are replaced with a soft decision based on distance to the query point [7]. Instead of defining the nearest neighbor as the closest feature vector, Goldberger et al. use a soft-neighborhood assignment. For a given feature vector f_i, the nearest neighbor f_j is picked at random with probability

$$p_{ij} = \frac{e^{-d(f_i,f_j)}}{\sum_k e^{-d(f_i,f_k)}}. \quad (5)$$

In other words, the probability that f_j is a nearest-neighbor of f_i decreases exponentially with the distance between them. Given this objective function, one can compute the probability, p_i, of a data point f_i within class C_i has a nearest neighbor within the same class:

$$p_i = \sum_{f_j \in C_i} p_{ij}. \quad (6)$$

The objective of NCA is to maximize the probability that each point has neighbors in the same class,

$$A_{nca} = \text{argmin} \sum_i p_i(A) \quad (7)$$

where the point probabilities depend implicitly on M. In words, the algorithm maximizes the expected number of classified input feature vectors under a probabilistic 1-NN classification. As Eq. (5) is continuous and differentiable so is the objective in Eq. (6), which can be maximized with standard hill-climbing algorithms such as gradient descent, or conjugate gradient.

The $O(N^2)$ cost of the calculation is mitigated because the exponential weighting falls off quickly so many distant pairs can be safely ignored.

LMNN

The last approach we investigated is large-margin nearest neighbor (LMNN) [10]. Similar to NCA, LMNN is also based on an optimization problem. However, instead of a smooth, non-convex objective, LMNN mimics the kNN leave-one-out classification error with a piecewise linear convex function. This minimization can be solved with well-studied semidefinite programs, which can be solved with standard optimiziation algorithms such as interior-point or sub-gradient descent. A key step in making the objective convex is to fix *target neighbors* for each input vectors prior to learning. These target neighbors must be of the same class and should be close under some reasonable metric. The objective tries to minimize the distance of an input vector to its target neighbors, while enforcing that no differently labeled inputs come closer than 1 unit from the target neighbor.

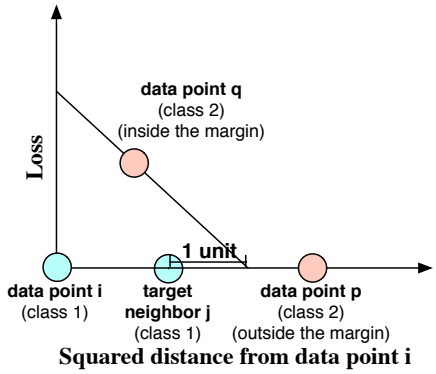

Figure 1. The loss function for LMNN. The loss (or error) increases rapidly for points not in class 1 that encroach too close to a point in the query class.

Partially inspired by support vector machines (SVM), the objective consists of two parts: One that forces target neighbors to be close, and a second that forces a margin between an input vector and differently-labeled vectors. Let $j \rightsquigarrow i$ denote that f_j is a target neighbor of f_i, then we write the objective as

$$\sum_{j \rightsquigarrow i} d(f_i, f_j) + \sum_{j \rightsquigarrow i} \sum_{k \notin C_i} [d(f_i, f_j) + 1 - d(f_i, f_k)]_+, \quad (8)$$

where C_i is the class of f_i and $[a]_+ = max(a, 0)$. The second term of this objective function pushes dissimilar points at least one unit away, as illustrated in Figure 1.

As the objective function in Eq. (8) is piece-wise linear, it can be efficiently minimized over large data sets $N > 60000$. One potential weakness of LMNN is the fact that the target neighbors are fixed before the optimization and their choice significantly impacts the final metric. In this paper we chose them to be the nearest neighbors under the Euclidean metric after whitening.

5 EVALUATION

We evaluate each metric algorithm by testing a kNN classifier's ability to recognize the correct album, artist or blog that describe each song. We do this by organizing all data by ID, and then selecting enough IDs so that we have more than 70% of all songs in a training set. The remaining data, slightly less than 30%, is a test set. The classifier's task is to look at each test point, and see if at least 2 of its 3 neighbors have the desired ID.

Thus we train a Mahalanobis matrix on a large fraction of our data, and test the matrix by measuring identification

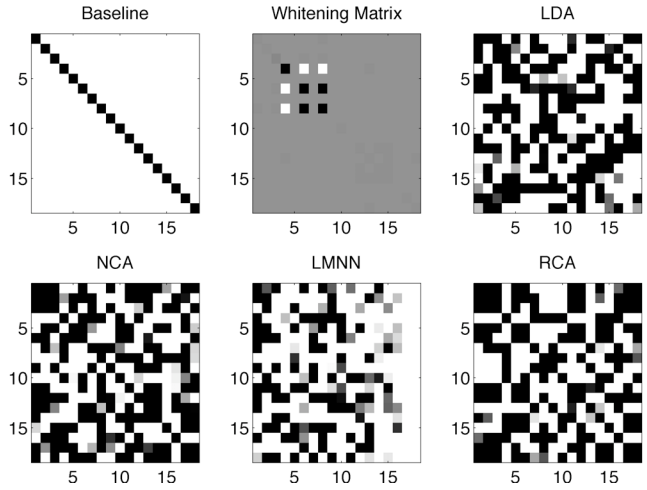

Figure 2. Summary of Mahalanobis matrices derived by each algorithm based on album similarity. The whitened matrix has both positive and negative terms centered around the gray of the background. The other matrices have been scaled so that white is at 0 and black represents the maximum value. All features are in the same order as described in Section 3.

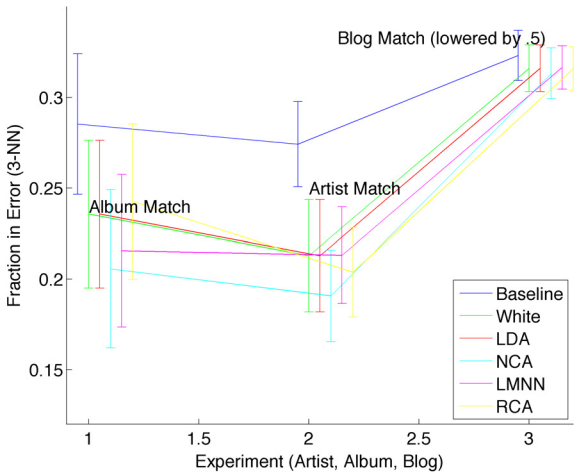

Figure 3. Summary of metric algorithms kNN performance. The results for each of the six metrics are offset horizontally, in the order shown in the legend, to facilitate comparison. Note, the performance of the blog-identification task was very poor, and we have reduced each blog error by 0.5 to fit them on the same graph as the others.

performance on data it has never seen. We do this for album and artist ID, and also try to predict the blog that mentions this song.

We found that adding a whitening stage before *all* algorithms improved their performance. Thus each of the four algorithms we investigated (LDA, NCA,LMNN, RCA) are preceded by a full whitening step. We also compare these algorithms to two strawman: a baseline using the original (unweighted) feature space, and the whitened feature space with no other processing.

We are interested in the robustness of these algorithms to noise. To test this, we measured performance as we added noisy features. Each feature is a zero-mean Gaussian random variable with a standard deviation that is 10% of the average for the real features. This level is arbitrary since all algorithms performed best, in testing, when the data is whitened first. This removes the level dependence on all but the baseline data.

6 RESULTS

Figure 2 shows all 6 Mahalanobis matrices. The four entries in the whitening matrix with the largest values are the four loudness features. Except for LDA, the other matrices make relatively small changes to the feature space.

Yet, these small changes in the Mahalanobis matrices have significant difference in performance. Figure 3 shows the performance of all six approaches on all three identification tasks. We performed 10 trials for each classification test, in each case choosing at random new (non-overlapping) training and testing sets. In these tests, NCA did best, but LMNN and RCA were close seconds on the album-match and artist-match tasks respectively. In our tests, both album and artist ID are relatively easy, but identifying the blogger who referenced the song was problematic. This suggests that album and artists are better defined than a blogger's musical interests.

Figure 4 shows the performance as we add noisy features. In this experiment we added noisy dimensions to simulate the effect of features that are not relevant to the musical queries. The error for the baseline quickly grows, while the other methods do better. This test was done for the album-recognition task. In this case NCA and RCA perform best, and this is significant because the computational cost of RCA is trivial compared to that of NCA and LMNN.

One explanation for the relative win of RCA over LMNN (and LDA) is that the later algorithms try to push different classes apart. This might not be possible, or even a good idea when there are as many classes as in our experiment. There just isn't any room for the classes to separate. Thus in our tests, especially when noise is added, the overlapping classes are not amenable to separation.

All of these algorithms have the ability to do feature selection and reduce the dimensionality of the feature space. LDA and RCA order the dimensions by their ability to predict the data clouds, so it's natural to cut off the smaller dimensions. Both NCA and LMNN are essentially optimization problems, and by posing the problem with a rectangular instead of a square A matrix one can calculate a low-

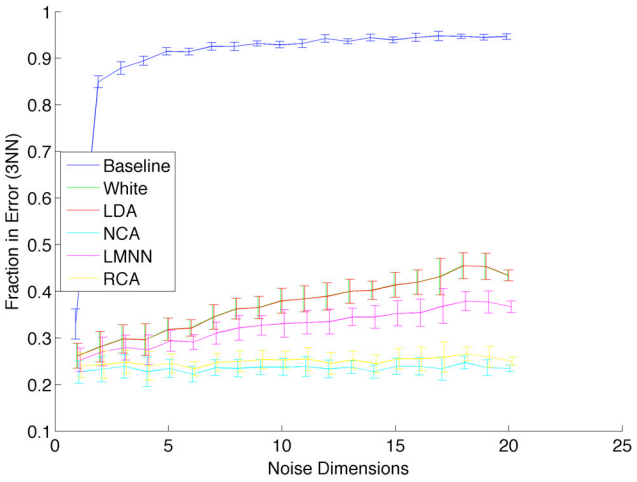

Figure 4. Summary of metric algorithms kNN performance with additional noisy features. We have augmented the original 18 features with additional purely noise features to measure the ability of each algorithm to ignore the noisy dimensions. The results for each of the six metrics are offset horizontally, in the order shown in the legend, to facilitate comparison.

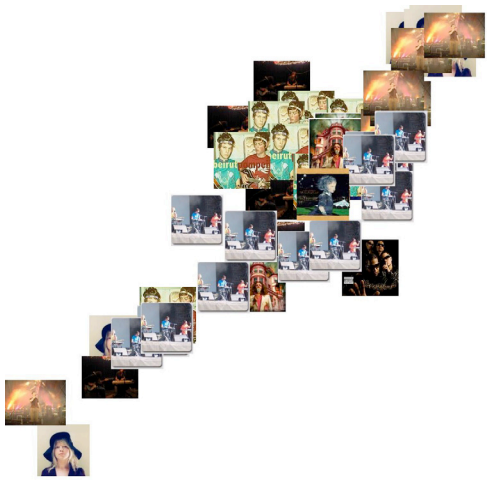

Figure 5. We used NCA to find the optimal projection of the 9 most common artists. Each image indicates the location of one or more songs by that artist in a two-dimensional space.

dimensional embedding. We illustrate this kind of dimensionality reduction in Figure 5. Dimensionality reduction can be important in a nearest-neighbor recognizer because one must store all the prototypes, and the dimensionality of the feature space directly links to the amount of memory needed to store each song's feature vector.

7 CONCLUSIONS

In this paper we have described and demonstrated 6 different means to embed acoustic features into a metric space. In the best-performing cases the algorithms use meta data about the songs—in our case album, artist, or blog IDs— to tune the space so that songs with the same ID are close to each other. With our data, more than 5000 songs described on music blogs, we found that all algorithms lead to a significant improvement in kNN classification and, in particular, NCA and RCA perform by far most robustly with noisy input features. More work remains to be done to verify that these results produce sensible playlists and pleasing transitions. We would also like to investigate which features are important for these problems.

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SUPPORT FOR MIR PROTOTYPING AND REAL-TIME APPLICATIONS IN THE CHUCK PROGRAMMING LANGUAGE

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ABSTRACT

In this paper, we discuss our recent additions of audio analysis and machine learning infrastructure to the ChucK music programming language, wherein we provide a complementary system prototyping framework for MIR researchers and lower the barriers to applying many MIR algorithms in live music performance. The new language capabilities preserve ChucK's breadth of control—from high-level control using building block components to sample-level manipulation—and on-the-fly re-programmability, allowing the programmer to experiment with new features, signal processing techniques, and learning algorithms with ease and flexibility. Furthermore, our additions integrate tightly with ChucK's synthesis system, allowing the programmer to apply the results of analysis and learning to drive real-time music creation and interaction within a single framework. In this paper, we motivate and describe our recent additions to the language, outline a ChucK-based approach to rapid MIR prototyping, present three case studies in which we have applied ChucK to audio analysis and MIR tasks, and introduce our new toolkit to facilitate experimentation with analysis and learning in the language.

1. INTRODUCTION

ChucK [13] began as a high-level programming language for music and sound synthesis, whose design goals included offering the musician user a wide breadth of programmable control—from the structural level down to the sample level—using a clear and concise syntax, and employing a set of abstractions and built-in objects to facilitate rapid prototyping and live coding. We have recently expanded the language to provide support for audio analysis and machine learning, with two primary goals: first, to offer real-time and on-the-fly analysis and learning capabilities to computer music composers and performers; and second, to offer music information retrieval (MIR) researchers a new platform for rapid prototyping and for easily porting algorithms to a real-time performance context. We address the former goal in [6]; here, we focus on the latter.

We begin in Section 2 by reviewing prototyping in MIR and motivating the need for additional shared tools between MIR and performance. We describe the ChucK language as it is used for music creation, including prototyping and live coding systems. In Section 3, we describe in some detail how we have incorporated analysis and learning into the language, with attention both to preserving the flexible and powerful control that makes ChucK suited for prototyping and experimentation, and to tightly and naturally integrating the new functionality with ChucK's synthesis framework. Sections 4 and 5 illustrate the new potential of ChucK as an MIR rapid prototyping workbench, introducing an example working pipeline for prototyping an MIR task and presenting three examples of how we have used ChucK to perform and teach music analysis. Finally, in Section 6 we discuss ongoing work to improve ChucK as an MIR tool and announce the release of a toolkit containing examples and supporting code for MIR researchers desiring to experiment with the language.

2. BACKGROUND AND MOTIVATION

2.1. MIR Prototyping

One primary goal in this work is to provide a new rapid prototyping environment for MIR research. Our definition of an MIR prototyping environment includes: the ability to design new signal processing algorithms, audio feature extractors, and learning algorithms; the ability to apply new and existing signal processing, feature extraction, and learning algorithms in new ways; and the ability to do these tasks quickly by taking advantage of high-level building blocks for common tasks, and by controlling the system either via a GUI or through concise and clear code. Furthermore, a prototyping system should encourage experimentation and exploration. There exist several programming environments for MIR that accommodate many of the above requirements, including Matlab, M2K[1], MARSYAS [12], CLAM [1], and Weka [15], and their popularity suggests that they meet the needs of many MIR research tasks. We propose that ChucK also meets all of these requirements and inhabits a complementary place in the palette of tools at the MIR researcher's disposal. In particular, ChucK features real-time support for analysis and synthesis, easy accommodation of concurrency and auditory feedback, and a syntax and running environment

[1] http://www.music-ir.org/evaluation/m2k/index.html

that encourage a development cycle rapid enough for live-coding, in which the performer modifies the code in a live performance [14]. As we discuss below, ChucK is therefore able to facilitate a rapid, feedback-driven, real-time and performance-enabled approach to MIR prototyping that is not a primary concern of other tools.

2.2. MIR and Music Performance

Research in MIR has primarily focused on analyzing and understanding recorded audio, static symbolic representations (e.g., MIDI or Humdrum files) and other non-performative musical representations and metadata. There is some exceptional work, notably real-time score-following and accompaniment systems such as [8], [11]. However, many other popular music information retrieval tasks, such as mood and style analysis and instrumentation and harmony identification, are directly pertinent to real-time interactive performance, and the relevance of core MIR work to live music remains under-exploited.

For one thing, a primary focus of MIR is building computer systems that understand musical audio at a semantic level (e.g., genre, rhythm, mood, harmony), so that humans can search through, retrieve, visualize, and otherwise interact with musical data in a meaningful way. Making sense of audio data at this higher level is also essential to *machine musicianship*, wherein the performing computer—like any musically trained human collaborator—is charged with interacting with other performers in musically appropriate ways [10].

Despite the shared need to extract useful features from audio, and to bridge the gap between low-level audio features and higher-level musical properties, there does not exist a shared tool set for accomplishing these tasks in both computer music and MIR. On one hand, most computer music languages do not readily accommodate analysis of audio *in the language*; for example, highly custom spectral analysis and processing tasks must be coded in C++ externals in order to be used in SuperCollider or Max/MSP. Externals' development overhead, enslavement to the audio and control rates and APIs of the associated music language, and unsuitability as recursive building blocks within larger analysis algorithms make them an unattractive choice for MIR researchers (not to mention musicians). On the other hand, most MIR toolkits and frameworks do not support real-time audio processing or synthesis. Those that do (namely MARSYAS [3] and CLAM [1]) do not offer the full-fledged synthesis and interaction support of a computer music *language*, so while popular among MIR researchers, they do not have widespread adoption among computer musicians.

Computer musicians would undoubtedly benefit from lower barriers to adapting state-of-the-art MIR algorithms for pitch tracking, beat tracking, etc. for their real-time performance needs. We additionally posit that MIR researchers can benefit from increased collaboration with musicians and composers, which does not suffer from the common challenges of copyright restrictions on obtaining data or releasing systems to the public, nor the difficulty and expense in obtaining reasonable ground truth to perform evaluations. Additionally, applying MIR research in music performance can offer researchers outside of industry a greater potential to directly impact people's experiences with music.

We realize that no single tool will meet the needs of everyone in computer music and MIR, but we propose to lower barriers to working at the intersection of these fields—in analysis-driven computer music and real-time, potentially performance-oriented MIR—via a full-fledged computer music language that allows for arbitrarily complex signal processing and analysis tasks within a user-extensible, object oriented framework.

2.3. ChucK

ChucK is a cross-platform, open-source computer music programming language [13] whose primary design goals include the precise programmability of time and concurrency, with an emphasis on encouraging concise, readable code. System throughput for real-time audio is an important consideration, but first and foremost, the language was designed to provide maximal control and flexibility for the audio programmer. In particular, key properties of the language are as follows:

- *Flexibility*: The programmer may specify both high-level (e.g., patching an oscillator to a filter, or initiating structural musical events) and low-level, time-based operations (e.g., inspecting and transforming individual samples) in a single unified language mechanism, without any need for externals.
- *Readability*: The language provides a strong correspondence between code structure, time, and audio building blocks; as a result, the language is increasingly being used a teaching tool in computer music programs, including at Princeton, Stanford, Georgia Tech, and CalArts.
- *Modularity*: Like many other languages, ChucK encapsulates behaviors of synthesis building blocks (filters, oscillators, etc.) using the "Unit Generator" model. Its object-oriented nature also supports modular user code.
- *A "do-it-yourself" approach*: By combining the ease of high-level computer music environments with the expressiveness of lower-level languages, ChucK supports high-level musical representations, as well as the prototyping and implementation of low-level, "white-box" signal-processing elements in the same language.
- *Explicit treatment of time*: There is no fixed control rate. It's possible to assert control on any unit generator at any point in time, and at any rate, in a sub-sample-precise manner.

- *Concurrency*: Parallel processes (called shreds) share a notion of time, so one can precisely synchronize and easily reason about parallel code modules according to each's treatment of time. Parallelism is easy and expressive.
- *On-the-fly programming*: Programs can be edited as they run; this functionality is supported and encouraged in the miniAudicle development environment[1].

The following example synthesis code illustrates several of these concepts. In it, an impulse train is synthesized, and its frequency is modified in a concurrent parallel function at an arbitrary "control rate."

```
// patch impulse generator to output
Impulse i => dac;

50::samp => dur impulsePeriod; // start at 50 samples
spork ~ sweepPeriod(); // run function in parallel

// generate an impulse every period
while( true ) {
    1.0 => i.next;        // fire impulse
    impulsePeriod => now; // advance time
}

// sweep the period from 50 to 200 samples
// updating every .1 second
fun void sweepPeriod() {
    while( impulsePeriod < 200::samp ) {
        impulsePeriod * 1.01 => impulsePeriod;
        .1::second => now;
    }
}
```

Figure 1: Simple concurrent ChucK code. Note that '… => now' acts to control program execution in time, and '=>' alone acts as a left-to-right assignment operator.

3. ADDITIONS TO CHUCK

3.1. Unit Analyzers

In previous work, we introduced a language-based solution to combining audio analysis and synthesis in the same high-level programming environment of ChucK [15]. The new analysis framework inherited the same sample-synchronous precision and clarity of the existing synthesis framework, while adding analysis-specific mechanisms where appropriate. The solution consisted of three key components. First, we introduced the notion of a Unit Analyzer (UAna), similar to its synthesis counterpart, the Unit Generator (UGen), but augmented with a set of operations and semantics tailored towards analysis. Next, we introduced an augmented dataflow model to express dependencies among unit analyzers (UAnae), and to manage caching and computation accordingly. Third, we ensured that the analysis framework made use of the existing timing, concurrency, and on-the-fly programming mechanisms in ChucK as a way to precisely control analysis processes.

```
// set up analysis network for mic input
adc => FFT fft =^ Centroid c => blackhole;
fft =^ RollOff r;

// (would set FFT window, size, etc. here)
// ...

// launch in parallel
spork ~ computeRolloff();

// compute centroid every 512 samples
while( true ) {
    c.upchuck();
    512::samp => now;
}

fun void computeRolloff() {
    // compute rolloff every 1 second
    while( true ) {
        r.upchuck();
        1::second => now;
    }
}
```

Figure 2: An example analysis patch

For example, it's possible to instantiate an FFT object for spectral analysis, instantiate spectral centroid and spectral rolloff objects that compute using the output of the FFT, and trigger the extraction of each feature at its own rate. Code for this task appears in Figure 2. The programmer has dynamic, sample-precise control over analysis parameters such as FFT/IFFT sizes, analysis windows, hop sizes, feature extraction rates, etc. The primary advantages of using ChucK for analysis are threefold:

- *Conciseness*: ChucK implicitly manages real-time audio and buffering, and the language is tailored for audio, so analysis system implementation time and code length are greatly reduced.
- *Rapid turnaround experimentation*: Through the application of on-the-fly programming and ChucK's concise audio programming syntax, one can quickly prototype systems and sub-systems, changing parameters as well as the underlying system structure, and experience the results almost immediately.
- *Concurrency*: Parallel management of feature extraction, even at multiple rates, is straightforward in ChucK. This sample-synchronous concurrency would be extremely challenging and/or inefficient in C++, Java, or a library built in these languages, due to their support for preemptive, thread-based concurrency, and the consequent need to contend with thread-instantiation, synchronization, bookkeeping, etc.

[1] http://audicle.cs.princeton.edu/mini/

These highly useful flexibilities come with a tradeoff: system performance and throughput. A system implemented in reasonable ChucK code would likely run much less efficiently than an optimized C++ implementation. In this regard, it may be desirable to leverage the flexibility and rapid experimentation abilities of ChucK to prototype systems and components, and if needed, then implement "production" code in a low-level language with an optimizing compiler. But for researchers experimenting with new MIR algorithms, such a prototyping stage can be instrumental in crafting new systems and testing the feasibility of new ideas.

3.2. Learning Framework

A natural consequence of ChucK's analysis capabilities is that analysis results can be treated as *features* whose relationship to high-level musical concepts can be learned via labeled examples and standard classification algorithms. Classification is a common and powerful technique in MIR, and it has been applied successfully to MIR problems such as genre [2], mood [7], and transcription [5]. Classification has also been widely used in computer music for tasks such as pitch tracking [9], and learning is a key component of the aforementioned accompaniment and score-following systems. ChucK's learning framework was designed with the recognition that a general tool for learning could be useful for accomplishing both MIR analysis tasks and creative compositional tasks, and with acknowledgment that classification of streaming audio is a task that should be natural to perform in real time.

We have built an object-oriented classification framework in ChucK, modeled on the architecture and naming conventions of Weka [16], a popular Java framework for applied machine learning. Classification algorithms such as k-nearest-neighbor and AdaBoost are implemented as ChucK classes which inherit from a parent Classifier class. Each Classifier can be trained on a dataset, represented by an Instances class, and trained Classifiers can be used to assign class predictions to new Instance objects. Because this infrastructure is all written in ChucK, users can easily add their own Classifier child classes in Chuck, as well as modify not just classification parameters but also the underlying algorithms on-the-fly. We have also implemented a suite of standard MIR features, listed in Table 1.

Note that the implementation of ChucK's analysis and learning capabilities preserves its suitability for rapid prototyping, and extends this capability to many MIR tasks. In particular, the user retains sample-level control over audio processing, allowing easy implementation of new feature extraction methods and signal processing algorithms directly in the language. Learning algorithms are also implemented in the language, affording the user arbitrary modification of these algorithms at any time, without the need to recompile. Additionally, the user can choose from standard MIR features and standard classifiers that have already been implemented, and use these out-of-the-box. Furthermore, the ability to code features and algorithms precisely in ChucK, within its object-oriented framework, means that the user can create new "white-box" features and classifiers for immediate usage. In short, ChucK meets the requirements we believe to be important to a prototyping environment.

FFT, IFFT, DCT, IDCT	LPC coefficients
Spectral centroid, spread, rolloff, and flux	Zero crossing rate
	RMS
Mel- and real-frequency cepstral coefficients	Cross- and auto-correlation

Table 1: MIR features in ChucK

4. AN MIR PROTOTYPING PIPELINE

Here we provide an example of how an MIR student or researcher might approach working with the MIR prototyping system of ChucK. Perhaps someone has an idea for an audio analysis task. One can use ChucK to write relevant code immediately: in particular, the programmer can 1) quickly instantiate and connect together essential unit analyzers (FFT's, feature extractors, etc.), then 2) specify any initial parameters, and 3) write control code, potentially in concurrent chuck processes (shreds). If a learning algorithm is part of the task, the programmer will additionally instantiate a classifier and set its parameters, and include control code for passing the extracted feature vectors to the classifier during training and testing stages. The programmer can ignore issues of audio buffering, dependencies among analysis results (e.g., spectral centroid requires an FFT), concurrency management, etc., as these are handled by ChucK, and instead focus on the content of the algorithm.

As soon as a basic skeleton for the algorithm is written, the programmer can immediately run the code and observe the output (which can easily be sonified, if desired). At this point, the programmer can augment or tweak the parameters *and* underlying structures of the system via on-the-fly programming. For example, the programmer can easily change STFT window type, size, zero-padding, and hop size, or add and remove the features used in classification, and immediately observe the consequences of these changes. It is in this focal stage of prototyping, wherein the programmer interactively tunes system performance and logic, that ChucK is most uniquely powerful as a prototyping tool, aided by its live-coding support in miniAudicle.

5. EXAMPLES

We now present three examples of music analysis systems that have been built in ChucK. The first is a trained vowel

identification system with a GUI interface, which illustrates the conciseness of interactive ChucK code and ease with which GUIs can be incorporated. The second is an implementation of a genre and artist classification system described in [2], which illustrates the ease of porting a complex MIR feature extraction and classification system to ChucK and applying it to real-time audio input. The third is a discussion of our experiences employing ChucK's MIR capabilities as an educational tool, illustrating its usefulness to MIR and computer music novices. The code for case studies 1 and 2 is publicly available (see Section 6).

5.1. Simple Visualizations and Classifier

In order to visualize spectral properties of a sound in real-time, a filter-based sub-band spectral analyzer was implemented in ChucK using MAUI, miniAudicle's framework for graphical widgets. The GUI appears in Figure 3(a); sliders update to indicate the power in each octave sub-band in real-time. The entire code for this task is 25 lines, 15 of which manage the display.

With a simple modification, this code and display can be used to train a classifier on any task for which sub-band power is a useful feature, then perform classification (and useful visual feedback) on new inputs. For example, code that uses 1/3-octave sub-bands from 200Hz to 4kHz to train a nearest-neighbor classifier to recognize each of 5 vowels spoken by the user (and silence), to classify the vowel of new inputs, and to construct the training and feedback graphical interface shown in Figure 3(b), requires under 100 lines of code. This includes the nearest-neighbor classifier coded from scratch (i.e., the KNN ChucK class was not used here). Clearly, ChucK allows for concise specification of signal processing analysis tasks and for real-time GUI display and input.

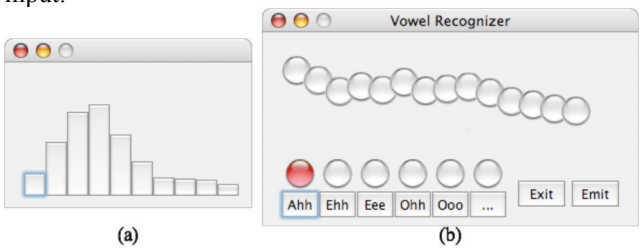

Figure 3: GUIs for (a) octave-band spectral analysis and (b) trained vowel recognition

5.2. Artist Classification

Of obvious concern to MIR researchers interested in ChucK is whether more complex systems are possible. To address this, we have implemented one of the genre and artist classification systems of Bergstra et al. described in [2]. It uses eight types of standard audio features, including FFT coefficients, real cepstral coefficients, mel-frequency cepstral coefficients, zero-crossing rate, spectral spread, spectral centroid, spectral rolloff, and LPC coefficients (all are available in ChucK). Means and variances are computed for each feature over a segment of consecutive frames, then classified by an AdaBoost classifier using decision stumps as the weak learners. Bergstra's system performed in the top two submissions for both genre and artist classification at MIREX 2005 [4].

We have embedded this classification approach in a simple interface for real-time, on-the-fly training and classification of audio. The user can specify via a keyboard or graphical interface that the incoming audio provides examples of a particular class (thereby initiating the construction of labeled Instances from features extracted from this audio), initiate the training rounds of AdaBoost using all available Instances, and then direct the trained classifier to classify the incoming audio and output the class predictions to the screen.

Our experimental on-the-fly application of this system reveals the impressive strength of the features and classifier; in fact, training on only 6 seconds of audio from each of two artists, the system is able to classify new songs' artists with over 80% accuracy. Furthermore, it is easy to use the keyboard interface to experiment with applying this system to new tasks, for example speaker identification and instrument identification, without modification to the algorithm or features.

5.3. ChucK as Introductory MIR Teaching Tool

We have used ChucK to teach audio analysis and basic MIR in introductory computer music courses at Princeton and Stanford. By leveraging the clarity and flexibility of representing analysis and MIR concepts in the language, as well as ChucK's rapid prototyping and on-the-fly programming capabilities, we were able to teach the following topics to students familiar with the basics of the language:

- Practical real-time, short-time Fourier analysis on audio signals (both recorded and live), with on-the-fly demonstration of the effects of window type, window size, FFT size, and hop size
- Standard spectral- and time-domain audio features, their extraction algorithms, and real-time exploration of relationships between audio signals and feature values
- Classification, including applications to speaker identification, vowel/consonant analysis, and physical gesture recognition
- Basic pitch and beat tracking using spectral processing

Additionally, at Stanford, these concepts were taught in conjunction with a large class project to design a computer-mediated performance. Although the use of analysis and MIR algorithms was not a project requirement, more than one-third of the class integrated the extraction of some high-level musical information as a

component of their final projects. These ranged from real-time algorithmic processes that employed extracted features to create compelling accompaniments to a live flutist, to real-time speech analysis that was transformed into gestures controlling a Disklavier, to amplitude and pitch event-triggered generative "sonic clouds." While these projects used very basic MIR components, it was encouraging to witness the students (most of whom had not worked with MIR or audio analysis before) eagerly and efficiently experiment in a 2–3 week period and craft successful musical performances from these investigations.

6. ONGOING WORK AND CONCLUSIONS

We have made available the ChucK learning infrastructure described in Section 3.2, code for the first and second examples described in Section 5, as well as code for several other example tasks, as part of the Small Music Information Retrieval toolKit (SMIRK)[1]. SMIRK will provide a permanent and growing open-source repository for music information retrieval infrastructure and examples using ChucK, targeted at researchers, educators, and students.

Several improvements to ChucK's MIR capabilities are currently underway. First, support for asynchronous file I/O will allow the reading and writing of large datasets in a way that does not interfere with audio production. Second, we are adding more classifiers and support for modeling, beginning with hidden Markov models.

Our recent additions to ChucK have greatly expanded its capabilities beyond the realm of sound synthesis and music performance. Through the integration of analysis and learning tools, we hope to lower the barriers to applying MIR algorithms in real-time settings and suggest a new prototyping paradigm for signal processing, feature extraction, and learning components of MIR systems. We are excited by the nascent potential for new creative and technical work by researchers, musicians, educators, and students using this new framework.

7. ACKNOWLEDGEMENTS

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[1] http://smirk.cs.princeton.edu

A COMPARISON OF SIGNAL-BASED MUSIC RECOMMENDATION TO GENRE LABELS, COLLABORATIVE FILTERING, MUSICOLOGICAL ANALYSIS, HUMAN RECOMMENDATION, AND RANDOM BASELINE

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ABSTRACT

The emergence of the Internet as today's primary medium of music distribution has brought about demands for fast and reliable ways to organize, access, and discover music online. To date, many applications designed to perform such tasks have risen to popularity; each relies on a specific form of music metadata to help consumers discover songs and artists that appeal to their tastes. Very few of these applications, however, analyze the signal waveforms of songs directly. This low-level representation can provide dimensions of information that are inaccessible by metadata alone. To address this issue, we have implemented signal-based measures of musical similarity that have been optimized based on their correlations with human judgments. Furthermore, multiple recommendation engines relying on these measures have been implemented. These systems recommend songs to volunteers based on other songs they find appealing. Blind experiments have been conducted in which volunteers rate the systems' recommendations along with recommendations of leading online music discovery tools (Allmusic which uses genre labels, Pandora which uses musicological analysis, and Last.fm which uses collaborative filtering), random baseline recommendations, and personal recommendations by the first author. This paper shows that the signal-based engines perform about as well as popular, commercial, state-of-the-art systems.

1 INTRODUCTION

The nature of online music distribution today is characterized by massive catalogs of music unbounded by physical constraints. As pointed out in [1], current technology has offered music listeners "massive, unprecedented choice in terms of what they could hear". The number of songs available on-line is in the billions, and many millions of users are continuing to flock from traditional means of obtaining music (e.g., CD stores) to online alternatives [11].

With such a vast amount of music available on the Internet, end users need tools for conveniently discovering music previously unknown to them (whether recently released or decades old). In the context of electronic music distribution, it is the goal of today's online discovery tools to automatically recommend music to human listeners. This is no simple task; a program must have an automated way of computing whether or not one song is, in some sense, similar to some other set of songs (i.e., to songs that are already liked by the user to whom the program is recommending new music). In accordance with this goal, we have designed and implemented three systems that use signal-based music similarity measures to recommend songs to users.

In this paper, we first discuss existing methods of automatic music recommendation, including a discussion of commercial, state-of-the-art systems that use them, in Section 2. Next, we discuss techniques for automatically computing signal-based music similarity, including a description of our own similarity measures, in Section 3; the optimization of the measures is discussed in Section 4. In Section 5, we discuss how these similarity measures have been used to design and implement three automatic, signal based music recommendation engines. Section 6 describes experiments in which volunteers have rated the recommendations of these systems, along with those of the popular systems described in Section 2, a baseline system, and human recommendations. We evaluate the results of these experiments in Section 7. We then state some general conclusions in Section 8.

2 STRATEGIES FOR AUTOMATIC MUSIC RECOMMENDATION

Three possible strategies of automatic music recommendation involve expert opinions, collaborative filtering, and musicological analysis. Recommendation by expert opinion often relies on the application of genre labels to songs and artists. The wide variety of music genre labels has arisen through a multifaceted interplay of cultures, artists, music journalists, and market forces to make up the complex hierarchies that are in use today [16]. Currently, the largest database of music that is organized by genre is Allmusic [1],

[1] http://www.allmusic.com/

where professional editors compose brief descriptions of popular musical artists, often including a list of similar artists [6]. In the context of automatic music recommendation, recent research has effectively pointed out significant deficiencies of the traditional genre labeling methodology. For one, as discussed in [16], there is no general agreement in the music community as to what kind of music item genre classification should be consistently applied: a single song, an album, or an artist. Second, as discussed in [14], there is no a general agreement on a single taxonomy between the most widely-used music databases on the Internet. Lastly, it is noted in [16] that the criteria for defining music genres have, for countless years, been inconsistent; some labels are geographically defined, some are defined by a precise set of musical techniques, while others arise from the lexical whims of influential music journalists.

Given these inconsistencies, musicological analysis aims to determine music similarity in a way that transcends conventional genre labels, focusing primarily on music theoretic description of the vocal and instrumental qualities of songs. This technique was spearheaded by the Music Genome Project (MGP) in 2000, whose research culminated in the music discovery website/tool Pandora [2] [10]. The automatic recommendation algorithm behind Pandora involves comparisons of very particular descriptions of songs. The description process involves analysis of songs by a team of professional music analysts, each song being represented by about 150 "genes," where each gene describes a musicological quality of the song. Perhaps the most apparent drawback of musicological analysis — especially in the context of Pandora — is that while the recommendation process is automated, the description aspect is not. It is this aspect that contributes to the relatively slow rate at which new content is added to the Pandora database.

Also designed within the context of online music discovery, collaborative filtering works according to the principle that if songs or artists you like occur commonly in other users' playlists, then you will probably also like the other songs or artists that occur in those playlists. According to [8], "if your collection and somebody else's are 80% alike, it's a safe bet you would like the other 20%". One of the most popular on-line recommendation engine to use collaborative filtering is Last.fm [3] , which boasts 15 million active users and 350 million songs played every month [12]. One problem with collaborative filtering systems is that they tend to highlight popular, mainstream artists. As noted in [8], Last.fm "rarely surprises you: It delivers conventional wisdom on hyperdrive, and it always seems to go for the most obvious, common-sense picks." In other words, collaborative filtering is not helpful for discovering lesser known music which a user might highly appreciate.

The past several years have seen considerable progress in the development of mathematical methods to quantify musical characteristics of song waveforms based on the content of their frequency spectra. In particular, these methods have enabled the extraction of features of a song's waveform that are correlated with the song's pitch, rhythmic, and timbral content. Timbre can be said to be the most important of these three elements when subjectively assessing musical similarity between a pair of songs; indeed, it may even be said that the global timbral similarity between two pieces of music is a reasonable — and often sufficient — estimate of their overall musical similarity [4].

These research efforts have also gone on to evaluate and test several different timbre-based music similarity measures applied to a number of signal-based music information retrieval tasks, including supervised and unsupervised classification of entire music databases and the segmentation and summarization of individual songs [16]. Following the lead of these efforts, we have applied signal-based measures of music similarity to the task of automatic recommendation of music. An automatic recommendation engine built on a signal-based music similarity measure would possess the advantages that current online music discovery tools merely trade off. It would boast the ability to describe and compare pieces of music based purely on their musical qualities, and would also facilitate the rapid addition of new content to a music database that does not require human intervention.

3 COMPUTING MUSIC SIMILARITY

Our signal based recommendation engines rely on the ability to automatically compute the similarity of two songs. First, the relevant information about each song — features — is computationally derived from its waveform data. Second, a compact representation of the song is obtained by modeling the distribution of its feature data using mixture and clustering algorithms. Third, a metric for comparing mixture models of songs is used to estimate the similarity between the feature distributions of two different songs. In effect, the timbral similarity between the two songs is mathematically computed.

As a whole, this music similarity measure framework allows a user to present a song query to the signal-based recommendation engine and receive a set of song recommendations (i.e., similar songs) drawn from a target music database. The similarities of the recommended songs to the query song are determined via signal processing alone, without human intervention. In this section, a general overview is given of the three similarity measures examined in this paper. Our implementations of these measures are based partly on those proposed in [9, 15, 17].

The music feature dataset extracted by the measures' analysis front-ends are the Mel-frequency cepstral coefficients (MFCC's). These perceptually-motivated features capture the "spectral shape" — and effectively, the timbral quality

[2] http://www.pandora.com/

[3] http://www.last.fm/

— of a music signal within a small frame of the waveform [18, 6]. In the literature, the MFCC feature set has already shown effective performance for various audio classification experiments [6, 18, 13, 3].

3.1 K-Means Clustering with Earth Mover's Distance

The first similarity measure used in our work was originally proposed by Logan and Salomon in their work in [13]. In this architecture, K-means clustering of a target song's feature vectors is performed during the statistical modeling stage, with each data cluster being subsequently fit with a Gaussian component to form a Gaussian Mixture Model (GMM). Also in line with what was proposed in [13], the distance metric stage of the first similarity measure incorporates the Earth Mover's Distance (EMD). The EMD expands the Kullback-Leibler divergence — a distance metric for comparing individual probability distributions — to the comparison of mixtures of distributions (in this case, GMM). For the remainder of the paper, this similarity measure combining K-means training of GMM's with the Earth Mover's Distance for GMM comparison is referred to by the shorthand term *KM+EMD*.

3.2 Expectation-Maximization with Monte Carlo Sampling

The second similarity measure that we have relied on uses the Expectation-Maximization (EM) algorithm to train the parameters of each GMM component. Aucouturier and Pachet introduced and refined the use of EM to model music feature distributions in [2, 3, 4]. This method makes use of vectors sampled directly from the GMM's of the two songs to be compared; the sampling is performed computationally via random number generation. This sampling process corresponds roughly to recreating a song from its timbre model [4], and is known as Monte Carlo Sampling (MCS). Using MCS in conjunction with GMM training via Expectation-Maximization is in line with what was originally proposed by Aucouturier and Pachet in [2, 3, 4]. For the remainder of the paper, the similarity measure based on this approach is referred to as *EM+MCS*.

3.3 Average Feature Vector with Euclidean Distance

In the early work of Tzanetakis and Cook [18], a simple way is presented to construct an averaged vector representation of a song's MFCC's. They propose that low-order statistics such as mean and variance should be calculated over segments called texture windows that are more meaningful perceptually. With respect to human auditory perception, the length of a so-called texture window roughly corresponds to the minimum duration of time required to identify a particular sound or music "texture" that corresponds to its overall timbral character. This has led us to test a simpler similarity measure which does not involve the training of a GMM. For each song, a single "average feature vector" is constructed from means and variances taken across the texture windows of the song's waveform. The song's representative vector may then be compared to that of another song by taking the Euclidean distance between them. The similarity measure based on this approach is referred to as *AV+EUC*.

4 PARAMETER OPTIMIZATION

In order to use any of the similarity measures discussed in Section 3, the values of several parameters must be selected. Perhaps most importantly are the dimensionality of the MFCC vectors (N) and the number of Gaussian components in a GMM (M). The parameter M is not applicable when using AV+EUC as a similarity measure. Other parameters include the sampling frequency of the song waveforms (f_s), the frame length (N_f), and for the case of EM+MCS, the distance sample rate (N_{DSR}). It has been hypothesized in [4] that these later three parameters are independent of N and M, and we have decided to use the values that were obtained in [4] and [5]; namely, f_s = 44,100 Hz (44.1 kHz), N_f = 1,102 samples (corresponding to a frame duration of 25 ms), and N_{DSR} = 2,000.

In order to optimize the first two parameters, two authors of this paper have subjectively evaluated the similarity of 200 song pairs that were randomly selected from a corpus containing approximately 10,000 songs spanning 40 different genres. Each author has rated each pair of songs using a one to four scale explained in Table 1; half ratings (e.g., 2.5) were also allowed. For the similarity measures KM+EMD and EM+MCS, N was varied from 5 to 25 in steps of 5, and M was varied from 5 to 30 in steps of 5. For the similarity measure AV+EUC, N was taken from the set {3, 4, 5, 8, 10, 15, 20, 23, 30, 40, 50}.

Two-fold cross validation has been used to evaluate each parameter configuration. The 200 song pairs are randomly divided into two disjoint subsets with 100 song pairs each. Similarity measures are computed for the each of the first 100 song pairs; these pairs are then sorted according to their similarities and grouped into ten bins with ten song pairs each. Each bin is then labeled with an average rating, according to the authors, of the ten songs in the bin, rounded to the nearest 0.5. Next, similarity measures are computed for the other 100 song pairs, and each is assigned a rating according to the bin from the first 100 song pairs into which the current song pair would fall. These automatically assigned ratings for the second subset of 100 songs are used to compute the average computer-to-human correlation for the current parameter configuration. The entire process is then repeated, swapping the two subsets of 100 songs, and the two correlations computed for each parameter configuration are averaged together. Correlation has been used as defined in [7].

Rating	Meaning	Description
4	"Extremely Similar"	If a person likes one of the songs, it would be rare they wouldn't like the other.
3	"Similar"	If a person likes one of the songs, it's fairly likely they would like the other.
2	"Not Similar"	Liking the first song does not increase or decrease the chances of liking the other.
1	"Totally Different"	It is highly unlikely that a person would like both songs at the same time.

Table 1: Subjective scale for rating music similarity.

The optimal values of N and M for KM+EMD were N=20 and M=15 leading to a computer-to-human correlation of 0.496. The optimal values of N and M for EM+MCS were N=5 and M=25 leading to a computer-to-human correlation of 0.547. The optimal value of N for AV+EUC was 4 leading to a computer-to-human correlation of 0.484. According to [7], these numbers represent medium to large correlations. Note that the correlation between the two authors was 0.613, a large correlation, and since it is unlikely that an automatic similarity measure would outperform humans, this could be considered a reasonable upper bound on the achievable correlations. For the remainder of the paper, the optimized configurations of the three approaches are referred to as *KM+EMD(20,15)*, *EM+MCS(5,25)* and *AV+EUC(4)*.

5 SIGNAL-BASED MUSIC RECOMMENDATION

Three self-sufficient music recommendation engines have been implemented, each incorporating one of the three optimized music similarity measures. The same corpus of 10,000 songs mentioned in Section 4 serves as the source from which each engine draws its recommendations. Each engine accepts a single music query from a user in the form of a digital file. The song's MFCC features are then extracted, and a representative mixture model or vector is computed from the feature distribution.

To begin the recommendation process, the distance between the query song model and the model of each song in the target music corpus is computed, resulting in a total of approximately 10,000 distances generated for the query. Since the corpus is organized by album, the songs in each album are then arranged in order of least to greatest distance from the query song (i.e., "most timbrally similar" to "least timbrally similar"). The most similar song is then chosen from each album; in this way, we are not allowing two recommendations from the same album. The representative songs from each album are sorted in order of least to greatest distance from the query, and three songs are selected at random from the top 2% of songs in the sorted list. During this process, any song selection bearing the same artist as one of the previous selections is discarded, and random selection is repeated as necessary. The final three song selections are considered to be the recommendations based on the query song.

The authors found it justified to present the user with only one song from each artist according to the reasonable assumption that most artists' songs are, generally speaking, timbrally and stylistically consistent. In the case that many of an artist's songs are computed to be extremely close to a query song, the respective artist would be overrepresented in the resulting recommendations. It suffices to assign the role of "gateway song" to the closest song from such an artist to introduce a user to the artist and their discography, and it gives other possibly relevant songs the chance to find a place in the recommendation set.

6 EXPERIMENTAL DESIGN

We have conducted experiments to compare our recommendation engines to today's leading online music discovery tools (i.e., Pandora, Last.fm, and Allmusic). 15 volunteers not involved with the research were recruited, and each volunteer was requested to submit one song based on which three recommendations would be generated by every system. Each volunteer was instructed to verify that Pandora.com, Last.fm, and Allmusic.com all recognize the title and artist of their chosen song prior to submission. The 15 submitted songs were well distributed in terms of their Allmusic genre classification — one belonged to the *Proto-Punk* genre, one to *Hip-Hop*, one to *Hard Rock*, one to *MPB* (Música Popular Brasileira), one to *Jazz*, one to *Alternative Rock*, one to *Indie Pop*, two to *Punk Rock*, two to *Classical*, and three to *Rock-n-Roll*.

To generate recommendations from Pandora, the title of the volunteer's song was submitted to the Pandora website, and the first three songs returned were used (unless any single artist happened to be repeated, in which case the latter song by the artist would be skipped and the next song by a new artist was used in its place). To generate recommendations from Last.fm, which uses artists as opposed to songs to generate suggestions, the artist of the volunteer's song was submitted and the first three recommended songs (also excluding songs from identical artists) were used. To generate recommendations from Allmusic, three songs were randomly chosen from the same narrow genre as the volunteer's submission (not allowing duplicate artists). As a baseline, we also created a simple engine that randomly chooses three songs from the entire corpus (not allowing duplicate artists). As an upper bound, the first author of this paper suggested three personal recommendations.

The three systems described in Section 5, the three online discover tools, the baseline system, and the first author each generated three recommendations based on every submitted song, so 24 total recommendations were generated for each volunteer. These recommendations were returned to the volunteer in a randomized order, without indicating which recommendation was produced by which method; in

the rare instance that multiple engines would choose the same song, that song would only be included in the list once. Each volunteer was then asked to rate each recommendation on a one to five scale explained in Table 2; half ratings were also allowed.

Rating	Description
5	"A top-notch recommendation"
4	"A good recommendation"
3	"An OK recommendation"
2	"Not a good recommendation"
1	"A very bad recommendation"

Table 2: Subjective scale for rating recommendations.

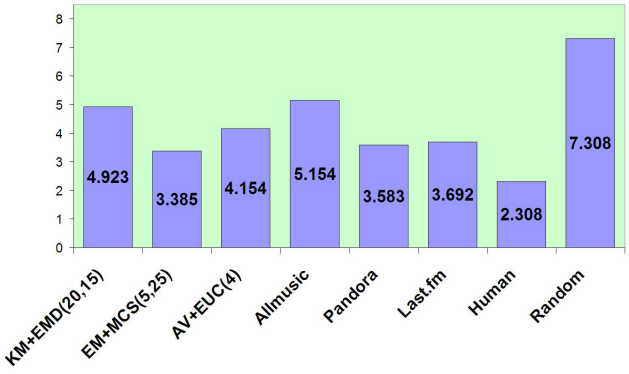

Figure 2: Average rankings of all music recommendation engines, computed across the entire set of volunteers.

7 RESULTS AND EVALUATION

Ultimately, 13 of the 15 volunteers submitted their subjective ratings of the recommendations for their query song. For each volunteer, the performance of each of the eight recommendation engines have been assessed by computing the average of the ratings given to the three songs recommended by that particular engine. These averages have also been used to determine the rank (from first to eighth place) of each engine; engines which tied were assigned equal ranks. To evaluate the performance of all eight recommendation methods across the entire set of volunteers, the ratings and rankings assigned by all volunteers for each method have been averaged; the results are shown in Figure 1 and Figure 2.

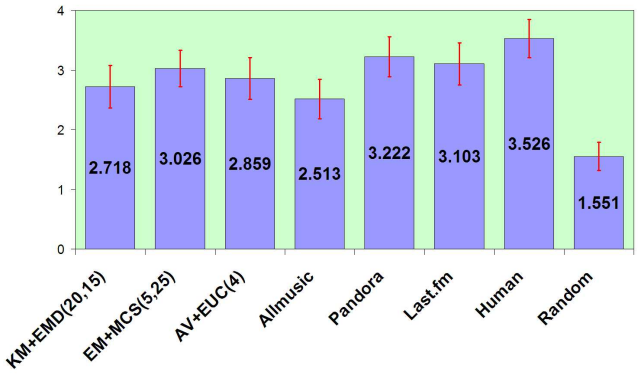

Figure 1: Average ratings for all music recommendation engines, computed across the entire set of volunteers.

It can be seen from Figures 1 and 2 that all of the software-based recommendation engines significantly outperform the baseline random recommender (which received the lowest average rating and worst average rank value), but none perform quite as well as the human recommender (who received the highest average rating and best average rank). According to average ratings, the order of automatic recommendation engines, from best to worst, is Pandora, Last.fm, EM+MCS(5, 25), AV+EUC(4), KM+EMD(20,15), and Allmusic. According to average ranks, the order, from best to worst, is EM+MCS(5, 25), Pandora, Last.fm, AV+EUC(4), KM+EMD(20,15), and Allmusic.

The red bars in Figure 1 represent 95% confidence intervals, assuming normal distrubutions for ratings. Note that the confidence interval for random recommendations has no overlap with that of any other approach; the closest gap is of size approximately 0.4. With the exception of Pandora compared to Allmusic, the confidence intervals of the automatic recommendation engines all overlap, and even for the one exception, the confidence intervals miss each other by only a few hundredths of a point. The top three automated systems - Pandora, Last.fm, and EM+MCS(5, 25) - have confidence intervals that partially overlap with that of the human recommender.

It is not surprising that the two professional online music tools - Pandora and Last.fm - are rated the highest by the 13 volunteers. Note, however, that our signal based recommendation engines trail closely behind, and in particular, EM+MCS(5,25) achieves an average rating only 6% lower than that of Pandora and 2.5% lower than that of Last.fm. In fact, based on average rank, EM+MCS(5,25) performs the best of all the automated systems, indicating that although its average rating is not as high, it beats the professional systems more often than it loses to them. Among the three signal-based recommendation engines, it is not surprising that EM+MCS(5,25) performs the best. The merits of a music similarity measure that utilizes expectation-maximization and Monte Carlo sampling have already been established in the literature [2, 3, 4].

8 CONCLUSIONS

This paper has shown that a signal-based recommendation engine can perform comparably to popular, state-of-the-art commercial music discovery applications when subjected to human evaluation. This fact further highlights the important role that timbre plays in subjective judgment of music similarity; a timbre similarity measure that relies on signal analysis alone appears to be approximately as robust a musical descriptor as musicological analysis or collaborative filtering, and moreso than conventional genre taxonomies.

The results also show that music recommendations given by a fellow human do not satisfy the sensibilities of a music consumer all of the time. Accurately predicting a person's musical tastes is highly dependent on several cultural, sociological, and psychoacoustic factors. Nevertheless, it may be seen that, acting independently, each recommendation engine — whether signal-based or not — produces significantly more accurate recommendations than a baseline random recommender. We can thus say that the particular aspects of music highlighted by each recommendation method are all integral parts of whatever holistic sense of music similarity a person may be said to possess.

Sun Microsystems' Paul Lamere, who is one of the leading researchers in music information retrieval, has dubbed the ideal music discovery engine the "celestial jukebox" [8]. It may be posited this ideal hypothetical engine would be one that somehow combines all the similarity measurement techniques evaluated in this paper, and others as well. Given the positive results discussed in this paper, there is little doubt in the minds of the authors that signal-based music similarity measures will be a *sine qua non* feature of the celestial jukebox of the future.

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MUSIC THUMBNAILER: VISUALIZING MUSICAL PIECES IN THUMBNAIL IMAGES BASED ON ACOUSTIC FEATURES

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ABSTRACT

This paper presents a principled method called *MusicThumbnailer* to transform musical pieces into visual thumbnail images based on acoustic features extracted from their audio signals. These thumbnails can help users immediately guess the musical contents of audio signals without trial listening. This method is consistent in ways that optimize thumbnails according to the characteristics of a target music collection. This means the appropriateness of transformation should be defined to eliminate ad hoc transformation rules. In this paper, we introduce three top-down criteria to improve memorability of thumbnails (generate gradations), deliver information more completely, and distinguish thumbnails more clearly. These criteria are mathematically implemented as minimization of brightness differences of adjacent pixels and maximization of brightness variances within and between thumbnails. The optimized parameters of a modified linear mapping model we assumed are obtained by minimizing a unified cost function based on the three criteria with a steepest descent method. Experimental results indicate that generated thumbnails can provide users with useful hints as to the musical contents of musical pieces.

1 INTRODUCTION

Music recommender systems are increasingly important in online music-distribution services to help users discover their favorite pieces among a huge music collection. For instance, recommender systems based on collaborative filtering [1, 2] recommend musical pieces to the user by taking into account someone else's ratings of those pieces. Content-based filtering systems [3, 4] select musical pieces that are similar to the user's favorites in terms of musical content (acoustic features). Recently, several hybrid systems that integrate these two techniques have been proposed to enable more accurate recommendations [5, 6].

An important problem that has not been resolved is that users cannot immediately grasp the musical contents of recommended pieces after these pieces are listed. Users have to listen to all listed pieces, including those they do not like, to find which pieces are worth listening to. This often prevents users from seamlessly listening to their favorite pieces. Worse still, trial listening is time-consuming because the information of audio signals (temporal media) is not simultaneously delivered to users whereas visual images (spatial media) can be easily skimmed through.

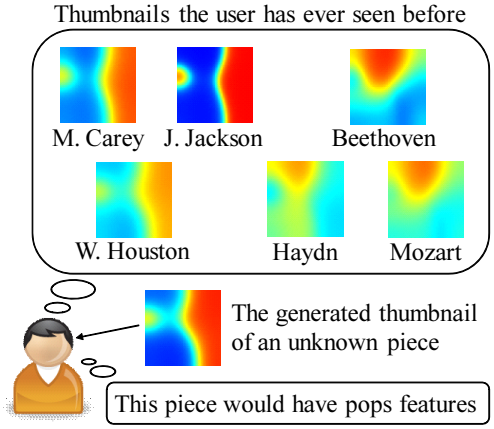

Figure 1. Expected scenario: A user can roughly guess the musical contents of unknown pieces by seeing only thumbnails and without time-consuming trial listening.

To solve this problem, we propose an audio-visual transformation method called *MusicThumbnailer* that generates compact images corresponding to the audio signals of individual pieces. This helps users guess the musical contents of audio signals without trial listening. For example, this method will work well in recommender systems, as shown in Fig. 1. Initially, users only glance at compact thumbnails attached to recommended pieces when they actually listen to these pieces. While accumulating this experience, users will unconsciously associate particular types of thumbnail with their preferred music. Users thus learn to understand the musical meanings of the thumbnails' features. Finally, users will be able to efficiently select audio signals of their desired pieces by using their eyes rather than their ears.

One advantage of our method is that the visual features (colors and patterns) of thumbnails are automatically optimized for a given collection. To achieve this, it is necessary to eliminate ad hoc rules that arbitrarily associate acoustic features with visual features, because these rules lack a principled justification that is consistent for different collections. In this paper, we define some top-down criteria on generated thumbnails from the viewpoint of usability, independently of the characteristics of music collections. We then mathematically represent these criteria in a unified cost function. Audio-visual associations are obtained in a self-organized way so that the cost function is minimized.

The rest of this paper is organized as follows. Section 2 introduces related work. Section 3 explains the principles of our audio-visual transformation method and its implementation. Section 4 reports on our experiment using the RWC Music Database [7]. Section 5 summarizes the key findings of this paper.

2 RELATED WORK

Many visualization methods have been proposed for spatial representation, organization, and browsing of music collections [8–16]. These methods typically locate musical pieces in a two- or three-dimensional space so that musical pieces that have similar musical contents (acoustic features) are arranged close to each other. This enables users to easily understand relationships between musical pieces because similarity in acoustic features can be observed as spatial distance. From a mathematical viewpoint, this kind of visualization can be interpreted as information compression of high-dimensional feature vectors according to some criteria. The self-organizing map (SOM) is often used for this purpose (e.g., Islands of Music [8]).

In a music-playback interface called Musicream [17], individual musical pieces are visualized as colored discs using an ad hoc rule. The disc color (hue and saturation) is determined from the color circle whose circumference and radius correspond to hue and saturation, respectively. Each piece is mapped into the circle according to its feature vector. Principal component analysis (PCA) is used to reduce the dimensionality of acoustic feature vectors to a two-dimensional vector on a plane. Musicovery [18] uses arbitrary rules for associating genres with colors specified in advance.

In the work described in this paper, we aimed to visualize individual pieces, rather than a collection, as compact images (thumbnails) without using ad hoc rules. In general, visual images are represented as super-high-dimensional vectors that contain the color values of numerous pixels. Therefore, our objective was to find an appropriate mapping from a low-dimensional acoustic space (several-tens dim.) to a high-dimensional visual space (several-thousands dim.). A unique feature of this problem lies in this drastic increase in degrees of freedom. To solve such an ill-posed problem by using an optimization method, it is necessary to incorporate some criteria on the appropriateness of the mapping.

3 MUSIC THUMBNAILER

This section describes our method of generating thumbnails of musical pieces based on acoustic features.

3.1 Problem Specification

Given a collection of musical pieces (audio signals) as input data, our objective is to output appropriate thumbnails (visual images) that reflect acoustic features extracted from these pieces. We first prepare some criteria to evaluate the appropriateness of the generated thumbnails as discussed later. In this paper, we focus on generating gray-scale thumbnails as the first step towards obtaining full-color thumbnails in the future. This means we have only to deal with the brightness of pixels contained in each thumbnail.

We first define constant values in advance. Let N be the number of musical pieces. Let S be the number of acoustic features taken into account. Let T be the number of pixels contained in each thumbnail, where T is the product of width, W, and height, H; i.e., $T = WH$.

The input data is given by $X = [\boldsymbol{x}_1 \boldsymbol{x}_2 \cdots \boldsymbol{x}_N]$, which is a collection of feature vectors extracted from all musical pieces. Here, $\boldsymbol{x}_n = (x_{n,1}, \cdots, x_{n,S})^T$ is an S-dimensional feature vector of piece n ($1 \leq n \leq N$), where $x_{n,s}$ ($1 \leq s \leq S$) represents the value of feature s in piece n.

The output data is represented as $Y = [\boldsymbol{y}_1 \boldsymbol{y}_2 \cdots \boldsymbol{y}_N]$, which is a set of brightness vectors of generated thumbnails. $\boldsymbol{y}_n = (y_{n,1,1}, \cdots, y_{n,1,H}, y_{n,2,1}, \cdots, y_{n,2,H}, \cdots, y_{n,W,1}, \cdots, y_{n,W,H})^T$ is a T-dimensional brightness vector of piece n, where $y_{n,w,h}$ ($1 \leq w \leq W, 1 \leq h \leq H$) is the brightness of pixel (w, h) in thumbnail n. The range of $y_{n,w,h}$ is given by $0 < y_{n,w,h} < 1$.

3.2 Top-down Criteria for Principled Transformation

To evaluate the appropriateness of the generated thumbnails, we introduce three top-down criteria from the viewpoint of user friendliness (usability) as follows:

1. *Memorability:* Each thumbnail should be easily remembered by users. The visual pattern is an important factor that affects the ease of remembering thumbnails, as shown in Fig. 2. This is related to how easily users can understand the musical meanings of visual patterns based on their memories. We assume that gradation images are suitable to our purpose.

2. *Informational delivery:* Each thumbnail should provide a large amount of information to users. This is achieved if thumbnail's pixels have a wide variety of brightness, as shown in Fig. 3. This enables users to associate a thumbnail with detailed information about the music content.

3. *Distinguishability:* Users should be able to easily distinguish thumbnail images of different pieces. To enable this, each thumbnail should have a distinctive visual pattern that explicitly reflects the content of music, as shown in Fig. 4. This enables users to efficiently find their favorite pieces based on thumbnails they recognize.

Note that these criteria mention neither specific genres of music nor specific colors of thumbnails.

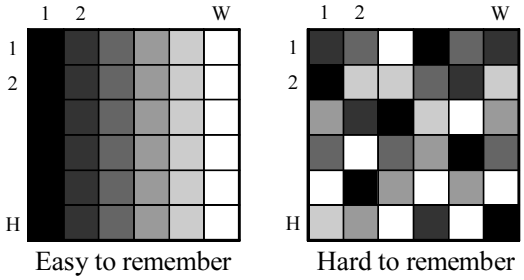

Figure 2. Difference in ease of remembering thumbnails. Both thumbnails have the same histogram of brightness.

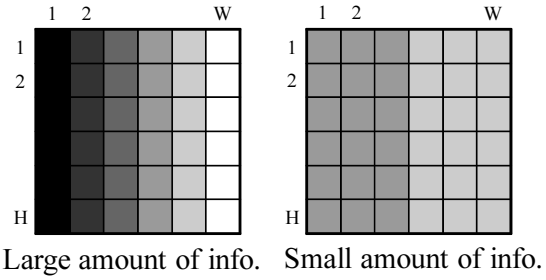

Figure 3. Difference of the amount of information obtained from thumbnails.

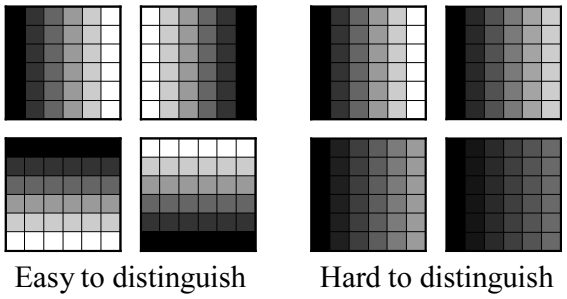

Figure 4. Difference in ease of distinguishing thumbnails of different pieces.

These top-down criteria enable us to design thumbnails in a non-ad-hoc way. In general, system designers tend to define ad hoc rules that directly associate musical contents of musical audio signals with specific colors of visual images (e.g., rock-red, jazz-green, and classic-blue). However, the appropriateness of these arbitrary rules cannot be guaranteed. In contrast, the criteria we proposed only regulate the appropriateness of transformation. The actual colors and patterns of thumbnails are optimized for a given collection in a self-organized way so that the criteria are best satisfied. Therefore, our approach is methodologically and mathematically sound.

A remaining problem here is how to mathematically implement these criteria. In this paper, we will present a simple implementation of the three criteria below.

3.3 Mathematical Formulation

From a mathematical point of view, the objective is to obtain optimal parameters of a mapping model that transforms a S-dimensional space to a T-dimensional space so that a cost function defined according to the three criteria is minimized. This transformation has a high degree of freedom; a higher-dimensional space can always preserve the complete information of an original space. We thus incorporate linear mapping into a mapping model. This is reasonable because linear mapping can strongly limit the degree of freedom in transformation. In addition, musical pieces which are close to each other in an acoustic space are expected to be mapped still close to each other in a visual space. This effect is suitable to achieve the expected scenario shown in Fig. 1. We define the mapping model as

$$Y = \mathit{Sig}(AX), \tag{1}$$

where A is a T-by-S transformation matrix to be optimized, which is given by

$$\begin{bmatrix} \boldsymbol{A}_{1,1}^T \\ \vdots \\ \boldsymbol{A}_{1,H}^T \\ \vdots \\ \boldsymbol{A}_{W,1}^T \\ \vdots \\ \boldsymbol{A}_{W,H}^T \end{bmatrix} = \begin{bmatrix} A_{1,1,1} & A_{1,1,2} & \cdots & A_{1,1,S} \\ \vdots & & & \vdots \\ A_{1,H,1} & A_{1,H,2} & \cdots & A_{1,H,S} \\ \vdots & & & \vdots \\ A_{W,1,1} & A_{W,1,2} & \cdots & A_{W,1,S} \\ \vdots & & & \vdots \\ A_{W,H,1} & A_{W,H,2} & \cdots & A_{W,H,S} \end{bmatrix} \tag{2}$$

and *Sig* is the sigmoid function given by

$$\mathit{Sig}(x) = \frac{1}{1+e^{-x}} \ \ (-\infty < x < \infty). \tag{3}$$

In Eq. (1), we applied the sigmoid function to each cell of matrix AX so that the value of each cell of matrix Y ranges from 0 to 1. Note that the differential of the sigmoid function is given by $\mathit{Sig}'(x) = \mathit{Sig}(x)\,(1 - \mathit{Sig}(x))$.

To evaluate the appropriateness of the mapping model, we define a cost function, C, as

$$C = C_s + \alpha_w C_w + \alpha_b C_b, \tag{4}$$

where C_s, C_w, and C_b are the costs corresponding to the three criteria described in Section 3.2, and α_w and α_b are the weighting parameters. We mathematically define these three costs below.

3.3.1 Minimization of adjacent-pixel distances

To generate gradation images, we focus on a necessary condition that the brightness values of adjacent pixels should be close to each other. One way to mathematically implement this condition is to minimize the differences of the brightness values of adjacent pixels included in each thumbnail.

However, this calculation is not efficient because it would have to be repeated for all thumbnails. To solve this problem, we directly define the cost function, C_s, for the transformation matrix, A, as follows:

$$C_s = \frac{1}{WHS} \sum_{w,h,s} D_{w,h,s}, \tag{5}$$

where $D_{w,h,s}$ is the average of the following eight distances:

$$D_{w,h,s} = \frac{1}{8} \sum_{i,j=\pm 1} \left(A_{w,h,s} - A_{w+i,h+j,s}\right)^2 \tag{6}$$

$$+ \frac{1}{8} \sum_{i=\pm 1} \left(A_{w,h,s} - A_{w+i,h,s}\right)^2 \tag{7}$$

$$+ \frac{1}{8} \sum_{j=\pm 1} \left(A_{w,h,s} - A_{w,h+j,s}\right)^2. \tag{8}$$

3.3.2 Maximization of within-thumbnail variances

To increase the amount of information delivered by a thumbnail, the brightness variance within the thumbnail should be maximized. We formulate this condition for each thumbnail and define the cost function, C_w, based on the average of brightness variances over all thumbnails:

$$C_w = -\frac{1}{N} \sum_{n} \frac{1}{WH} \sum_{w,h} \left(y_{n,w,h} - \bar{y}_n\right)^2, \tag{9}$$

where $\bar{y}_n$ is an average of the brightness values of wh pixels within a thumbnail of piece n, given by

$$\bar{y}_n = \frac{1}{WH} \sum_{w,h} y_{n,w,h}. \tag{10}$$

Note that maximization of within-thumbnail variances is equivalent to minimization of the cost function C_w.

3.3.3 Maximization of between-thumbnail variances

To enable users to clearly distinguish generated thumbnails, the brightness vectors of these thumbnails should be far from each other. A simple way to satisfy this condition is to maximize the brightness variance between all thumbnails, where the between-thumbnail variance is calculated in each position, (w, h). We define the cost function, C_b, based on the average of brightness variances over all positions as

$$C_b = -\frac{1}{WH} \sum_{w,h} \frac{1}{N} \sum_{n} \left(y_{n,w,h} - \bar{y}_{w,h}\right)^2, \tag{11}$$

where $\bar{y}_{w,h}$ is an average of the brightness values of n pixels in the same position (w, h), given by

$$\bar{y}_{w,h} = \frac{1}{N} \sum_{n} y_{n,w,h}. \tag{12}$$

Note that maximization of between-thumbnail variances is equivalent to minimization of the cost function C_b.

3.4 Parameter Optimization

To minimize the total cost function, C, as an optimization method, we use a steepest descent method that iteratively updates the parameters until the cost reduction is converged. The updating formula is given by

$$A_{w,h,s} \leftarrow A_{w,h,s} - \eta \frac{\partial C}{\partial A_{w,h,s}}, \tag{13}$$

where η is a learning parameter ($0 < \eta < 1$) and $\frac{\partial C}{\partial A_{w,h,s}}$ is decomposed into three terms:

$$\frac{\partial C}{\partial A_{w,h,s}} = \frac{\partial C_s}{\partial A_{w,h,s}} + \alpha_w \frac{\partial C_w}{\partial A_{w,h,s}} + \alpha_b \frac{\partial C_b}{\partial A_{w,h,s}}. \tag{14}$$

3.4.1 Derivation of updating formula

We now explain how to derive the updating formula. The first term in Eq. (14) is obtained by

$$\frac{\partial C_s}{\partial A_{w,h,s}} = \frac{2}{WHS} \left(A_{w,h,s} - \bar{A}_{w,h,s}\right), \tag{15}$$

where $\bar{A}_{w,h,s}$ is an average of the values in the vicinity of $A_{w,h,s}$, given by

$$\bar{A}_{w,h,s} = \frac{A_{w,h\pm 1,s} + A_{w\pm 1,h,s} + A_{w\pm 1,h\pm 1,s}}{8}. \tag{16}$$

The second and third terms are calculated as

$$\frac{\partial C_w}{\partial A_{w,h,s}} = \frac{\partial C_w}{\partial y_{n,w,h}} \cdot \frac{\partial y_{n,w,h}}{\partial A_{w,h,s}}, \tag{17}$$

$$\frac{\partial C_b}{\partial A_{w,h,s}} = \frac{\partial C_b}{\partial y_{n,w,h}} \cdot \frac{\partial y_{n,w,h}}{\partial A_{w,h,s}}, \tag{18}$$

where

$$\frac{\partial C_w}{\partial y_{n,w,h}} = -\frac{2}{NWH} \left(y_{n,w,h} - \bar{y}_n\right), \tag{19}$$

$$\frac{\partial C_b}{\partial y_{n,w,h}} = -\frac{2}{NWH} \left(y_{n,w,h} - \bar{y}_{w,h}\right), \tag{20}$$

$$\frac{\partial y_{n,w,h}}{\partial A_{w,h,s}} = \boldsymbol{Sig'}(\boldsymbol{A}_{h,w}^T \boldsymbol{x}_n) x_{n,s} \tag{21}$$

$$= \boldsymbol{Sig}(\boldsymbol{A}_{h,w}^T \boldsymbol{x}_n) \left(1 - \boldsymbol{Sig}(\boldsymbol{A}_{h,w}^T \boldsymbol{x}_n)\right) x_{n,s}. \tag{22}$$

3.4.2 Visual effects of updating formula

Next, we will discuss the visual effects of the three terms of Eq. (14) in the mathematically derived updating formula. Eq. (15) means the first term tries to make transformation coefficients close to their smoothed versions. Eq. (16) corresponds to a visual processing algorithm that is used for smoothing images by using a convolution matrix. Eq. (19) and Eq. (20) mean the second and third terms try to make the brightness value of each pixel far from the within- and between-thumbnail averages. This enhances the dynamic range of each thumbnail and the variety of generated thumbnails. These effects intuitively match our expectation.

4 EXPERIMENT

This section reports on a thumbnail-generation experiment done to evaluate the usefulness of MusicThumbnailer.

4.1 Experimental Conditions

As a music collection, we used the "RWC Music Database: Music Genre" (RWC-MDB-G-2001) [7], which consists of 100 pieces ($N = 100$) in total with three pieces prepared for each of 33 genres and one for a cappella. This database is divided into 10 main genre categories (popular, rock, dance, jazz, Latin, classical, march, world, vocal, and traditional Japanese music) and 33 subcategories (pops, ballad, rock, heavy metal, rap/hip-hop, house, techno, funk, soul/R&B, big band, modern jazz, fusion, bossa nova, samba, reggae, tango, baroque, classic, romantic, modern, brass band, blues, folk, country, gospel, African, Indian, flamenco, chanson, canzone, traditional-style Japanese popular music, Japanese folk music, and ancient Japanese court music).

To extract acoustic features, we used the MARSYAS [19]. We obtained a 42-dimensional feature vector for each piece, which consists of the average and variance of local spectral features (centroid, rolloff, and flux) and zero-crossings across the entire piece (8 dimensions), average and variance of Mel-frequency cepstral coefficients (MFCC) across the entire piece (26 dimensions), and rhythmic content features reflecting periodicity in beat (8 dimensions). We then used PCA to reduce the dimensionality; the 42-dimensional space was transformed into a 20-dimensional space still accounting for 95% of the variance of the original data ($S = 20$).

The size of thumbnails was 50×50 ($W = H = 50, T = 2500$). In this experiment, gray-scale thumbnails were converted to full-color ones according to a lookup table shown in Table 1 (called "Jet" color scheme in MATLAB) from an aesthetic point of view. Note that this is just for convenience because we should essentially eliminate such an ad hoc rule. The values of parameters, α_w, α_b, and η were empirically set to 0.4, 0.4, and 0.1.

4.2 Experimental Results

The experimental results indicate that the generated thumbnails are useful for guessing the musical contents, as shown in Fig. 5. At the main-category level, thumbnails of popular/rock pieces, those of classical (orchestra) pieces, and those of marches seem to be close to each other, respectively. In the dance category, similar thumbnails were obtained in each subcategory, where we can find the similarity between rap/hip-hop and funk. The thumbnails of funk pieces somewhat resemble those of reggae and African pieces across the main category. This corresponds with the fact that funk music has been developed while absorbing the characteristics of African music and reggae. In the categories of Latin, world, and traditional Japanese, we can roughly say that similar thumbnails were obtained in each subcategory. In the vocal category, each subcategory yielded the similar thumbnails, which have especially unique patterns among the database. In the jazz category, however, there were comparatively wide variations in thumbnails of each category. Moreover, fusion pieces tend to reflect and mix the styles of other pieces.

Table 1. Lookup table for converting brightness to RGB.

Brightness	0.00	0.33	0.66	1.00
RGB	(0,0,1)	(0,1,1)	(1,1,0)	(1,0,0)

5 CONCLUSION

We presented an audio-visual transformation method called MusicThumbnailer that generates thumbnail images reflecting acoustic features of audio signals. To achieve principled transformation free from ad hoc rules, we designed three top-down criteria regarding memorability, informational delivery, and distinguishability. These criteria are used to evaluate the appropriateness of generated thumbnails from the viewpoint of usability rather than to associate specific acoustic features with actual colors and patterns of thumbnails. From a mathematical viewpoint, we formulated this problem as a constrained minimization of a cost function based on the three criteria. The experiment showed promising results as to the usefulness of MusicThumbnailer.

Many issues still remain regarding the refinement of our method through subjective experiments. First, we plan to introduce a new criterion to preserve the topological relations of feature vectors in audio-visual transformation. Then, we will improve the mathematical implementation of each criterion and attempt to use a more sophisticated optimization algorithm that can achieve fast convergence while avoiding the local-minimum problem. Several experiments using different features and collections would be important. An interesting application of our method would be to generate a visual effect (a temporal sequence of visual images) that dynamically represents the local musical contents in a musical piece. This can be done by interpreting feature vectors extracted from individual time frames in a musical piece in the same way as for those extracted from individual pieces in a music collection. Such a visualizer will give users a practical overview of structures within a musical piece.

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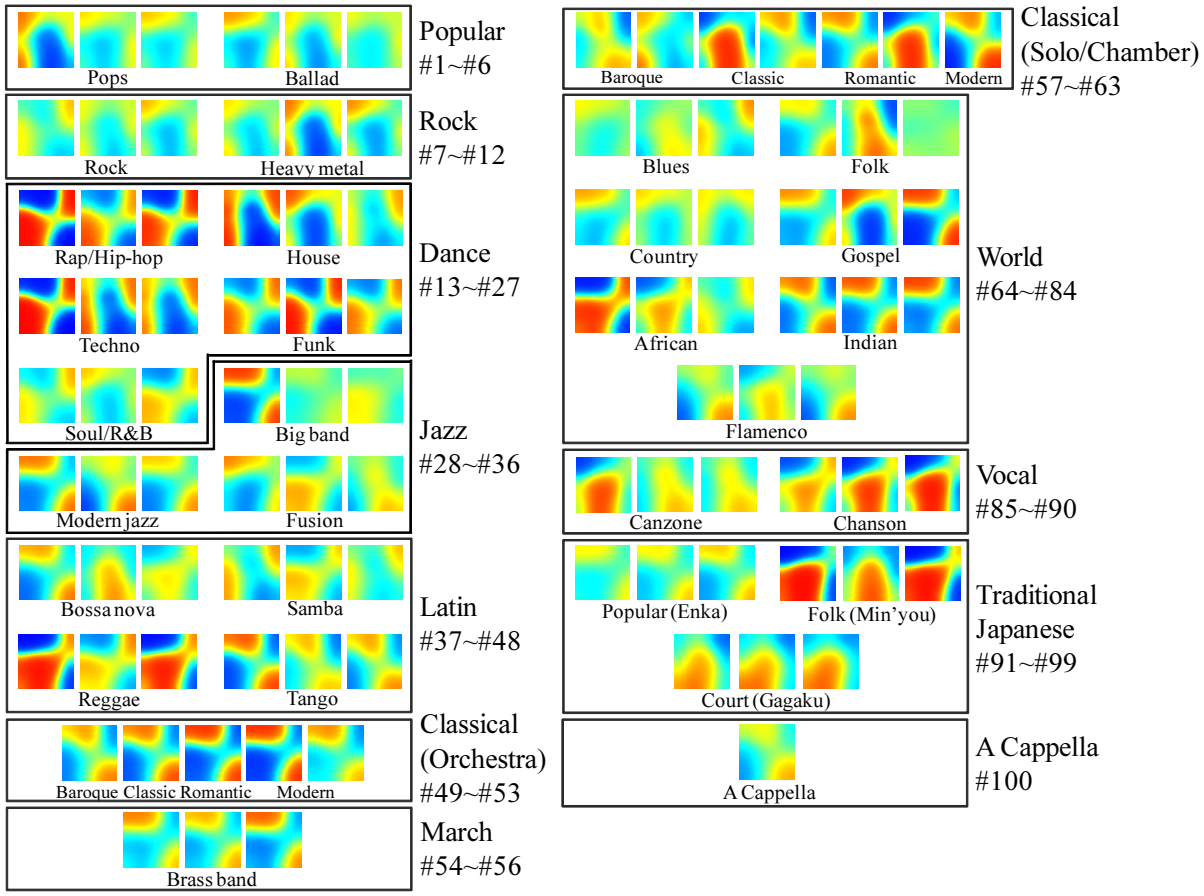

Figure 5. Experimental results of generating thumbnail images of 100 pieces included in RWC-MDB-G-2001 [7].

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DEVELOPMENT OF AN AUTOMATIC MUSIC SELECTION SYSTEM BASED ON RUNNER'S STEP FREQUENCY

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ABSTRACT

This paper presents an automatic music selection system based on runner's step frequency. Recent development of portable music players like iPod has increased the number of those who listen to music while exercising. However, few systems which connect exercises with music selection have been developed. We propose a system that automatically selects music suitable for user's running exercises. Although many parameters can be taken into account, as a first step we focus on runner's step frequency. This system selects music with tempo suitable for runner's step frequency and when runner's step frequency changes, it executes another music selection. The system consists of three modules: step frequency estimation, music selection, and music playing. In the first module, runner's step frequency is estimated from data derived from an acceleration sensor. In the second module, appropriate music is selected based on the estimated step frequency. In the third module, the selected music is played until runner's step frequency changes. In the experiment, subjects ran on a running machine at different paces listening to the music selected by the proposed system. Experimental results show that the system can estimate runner's SPM accurately and on the basis of the estimated SPM it can select music appropriate for users' exercises with more than 85.0% accuracy, and makes running exercises more pleasing.

1 INTRODUCTION

Recently, people can collect massive volumes of digital music data easily due to efficient audio compression techniques like MP3 and online music distribution services. Portable music players like iPod have enabled people to listen to music while doing other things: driving, browsing, or sporting. These circumstances have brought great demand for a system that selects music data suitable for their listening environments from among massive volumes of music data.

There are some commercial products concerning automatic music selection. Portable music players for exercising such as NW-S203F [13] and Nike+iPod Sport Kit [2] have attracted public attention. NW-S203F has an acceleration sensor, with which it can count steps, measure distance traveled, and track calories burned. Nike+iPod Sport Kit consists of a wireless sensor attached to a shoe and a receiver plugged into a portable music player. The sensor sends information such as pace, distance, and calories burned to the receiver, and the music player can store and display this data. These products function not only as portable music players but also as exercise equipment. However, these functions are independent and the information which is obtained while exercising does not influence music selection.

There are also some previous studies on automatic music selection systems using user's information. MPTrain [9, 10, 11] is designed as a mobile and personal system (hardware and software) that users wear while exercising such as walking, jogging or running. MPTrain's hardware includes a set of physiological sensors wirelessly connected to a mobile phone carried by the user. MPTrain does not take any action until about 10 seconds before the end of the current song. It then determines whether the user needs to increase, decrease or keep the running pace, by comparing the user's current heart-rate with the desired heart-rate from the desired workout for that day. Once it has determined the action to take, it searches the user's digital music library (DML) for the optimal song to play. The context-aware mobile music player PersonalSoundtrack described in [4] works with its owner's library to select music in real-time based on a taxonomy of attributes and contextual information derived from an accelerometer connected wirelessly to a laptop carried under the arm. On the basis of user feedback and analysis, a hand-held device is implemented for testing in less constrained mobile scenarios.

These systems select music automatically based on user's listening environments, however, these studies do not investigate in detail how the musical tempo influences user's pleasingness, nor do an objective evaluation to show how accurately their system can estimate runner's step frequency.

In this paper, we propose an automatic music selection system that interacts with user's running exercises. Many parameters can be taken into account, but as a first step we focus on runner's step frequency. We conducted a preliminary experiment to investigate how actual musical tempo

influences runner's pleasingness in the relation with his step frequency. On the basis of the result, we develop a system that automatically selects music with appropriate tempo based on runner's step frequency. This paper is structured as follows: in Section 2 we investigate how actual BPM influences user's pleasingness. In Section 3 we propose an automatic music selection system reflecting runner's step frequency. In Section 4 we present experiments to evaluate the proposed system. Finally in Section 5 we offer concluding remarks and discuss future work.

2 PRELIMINARY EXPERIMENT

In this paper, the developed system which automatically selects music by tempo is described. Tempo is defined as speed at which a musical composition is played [12], and represented by BPM (Beat Per Minute). Some experiments have shown that musical tempo has a great effect on people in the areas of performance of track athletes [3] and on general spatial awareness, arousal and mood [7]. This previous paper [1] studies the influence of the tempo of various pieces of music on SPM (Step Per Minute) of users running while listening to them. It concludes that there is no correlation between tempo and SPM, but there is significant correlation between tempo and subjective rating. These experiments imply that musical tempo is strongly connected with human perception.

In order to investigate how actual BPM influences user's pleasingness, we conducted a preliminary experiment. In the preliminary experiment, six subjects (university students) ran at constant step frequency (150 SPM) listening to two different kinds of music: one is calm and melodious and has a weak beat (Track1), while the other is rhythmical and has a strong beat (Track2). They were composed for the experiment, and BPM of each song was changed at each time :150±5, 10, 35, 40, 45, 70, and 80.

Figure 1 indicates the result of the experiment. Correlation coefficient between the difference between SPM and BPM and evaluated pleasingness of Track1 was -0.589 and that of Track2 was -0.634. This result leads us to the conclusion that people tend to feel pleased when BPM is near their SPM.

3 AUTOMATIC MUSIC SELECTION SYSTEM

3.1 Overview of the proposed system

In this section, we propose a system which selects music by BPM based on the result of the preliminary experiment. Figure 2 indicates an overview of the proposed system. The proposed system consists of three modules: step frequency estimation, music selection, and music playing. First, it estimates runner's step frequency based on data derived from an acceleration sensor (step frequency estimation module).

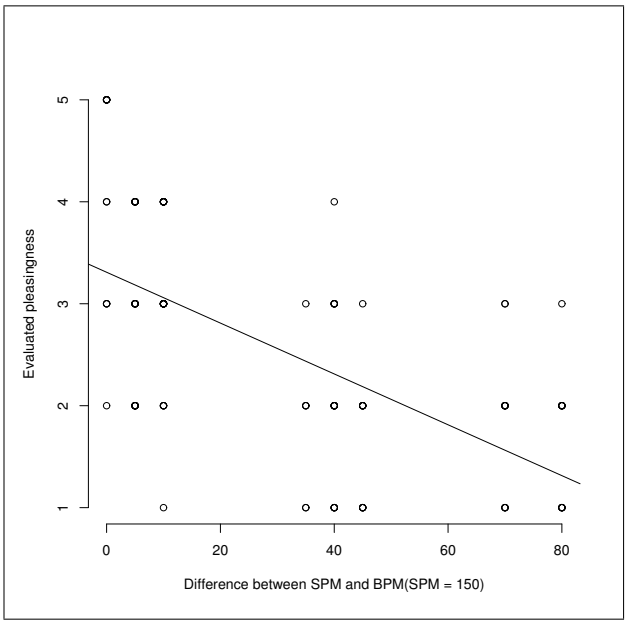

Figure 1. Evaluated pleasingness as a function of difference between SPM and BPM

Next, it selects music based on the estimated step frequency (music selection module). Finally, it plays the selected music (music playing module). We explain each module in detail in the following sections.

3.2 Step Frequency Estimation Module

Our preliminary experimental results show that human step frequency ranges from 70 to 220 SPM. We set sampling frequency of an acceleration sensor at 100Hz, in order to detect steps even if the rate is 600 SPM. We assume that the user takes a step when the acceleration sensor indicates the maximum value. We can calculate an interval between two steps Δt as Equation (1).

$$\Delta t = t_n - t_{n-1} \tag{1}$$

where t_n is the time (sec) the user takes n steps. We can estimate SPM as Equation (2).

$$\text{SPM} \simeq \frac{60}{\Delta t} \tag{2}$$

SPM is updated every time the acceleration sensor indicates the maximum value.

3.3 Music Selection Module

We assume BPM does not change throughout the entire piece and group pieces of music according to its BPM. Each group consists of pieces of music whose BPM are between n and

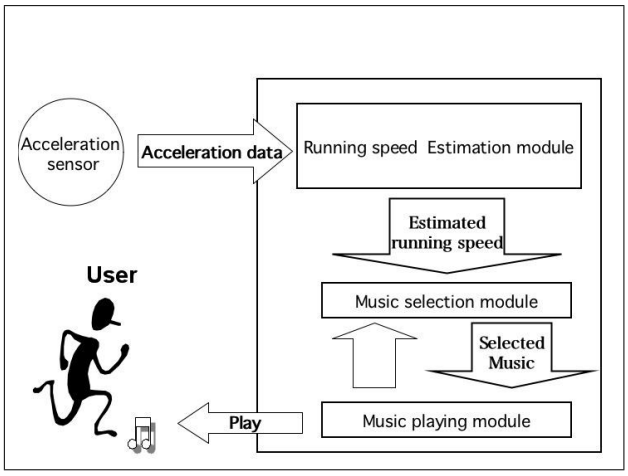

Figure 2. An overview of the proposed system

Figure 3. A mobile acceleration sensor:WAA-001

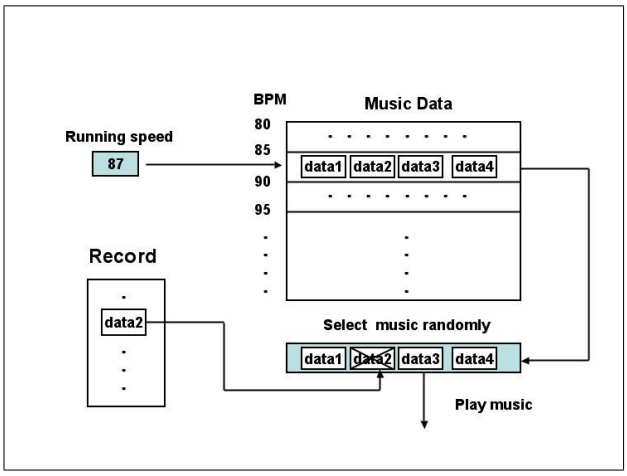

Figure 4. A description of the music selection module

n+4, which is labeled as n. Music selection module searches for the music group whose label n is the closest in value to user's SPM and selects one piece of music at random from the group. The pieces of music selected by the system in the last ten selections are logged, and are not selected again to avoid repetition.

We give an example in Figure 4. The input (step frequency) to the music selection module is 87 and data1, data2, data3 and data4 are music data. In this case, data1, data2, data3 and data4 are candidates because they are included in the music group labeled as 85, which is the closest in value to 87. Data2 is not selected because it is in the record. The system selects one at random from data1, data3 and data4. In this example, data3 is selected and input into the music playing module.

3.4 Music Playing Module

In the music playing module, the music selected in the previous module is played. When runner's step frequency changes, the system executes another music selection using current SPM. This function enables the runner to always listen to music with tempo suitable for his step frequency without manual control. Let P be the estimated SPM, $\overline{P}$ be the average of the last 10 P, $bNow$ be BPM of the playing music, E be the allowable gap between BPM and SPM, $Count$ be a counter, and $uBound$ be upper bound of the times P or $\overline{P}$ can be out of range of $bNow$. Then the algorithm $Play$ is described as follows:

step (S-1):initialize the counter
$Count$ = 0

step (S-2):input runner's step frequency
Input P and $\overline{P}$.

step (S-3):calculate the gap
If $|P - bNow| > E$ and $|\overline{P} - bNow| > E$

then Count++;
else if $|P - bNow| \leq E$ and $|\overline{P} - bNow| \leq E$
then Count−−;
step (S-4):judge the close condition
If $Count > uBound$
then stop music and select another piece of music.
step (S-5):repeat
Back to (S-2).

First, we assign a beginning value "0" to $Count$. When both P and $\overline{P}$ are less than $bNow - E$ or more than $bNow + E$, we increment $Count$. When both P and $\overline{P}$ are more than or equal to $bNow - E$ which is less than or equal to $bNow + E$, we decrement $Count$. Otherwise, $Count$ is not changed. If $Count$ exceeds the upper bound, the system stops the playing music, and executes another music selection using the current SPM (music selection module). Using both P and $\overline{P}$, this module can detect changes of runner's step frequency exactly and avoid frequent changes of music. By default E is 10 and $uBound$ is 20. The user can change these values. In the proposed system, a phase-synchronization[8] method is not implemented, and the next music just cuts in. This method will be implemented in future work.

4 EXPERIMENT

We carried out two experiments in order to confirm availability of the proposed system. In the first experiment, we determined how accurate the estimated SPM was and how long it took for music to change. In the second experiment, five subjects used and evaluated the proposed system. Music data used in the experiments was taken from the RWC Music Database: Popular Music [5, 6]. We used 100 pieces of music and BPM of each piece was calculated manually in advance. The experiment platform was Windows XP professional SP2 with an intel Pentium M 2.13GHz and 2GB memory.

4.1 Experiment to determine the accuracy of SPM estimation

In the first experiment, one subject ran on a running machine (Figure 5) changing his SPM. First, he ran at 80SPM and when music was played, he started to run at 120SPM. When the music changed, he started to run again at 80SPM. He repeated this until the music changed nine times. He also ran at 80 - 160SPM, 120 - 160SPM, and 120 - 200SPM. To determine the accuracy of SPM estimation, we defined precision ratio as Equation (3).

$$precision_k = \frac{\sum_{i=1}^{N} I_k(rSPM_i, eSPM_i)}{N} \qquad (3)$$

where k is the allowable range, N is the number of music selection, $eSPM$ is the estimated SPM, $rSPM$ is the actual SPM, and $I_k(x, y)$ is the function which returns 1 if $|y - x| \leq k$, otherwise 0. We calculated both $precision_5$ and $precision_{10}$.

Table 1 shows the precision ratio of each set. Table 2 shows the average precision ratio at each SPM and how long it took for music to change. One can observe in Table 1 that $Precision_{10}$ is more than 80% in all cases, while $Precision_5$ is too low at 80 - 120SPM, which results from the low precision ratio of 80SPM as indicated in Table 2. This might be due to the difficulty in running just at 80SPM since 80SPM is too slow for most people to run at keeping their step frequency fixed.

Table 2 indicates that the proposed system can select the next music at every SPM within 20 seconds after runner's SPM changes. The lower SPM is, the longer it takes for music to change. The user can shorten the time by changing the value of $uBound$, however, too small $uBound$ may lead to frequent changes of music at higher SPM. In order that it may take the same amount of time for music to change at every SPM, we should make $uBound$ change automatically associated with SPM in future work.

SPM	80-120	80-160	120-160	120-200
$Precision_{10}(\%)$	100.00	100.00	90.91	81.82
$Precision_5(\%)$	50.00	90.00	81.82	81.82

Table 1. The precision ratio of music selection of each set

SPM	80	120	160	200	Over all
$Precision_{10}(\%)$	100.00	87.50	90.00	83.33	90.21
$Precision_5(\%)$	40.00	81.25	90.00	83.33	73.65
Time for music change(sec)	19.90	14.70	11.56	9.32	13.87

Table 2. The precision ratio at each SPM and the time for music change (average)

4.2 Experiment to investigate how the user feels

In the second experiment, we investigated whether the user felt pleased with the music selected by the proposed system. Five subjects (university students) ran on a running machine at different paces listening to the music selected by the proposed system and continued to run until music changed five times. They also ran at different paces listening to the music selected by a random selection system. In the random selection system, music was selected randomly. Music data used in the random selection system was the same as in the proposed system. In both cases, they answered how suitable each music had been for exercises ("suitability") and how pleasant the whole exercise had been ("pleasingness") on a scale of one to five (Table 3, Table 4). Table 5 illustrates a

Evaluation	Criteria
5	very suitable
4	suitable
3	no opinion
2	unsuitable
1	not suitable at all

Table 3. Evaluation criteria for "suitability"

Evaluation	Criteria
5	very pleasing
4	pleasing
3	no opinion
2	boring
1	very boring

Table 4. Evaluation criteria for "pleasingness"

comparison with the random selection system. In both criteria, the proposed system achieved higher scores than the random selection system. The high score of "suitability" indicates that the proposed system can select music suitable for runner's step frequency. The high score of "pleasingness" indicates that the proposed system can give pleasure to people listening to music while running.

	Suitability	Pleasingness
The proposed system	4.3	4.4
Random	2.7	2.6

Table 5. Evaluated "Suitability" and "Pleasingness" (average)

The distribution of the number of pieces of music judged suitable by the users in one experimental session is shown in Table 6. In Table 6, we can find that 86.0% of the music selected by the proposed system was evaluated as suitable for their exercises, while only 35.0% of the music selected by the random selection system was evaluated as suitable. These results suggest that the proposed system can select music suitable for user's running exercises.

Table 7 shows the relation between BPM and "suitability" of the music selected by the proposed system. Table 7 implies that BPM is positively correlated with "suitability". One subject remarked that he had felt comfortable while listening to the music selected by the proposed system especially when running at high SPM. It suggests that the proposed system is especially effective when the user is running at high SPM. This may be partly because at high SPM, phase-shifts are less obtrusive than at low SPM. More detailed research will be needed for a complete understanding of this reason.

Figure 5. A scene of the experiment

Suitability	1	2	3	4	5
Proposed System (%)	0.0	0.0	13.0	43.0	43.0
Random (%)	20.0	33.0	13.0	28.0	7.0

Table 6. The distribution of the number of pieces of music judged suitable by the users

5 CONCLUSION

In this paper, we proposed an automatic music selection system for those who listen to music while running. We focused on runner's step frequency and succeeded in developing a system that automatically selects music appropriate for running exercises by making a correlation between musical tempo and runner's step frequency. Experimental results show that the system can estimate runner's SPM accurately, and on the basis of the estimated SPM it can select music appropriate for users' exercises with more than 85.0% accuracy, and makes running exercises more pleasing. To improve the proposed system, we need to use cross-fading technique to make transition between beats of both pieces of music seamless and have music phase-synchronized to increase user satisfaction. Future work will be dedicated to use other musical elements: musical genre, musical key, meter, timbre, melody, and harmony. And taking this step a bit further, we intend to correlate blood pressure, bodily temperature, or some other health information with various musical elements to make exercises not only amusing but also effective to improve human health.

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Suitability	1	2	3	4	5
Averages of BPM	-	-	115	119	153

Table 7. The relation between BPM and "suitability" of the music selected by the proposed system

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MUSIC GENRE CLASSIFICATION: A MULTILINEAR APPROACH

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ABSTRACT

In this paper, music genre classification is addressed in a multilinear perspective. Inspired by a model of auditory cortical processing, multiscale spectro-temporal modulation features are extracted. Such spectro-temporal modulation features have been successfully used in various content-based audio classification tasks recently, but not yet in music genre classification. Each recording is represented by a third-order feature tensor generated by the auditory model. Thus, the ensemble of recordings is represented by a fourth-order data tensor created by stacking the third-order feature tensors associated to the recordings. To handle large data tensors and derive compact feature vectors suitable for classification, three multilinear subspace techniques are examined, namely the Non-Negative Tensor Factorization (NTF), the High-Order Singular Value Decomposition (HOSVD), and the Multilinear Principal Component Analysis (MPCA). Classification is performed by a Support Vector Machine. Stratified cross-validation tests on the GTZAN dataset and the ISMIR 2004 Genre one demonstrate the advantages of NTF and HOSVD versus MPCA. The best accuracies obtained by the proposed multilinear approach is comparable with those achieved by state-of-the-art music genre classification algorithms.

1 INTRODUCTION

To manage large music collections, tools able to extract useful information about musical pieces directly from audio signals are needed. Such information could include genre, mood, style, and performer [13]. Aucouturier and Pachet [1] indicate that music genre is probably the most popular description of music content. Classifying music recordings into distinguishable genres is an attractive research topic in Music Information Retrieval (MIR) community.

Most of the music genre classification techniques, employ pattern recognition algorithms to classify feature vectors, extracted from short-time recording segments into genres. In general, the features employed for music genre classification are roughly classified into three classes: timbral texture features, rhythmic features, and pitch content features [20]. Commonly used classifiers are Support Vector Machines (SVMs), Nearest-Neighbor (NN) classifiers, or classifiers, which resort to Gaussian Mixture Models, Linear Discriminant Analysis (LDA), etc. Several common audio datasets have been used in experiments to make the reported classification accuracies comparable. Notable results on music genre classification are summarized in Table 1.

Reference	Dataset	Accuracy
Bergstra *et al*. [4]	GTZAN	82.50%
Li *et al*. [12]	GTZAN	78.50%
Lidy *et al*. [14]	GTZAN	76.80%
Benetos *et al*. [3]	GTZAN	75.00%
Holzapfel *et al*. [6]	GTZAN	74.00%
Tzanetakis *et al*. [20]	GTZAN	61.00%
Holzapfel *et al*. [6]	ISMIR2004	83.50%
Pampalk *et al*. [19]	ISMIR2004	82.30%
Lidy *et al*. [13]	ISMIR2004	79.70%
Bergstra *et al*. [4]	MIREX2005	82.34%
Lidy *et al*. [14]	MIREX2007	75.57%
Mandel *et al*. [16]	MIREX2007	75.03%

Table 1. Notable classification accuracies achieved by music genre classification approaches.

Recently, within MIR community, genre has been criticized as being a hopelessly, ambiguous, and inconsistent way to organize and explore music. Consequently users' needs would be better addressed by abandoning it in favor of more generic music similarity-based approaches [17]. From another point of view, end users are more likely to browse and search by genre than either artist similarity or music similarity by recommendation [11]. Furthermore, Aucouturier *et al*. [2] have observed that recent systems, which assess audio similarity using timbre-based features, have failed to offer significant performance gains over early systems and in addition their success rates make them unrealistic for practical use. It is clear that new approaches are needed to make automatic genre classification systems viable in practice. McKay *et al*. [17] argues on the importance of continuing research in automatic music genre classification and

encourages the MIR community to approach the problem in a inter-disciplinary manner.

The novel aspects of this paper are as follows. First, we use bio-inspired multiscale spectro-temporal features for music genre classification. Motivated by the fact that each sound is characterized by slow spectral and temporal modulations and investigations on the auditory system [22], we use the auditory model proposed in [21] in order to extract features that map a given sound to a high-dimensional representation of its spectro-temporal modulations. The auditory high-dimensional representation can be viewed as a high-order tensor that is defined on a high-dimensional space. Note that in the field of multilinear algebra, tensors are considered as the multidimensional equivalent of matrices or vectors [8]. In addition, cortical representations are highly redundant [18]. Therefore, it is reasonable to assume that the tensors are confined into a subspace of an intrinsically low dimension.

Feature extraction or dimensionality reduction thus aims to transform such a high-dimensional representation into a low-dimensional one, while retaining most of the information related to the underlying structure of spectro-temporal modulations. Subspace methods are suitable for the aforementioned goal. Indeed subspace methods, such as Principal Component Analysis (PCA), Linear Discriminant Analysis (LDA), and Non-Negative Matrix Factorization (NMF) have successfully been used in various pattern recognition problems. The just mentioned methods deal only with vectorized data. By vectorizing a typical three-dimensional cortical representation of 6 scales $\times$ 10 rates $\times$ 128 frequency bands results in a vector having dimension equal to 7680. Many pattern classifiers, can not cope with such a high-dimensionality given a small number of training samples. In addition, handling such high-dimensional samples is computationally expensive. Therefore, eigen-analysis or Singular Value Decomposition cannot be easily performed. Despite implementation issues, it is well understood that reshaping a 3D cortical representation into a vector, breaks the natural structure and correlation in the original data. Thus, in order to preserve natural structure and correlation in the original data, dimensionality reduction operating directly on tensors rather than vectors is desirable. State-of-the-art multilinear dimensionality reduction techniques are employed, e.g. Non-Negative Tensor Factorization (NTF) [3], High-Order Singular Value Decomposition (HOSVD) [9], and Multilinear Principal Component Analysis (MPCA) [15] in order to derive compact feature vectors suitable for classification. Classification is performed by an SVM.

Stratified cross-validation tests on two well-known datasets, the GTZAN dataset and the ISMIR2004Genre dataset, demonstrate that the effectiveness of the proposed approach is compared with that of state-of-the-art music genre classification algorithms on the GTZAN dataset, while its accuracy exceeds 80% on the ISMIR2004Genre one.

In Section 2, the computational auditory model and the cortical representation of sound are described. Basic multilinear algebra and multilinear subspace analysis techniques are briefly introduced in Section 3. The application of multilinear subspace analysis to cortical representations is discussed in this section as well. Datasets and experimental results are presented in Section 4. Conclusions are drawn and future research direction are indicated in Section 5.

2 COMPUTATIONAL AUDITORY MODEL AND CORTICAL REPRESENTATION OF SOUND

The computational auditory model, proposed in [21], is inspired by psychoacoustic and neurophysiological investigations in the early and central stages of the human auditory system. An acoustic signal is analyzed by the human auditory model and a four-dimensional representation of sound is obtained, the so-called *cortical representation*. The model consists of two basic stages. The first stage converts the acoustic signal into a neural representation, the so-called *auditory spectrogram*. This representation is a time-frequency distribution along a tonotopic (logarithmic frequency) axis. At the second stage, the spectral and temporal modulation content of the auditory spectrogram is estimated by multiresolution wavelet analysis. For each frame, the multiresolution wavelet analysis is implemented via a bank of two-dimensional Gabor filters, that are selective to spectrotemporal modulation parameters ranging from slow to fast temporal rates (in Hertz) and from narrow to broad spectral scales (in Cycles/Octave). Since, for each frame, the analysis yields a scale-rate-frequency representation, thus the analysis results in a four-dimensional representation of time, frequency, rate, and scale for the entire auditory spectrogram. Mathematical formulation and details about the auditory model and the cortical representation of sound can be found in [18, 21].

Psychophysiological evidence justifies the choice of $scales \in \{0.25,\ 0.5,\ 1, 2, 4, 8\}$ (Cycles / Octave) and positive and negative $rates \in \{\pm 2, \pm 4, \pm 8, \pm 16, \pm 32\}$ (Hz) to represent the spectro-temporal modulation of sound. The cochlear model, employed by the first stage, has 128 filters with 24 filters per octave, covering $5\frac{1}{3}$ octaves along the tonotopic axis. For each sound recording, the extracted four-dimensional cortical representation is averaged along time and the average rate-scale-frequency cortical representation is thus obtained, that is naturally represented by a third-order tensor. Accordingly, the feature tensor $\mathcal{D} \in \mathbb{R}_{+}^{I_1 \times I_2 \times I_3}$, where $I_1 = I_{scales} = 6$, $I_2 = I_{rates} = 10$, and $I_3 = I_{frequencies} = 128$ results. During the analysis, we have used the NSL Matlab toolbox [1] .

[1] http:// www.isr.umd.edu/CAAR/pubs.html

3 MULTILINEAR SUBSPACE ANALYSIS

Recently, extensions of linear subspace analysis methods to handle high-order tensors have been proposed. In this section, three multilinear subspace analysis methods are briefly addressed. To begin with, a short introduction in multilinear algebra is given.

3.1 Multilinear Algebra Basics

In the field of multilinear algebra, tensors are considered as the multidimensional equivalent of matrices (second-order tensors) and vectors (first-order tensors) [8]. Throughout this paper, tensors are denoted by calligraphic letters (e.g. $\mathcal{D}$), matrices by uppercase boldface letters (e.g. $\mathbf{U}$), and vectors by lowercase boldface letters (e.g. $\mathbf{u}$).

A high-order real valued tensor $\mathcal{D}$ of order N is defined over the tensor space $\mathbb{R}^{I_1 \times I_2 \times \ldots \times I_N}$, $I_i \in \mathbb{Z}$, $i = 1, 2, \ldots, N$. Each element of tensor $\mathcal{D}$ is addressed by N indices, $\mathcal{D}_{i_1,i_2,i_3,\ldots,i_N}$. Basic operations can be defined on tensors. The mode-n vectors are column vectors of matrix $\mathbf{D}_{(n)} \in \mathbb{R}^{I_n \times (I_1 \ldots I_{n-1} I_{n+1} \ldots I_N)}$ that results by mode-n unfolding the tensor $\mathcal{D}$.

The symbol $\times_n$ stands for the mode-n product between a tensor and a matrix. The mode-n product of a tensor $\mathcal{D} \in \mathbb{R}^{I_1 \times I_2 \times \ldots \times I_N}$ by a matrix $\mathbf{U} \in \mathbb{R}^{J_n \times I_n}$, denoted by $\mathcal{D} \times_n \mathbf{U}$, can be computed via the matrix multiplication $\mathbf{B}_{(n)} = \mathbf{U}\mathbf{D}_{(n)}$, followed by re-tensorization to undo the mode-n unfolding.

The inner product of two tensors $\mathcal{A}$ and $\mathcal{B} \in \mathbb{R}^{I_1 \times I_2 \times \ldots \times I_N}$ is denoted as $< \mathcal{A}, \mathcal{B} >$. The Frobenius norm of a tensor $\mathcal{A}$ is defined as $||\mathcal{A}||_F = \sqrt{< \mathcal{A}, \mathcal{A} >}$.

An N-order tensor $\mathcal{D}$ has rank 1, when it is decomposed as the Kronecker product of N vectors $\mathbf{u}^{(1)}, \mathbf{u}^{(2)}, \ldots, \mathbf{u}^{(N)}$, i.e. $\mathcal{D} = \mathbf{u}^{(1)} \otimes \mathbf{u}^{(2)} \otimes \ldots \otimes \mathbf{u}^{(N)}$. The rank of an arbitrary N-order tensor $\mathcal{D}$, $R = rank(\mathcal{D})$, is the minimal number of rank-1 tensors that yield $\mathcal{D}$, when linearly combined.

3.2 Non-Negative Tensor Factorization

NTF using Bregman divergences is proposed in [3]. The NTF algorithm is a generalization of the NMF algorithm [10] for N-dimensional tensors. NTF is able to decompose a tensor $\mathcal{D} \in \mathbb{R}_+^{I_1 \times I_2 \times \ldots \times I_N}$ into a sum of k rank-1 tensors:

$$\mathcal{D} = \sum_{j=1}^{k} \mathbf{u}_1^j \otimes \mathbf{u}_2^j \otimes \cdots \otimes \mathbf{u}_N^j \tag{1}$$

where $\mathbf{u}_i^j \in \mathbb{R}_+^{I_i}$. In [3], the NTF is performed by minimizing various Bregman divergences, when auxiliary functions are employed. Furthermore, NTF algorithms using multiplicative update rules, for each specific Bregman divergence are proposed.

In this paper, the NTF algorithm with the Frobenius norm is used. In order to apply the NTF algorithm for an N-order tensor, N matrices $\mathbf{U}^{(i)} \in \mathbb{R}_+^{I_i \times k}$, $i = 1, 2, \ldots, N$ should be created and initialized randomly with non-negative values. Let $*$ stand for the Hadamard product and $\odot$ denote the Khatri-Rao product. The following update rule in matrix form is applied to each $\mathbf{U}^{(i)}$:

$$\mathbf{U}^{(i)} = \tilde{\mathbf{U}}^{(i)} * \frac{\mathbf{D}_{(i)}\mathbf{Z}}{\tilde{\mathbf{U}}^{(i)}\mathbf{Z}^T\mathbf{Z}} \tag{2}$$

where $\mathbf{Z} = \mathbf{U}^{(N)} \odot \ldots \odot \mathbf{U}^{(i+1)} \odot \mathbf{U}^{(i-1)} \odot \ldots \odot \mathbf{U}^{(1)}$ and $\tilde{\mathbf{U}}^{(i)}$ refers to the matrix before updating. It is worth noting, that operators such as Khatri-Rao product preserve the inner structure of data.

3.3 High Order Singular Value Decomposition

HOSVD was proposed by Lathauwer *et al*. in [9] as a generalization of Singular Value Decomposition (SVD) applied to matrix for high-order tensors. Every tensor $\mathcal{D}$ can be expressed as:

$$\mathcal{D} = \mathcal{S} \times_1 \mathbf{U}^{(1)} \times_2 \mathbf{U}^{(2)} \times_3 \ldots \times_N \mathbf{U}^{(N)}. \tag{3}$$

Each $\mathbf{U}^{(n)}$ is a unitary matrix containing the left singular vectors of the mode-n unfolding of tensor $\mathcal{D}$. Tensor $\mathcal{S} \in \mathbb{R}^{I_1 \times I_2 \times \ldots \times I_N}$, known as core tensor, has the properties of all orthogonality and ordering. The HOSVD of a tensor $\mathcal{D}$ according to equation (3) is computed as follows.

1. Compute matrix $\mathbf{U}^{(n)}$ by computing the SVD of the matrix $\mathbf{D}_{(n)}$ and setting $\mathbf{U}^{(n)}$ to be its left singular matrix, $n = 1, 2, \ldots, N$.

2. Solve for the core tensor:

$$\mathcal{S} = \mathcal{D} \times_1 \mathbf{U}^{(1)^T} \times_2 \mathbf{U}^{(2)^T} \times_3 \ldots \times_N \mathbf{U}^{(N)^T}. \tag{4}$$

HOSVD results in a new ordered orthogonal basis for representation of the data in subspaces spanned by each mode of the tensor. Dimensionality reduction in each subspace is obtained by projecting data on principal axes and keeping only the components that correspond to the largest singular values.

3.4 Multilinear Principal Component Analysis

Recently, Lu *et al*. [15] proposed the Multilinear Principal Component Analysis as a multilinear equivalent of PCA. Similarly to PCA, let $\{\mathcal{D}_m \in \mathbb{R}^{I_1 \times I_2 \times \ldots \times I_N}, m = 1, 2, 3, \ldots, M\}$ be a set of M tensor samples. The total scatter of these tensors is defined as:

$$\Psi_{\mathcal{D}} = \sum_{m=1}^{M} ||\mathcal{D}_m - \overline{\mathcal{D}}||_F^2 \tag{5}$$

where $\overline{\mathcal{D}}$ is the mean tensor. MPCA aims to define a multilinear transformation that maps the original tensor space $\mathbb{R}^{I_1 \times I_2 \times \ldots \times I_N}$ onto a tensor subspace $\mathbb{R}^{P_1 \times P_2 \times \ldots \times P_N}$ with $P_n < I_n, n = \{1, 2, \ldots, N\}$ such that most of the variation observed in the original tensor objects is captured, assuming that the variation can be measured by the total tensor scatter. Details and an algorithm for MPCA can be found in [15].

3.5 Multilinear Dimensionality Reduction of Cortical Representations

In each step of a stratified cross-validation test, the dataset used in the experiments (see Section 4) is split into two subsets, one used for training and another used for testing. As mentioned in Section 2, each recording is represented by a third-order feature tensor $\mathcal{D} \in \mathbb{R}^{I_{scales} \times I_{rates} \times I_{frequencies}}$ defining its average cortical representation. Thus, by stacking the third order feature tensors, associated to training recordings, a fourth order data tensor $\mathcal{T} \in \mathbb{R}^{I_{samples} \times I_{scales} \times \times I_{rates} \times I_{frequencies}}$ is obtained, where $samples$ is the number of training set recordings.

3.5.1 Multilinear Dimensionality Reduction by NTF

The resulting data tensor $\mathcal{T}$ is approximated by k rank-1 tensors obtained by NTF. Without loss of generality, NTF approximation is expressed in matrix form as:

$$\begin{aligned} \mathbf{T}_{(1)} &= \mathbf{U}^{(1)}(\mathbf{U}^{(4)} \odot \mathbf{U}^{(3)} \odot \mathbf{U}^{(2)})^T \Longleftrightarrow \\ \mathbf{T}_{(1)}^T &= (\mathbf{U}^{(4)} \odot \mathbf{U}^{(3)} \odot \mathbf{U}^{(2)})\mathbf{U}^{(1)^T} \end{aligned} \quad (6)$$

where $\mathbf{T}_{(1)} \in \mathbb{R}^{samples \times (scales \cdot rate \cdot frequencies)}$ is the mode-1 unfolding of tensor $\mathcal{T}$, $\mathbf{U}^{(1)} \in \mathbb{R}^{samples \times k}$, $\mathbf{U}^{(2)} \in \mathbb{R}^{scales \times k}$, $\mathbf{U}^{(3)} \in \mathbb{R}^{rates \times k}$, and $\mathbf{U}^{(4)} \in \mathbb{R}^{frequencies \times k}$. From (6), it is clear that every column of $\mathbf{T}_{(1)}^T$, i.e. vectorized cortical representation of a sound, is a linear combination of the basis vectors, which span the columns, of the basis matrix $\mathbf{W} = \mathbf{U}^{(4)} \odot \mathbf{U}^{(3)} \odot \mathbf{U}^{(2)}$ with coefficients taken from the columns of coefficient matrix $\mathbf{U}^{(1)^T}$. Performing Gram-Schmidt orthogonalization on basis matrix $\mathbf{W}$, an orthogonal basis matrix $\mathbf{Q}$ can be obtained. The orthogonalized bases span the same space as that of learned bases. The above step was employed, because previous research [5] has shown that orthogonality increases the discriminative power of the projections. Thus, the suitable features for classification are derived from the projection $\tilde{\mathbf{x}}_i = \mathbf{Q}^T \mathbf{d}_i$, where $\mathbf{d}_i$ is the vectorized cortical representation of the ith recording of the dataset.

3.5.2 Multilinear Dimensionality Reduction by HOSVD

The resulting data tensor $\mathcal{T}$ is decomposed to its mode-n singular vectors using the algorithm described in Subsection 3.3. Then, the singular matrices $\mathbf{U}^{(scales)}$, $\mathbf{U}^{(rates)}$, and $\mathbf{U}^{(frequencies)}$ are obtained. These matrices are orthonormal ordered matrices, which contain the subspace of singular vectors. In order to produce a subspace that approximates the original data, each singular matrix is truncated by setting a threshold so as to retain only the desired principal axes for each tensor mode.

Features suitable for classification are derived as follows. Each feature tensor $\mathcal{D}_i$ corresponding to ith recording of the dataset is projected onto the truncated orthonormal axes $\hat{\mathbf{U}}^{(scales)}$, $\hat{\mathbf{U}}^{(rates)}$, and $\hat{\mathbf{U}}^{(frequencies)}$ and a new feature tensor $\hat{\mathcal{X}}_i$ is derived:

$$\hat{\mathcal{X}}_i = \mathcal{D}_i \times_1 \hat{\mathbf{U}}^{(scales)} \times_2 \hat{\mathbf{U}}^{(rates)} \times_3 \hat{\mathbf{U}}^{(frequencies)}. \quad (7)$$

The actual features for classification $\tilde{\mathbf{x}}_i$ are derived from the vectorization of $\hat{\mathcal{X}}_i$.

3.5.3 Multilinear Dimensionality Reduction by MPCA

In a similar manner to HOSVD, a multilinear transformation that maps the original tensor space $\mathbb{R}^{I_1 \times I_2 \times I_3}$ onto a tensor subspace $\mathbb{R}^{P_1 \times P_2 \times P_3}$ with $P_n < I_n$, $n = \{1, 2, 3\}$, such that the subspace captures most of the variation observed in the original tensor objects is obtained by MPCA on $\mathcal{T}$. Features for classification, $\tilde{\mathbf{x}}_i$, are derived from the vectorized form of the projected $\mathcal{D}_i$ using the multilinear transformation obtained by MPCA.

4 EXPERIMENTAL RESULTS

Experiments are performed on two different datasets widely used for music genre classification [4, 6, 12, 13, 19, 20]. The first dataset, abbreviated as GTZAN, consists of following ten genre classes: Blues, Classical, Country, Disco, HipHop, Jazz, Metal, Pop, Reggae, Rock. Each genre class contains 100 audio recordings 30 seconds long. The second dataset, abbreviated as ISMIR 2004 Genre, is from the ISMIR 2004 Genre classification contest and contains 1458 full audio recordings distributed over six genre classes as follows: Classical (640), Electronic (229), JazzBlues (52), MetalPunk (90), RockPop (203), World (244), where the number within parentheses refers to the number of recordings belong to each genre class.

Features are extracted from the cortical representation of sound using the aforementioned multilinear subspace analysis techniques. The value of parameter k in NTF algorithm was set to 150 and 140 for the GTZAN dataset and the ISMIR 2004 Genre one, respectively. The number of retained principal components for each subspace, when HOSVD is employed for feature extraction, is set to be 5 out of 6 for rate, 7 out of 10 for scale, and 12 out of 128 for frequency. The features extracted by MPCA capture 98% of the total variation in each mode. In Figures 1 and 2, the number of retained principal components in each subspace is shown as

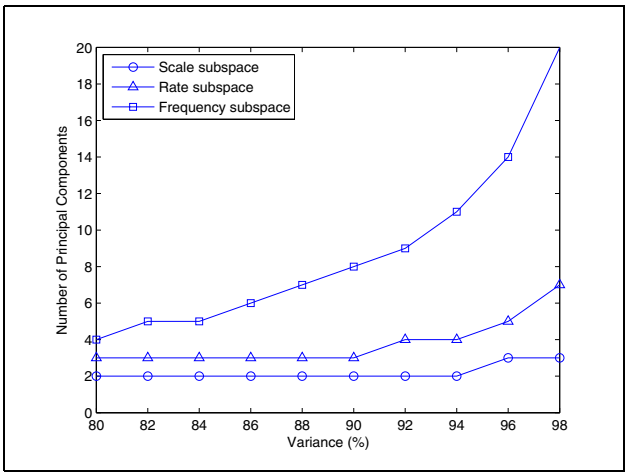

Figure 1. Total number of retained principal components in each subspace (e.g. scale, rate, and frequency) as a function of the portion of variance retained for the GTZAN dataset.

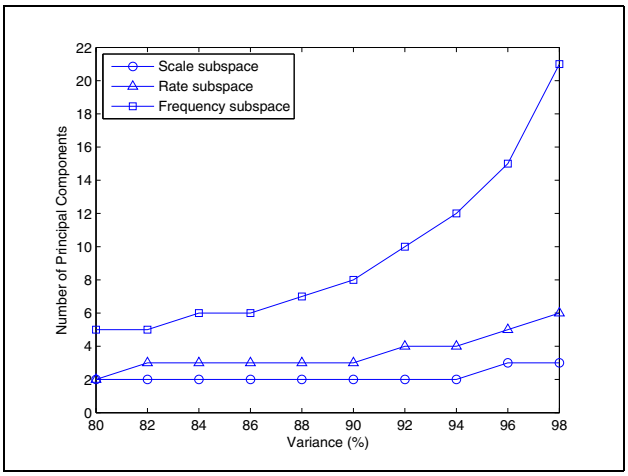

Figure 2. Total number of retained principal components in each subspace (e.g. scale, rate, and frequency) as a function of the portion of variance retained for the ISMIR2004Genre dataset.

a function of the portion of variance retained for the GTZAN and ISMIR2004 Genre dataset, respectively.

Classification was performed by SVM with an RBF kernel. In order to tune the RBF kernel parameters, a grid search algorithm similar to algorithm proposed in [7] was used. Linear and polynomials kernels were also considered, but they achieve poor performance.

The classification accuracies reported in Table 2 are the mean accuracies obtained by 10-fold stratified cross-validation on the full datasets. They are obtained by the various multilinear subspace analysis techniques followed by SVM. The row marked by NTF contains the classification results achieved by features extracted using NTF for both datasets. The row marked by HOSVD contains the classification results achieved by features extracted using HOSVD, while the row marked by MPCA contains the classification results achieved by features extracted using MPCA. The effectiveness of NTF and HOSVD as feature extraction techniques is self-evident in both datasets.

	GTZAN	**ISMIR2004Genre**
NTF	78.20%(3.82)	80.47%(2.26)
HOSVD	77.90% (4.62)	80.95% (3.26)
MPCA	75.01% (4.33)	78.53% (2.76)

Table 2. Classification accuracy on GTZAN and ISMIR2004Genre datasets. The accuracy is calculated by ten-fold stratified cross-validation. The number within parentheses is the corresponding standard deviation.

On the GTZAN dataset, the best classification accuracy outperforms the rates reported by Tzanetakis *et al.* [20] (61.0%), Lidy *et al.* [14] (76.8%), Holzapfel *et al.* [6] (74%), and it is comparable to the rate achieved by Li *et al.* [12] (78.5%). Bergstra *et al.* in [4] reported a classification accuracy equal to 82.5% on the GTZAN database, but they do not disclose details on the experimental setup (e.g. the number of folds).

On the ISMIR2004Genre dataset classification, accuracies achieved for features extracted by NTF and HOSVD are comparable and exceed 80%. It is not possible to compare directly our results with the results obtained by other researchers on this dataset, because of the quite different experimental settings [6, 13, 19].

5 CONCLUSIONS - FUTURE WORK

In this paper, the problem of automatic music genre classification is examined in a multilinear framework. Features have been extracted from the cortical representation of sound using three multilinear subspace analysis techniques. The best classification accuracies reported in this paper are comparable with the best accuracies obtained by other state-of-the-art music genre classification algorithms. The effectiveness of spectro-temporal features obtained by NTF and HOSVD has been demonstrated. It is true, that multilinear techniques applied in straightforward manner, although they provide a more accurate representation to be exploited by the subsequent classifier, do not yield a recognition accuracy much higher than state-of-the-art linear algebra techniques do. Therefore, more effort is required toward addressing the small sample case in the multilinear algebra as well. It is worth noting that the multilinear dimensionality reduction techniques employed in the paper are unsupervised. In the future, supervised multilinear subspace analysis techniques

based on NTF will be developed and tested for the automatic music genre classification.

Finally, in our experiments, we have considered that each song belongs to only one genre class. Obviously, it is realistic to use overlapping class labels for labelling music by style [4]. In general, high-order tensors are structures that are suitable for a such multi-labelling classification problem.

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MCIPA: A MUSIC CONTENT INFORMATION PLAYER AND ANNOTATOR FOR DISCOVERING MUSIC

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ABSTRACT

In this paper, we present a new tool for intra-document browsing of musical pieces. This tool is a multimedia player which represents the content of a musical piece visually. Each type of musical content (structure, chords, downbeats/ beats, notes, events) is associated which a distinct visual representation. The user sees what he/ she is listening too. He can also browse inside the music according to the visual content. For this, each type of visual object has a dedicated feedback, either as an audio-feedback or as a playhead feedback. Content information can be extracted automatically from audio (using signal processing algorithms) or annotated by hand by the user. This multimedia player can also be used as an annotator tool guided by the content.

1 INTRODUCTION

Content description of music based on audio signal analysis has been the subject of many researches over the last few years. Applications such as query over large music collections for specific music characteristics (melody, genre, mood, singer type ...) or search-by-similarity (based on melody, chord progression or timbre ...) have now become possible. However, few works address the problem of using content-information to guide the user during its listening, to allow him/ her to have a better understanding of the musical content, to help him/ her to browse inside a track by the content (intra-document browsing) or to help him comparing several pieces of music visually.

On the other side, the development of music content-extraction algorithms relies for a large part on the availability of annotated music-audio. These annotations are used to learn concepts such as audio structure, audio chords or to check the performances of developed algorithms (multi-pitch or beat-positions estimation algorithms). Tools that allow annotating music audio files in terms of specific music characteristics (chords, beats, structure) are still missing.

In this paper, we present a tool which has been developed in order to allow both user interaction with the music content and annotation of the music content. The paper is organized as follows. In part 2, we give an overview of currently existing music player and audio annotation tools. From this overview we give a set of requirements for our tool. In part 3, we detail the various parts of our tool: its general architecture, the various types of content described and explain how the tool is used for annotation. In part 4, we give technical details about the development of our tool.

2 RELATED WORKS AND REQUIREMENTS

Computer based media players are numerous (iTunes, Windows Media Player, Real Player, Win Amp ...). However, currently none of them propose a visualization of the content of the music track in order to allow browsing of the document by its content. On the opposite, there exist several tools dedicated to the annotation of audio. **"AS (AudioScuplt) Annotation"** [12] is a free software developed by Ircam for Mac-OS-X . The main paradigm of this software is annotation over the visualization of the spectrogram. Many algorithms are integrated into the software such as transient detection, pitch detection ... Annotation is done over the spectrogram using temporal markers or by drawing midi notes over a note-scaled spectrogram. **"Sonic Visualizer"** [5] is an open source software (Linux/ Mac-OS-X/ Windows) developed by Queen Mary University of London. It is also based on annotation over the visualization of either the waveform or the spectrogram. The various annotations can be super-imposed using a set of visual masks. Content analysis is performed using external plug-ins (Vamp format) so that the user can plug-in its favourite content-analyzer. **"CLAM annotator"** [1] is an open source software (Linux/ Mac-OS-X/ Windows) developed by IUA-UPF. It is a framework for developing graphical interfaces and signal analysis algorithms. **"MUCOSA"** [11] also developed by IUA-UPF is an online system for global (a single description assigned to the whole file duration) annotations of music tracks. **"Wavesurfer"** [19] is an open source software (Linux/ Mac-OS-X/ Windows) developed by KTH. It is based on waveform and spectrogram visualization. Some algorithms for energy, pitch or formant estimation are included. Annotation is done by placing markers over the file duration. Some plug-ins for music content analysis (such as beat analysis [9]) have been developed. Wavesurfer has functionalities for browsing by content (markers) over the file duration. **"Praat"** [3] is an open source software (Linux/ Mac-OS-X/ Windows) developed by IPS. It includes many signal analysis algorithms (pitch, formant) but is mostly dedicated to speech analysis. **"Acousmographe"** [10] is a Windows XP software developed by the GRM. It is mainly based on annotation over spectrogram and includes a wide range of graphical tools in order to facilitate annotation reading (specific shape and colour for each annotation). **"Transcriber"** [6] is an open source software (Linux/ Mac-OS-X/ Windows) developed by the French DGA. It is mostly dedicated to the transcription of speech to text. It has functionalities for browsing by content (text) over the file duration. **"Audac-**

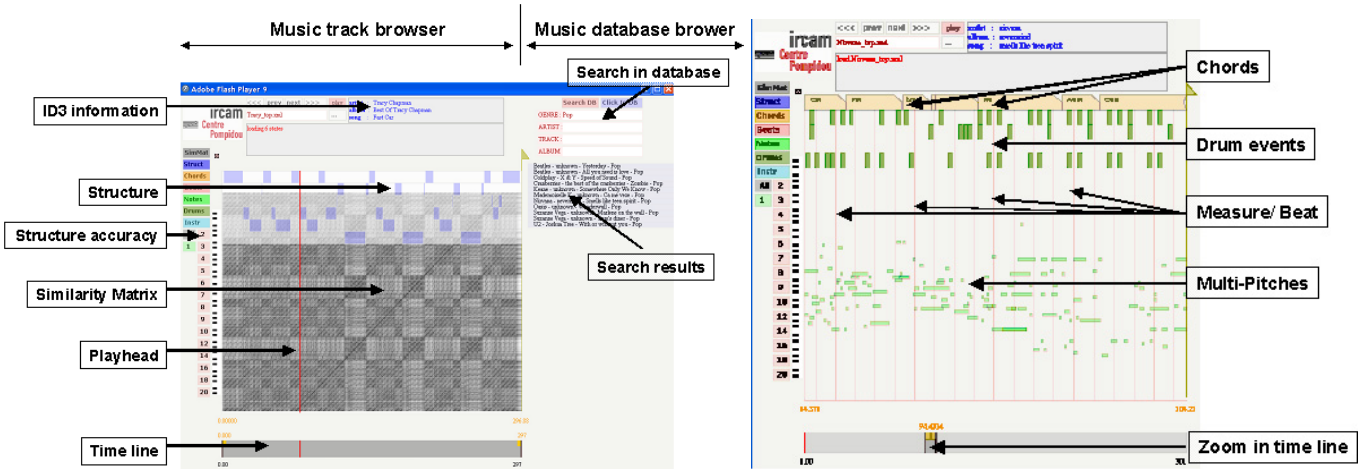


Figure 1. MCIpa interface: [left] large view of the interface with music track browser (similarity matrix and structure representation) and music database browser, [right] detailed view of the music track browser (chord progression, drum events, downbeat/ beat positions and multi-pitch).

ity" [2] [13] is an open source audio multi-tracks recorder with MIDI integration. Because of the audio/ MIDI synchronization it is used for annotation although no specific tools are dedicated to annotation.

2.1 Discussion

According to the previous list, there are currently no tools dedicated specifically to the annotation of music in terms of music content (structure, chords ...). Most of the tools indeed aim to annotate generic audio or speech and propose a representation of the signal (waveform or spectrogram) to help this annotation. A signal representation is useful to annotate a file in low-level audio terms (transients, onsets, fundamental frequency ...) but not to annotate it in high-level music terms (structure, chords ...). Our tool only concentrates on high-level music description. We mean by high-level music description a description that an every-day user can understand (a spectrogram still requires some knowledge about signal processing). Our point of view is that the visualization of one type of music content can help the annotation of another type of music content. Therefore the ground visualization of our interface is the music content description itself (by default it is a similarity matrix representing the global structure of the track) and not a signal representation. However some functionalities present in the list of softwares described above are also interesting in our case: - visual masking and transparency system (AS-Annotation, Sonic Visualizer, Acousmographe), - use of specific colour and shape for each type of annotations (Acousmographe), - separation between the graphical interface and the content-extraction tools (Sonic Visualizer, Wavesurfer), - possibility to quickly browse the files by annotations (Wavesurfer, Transcriber). In our tool, each type of music content is displayed on the same panel and has a specific colour and shape. A large use is made of visual masking and transparency. The content-extraction algorithms are not part of the tool and are provided to it by a set of XML files. These files can be generated by any content-extraction algorithms or annotated by hand. The tool acts either as a media player with music-content visualization or as an annotator of music-content. In the following of this paper, we call **Music Content Information (MCI) object** a description of the music-content that has a specific visual representation, a specific interface-feedback, a dedicated content-extraction algorithm and a dedicated file format.

2.2 Semantic HIFI intra-document player

In the Semantic HIFI project, a first intra-document-player based on the visualization of/ and the interaction with the musical structure has been integrated into the remote-controller of an HIFI system (see Fig. 2). User testing of this interface has been performed and are presented in [4]. The results of this study show that users found the interface "interesting, useful and innovating". However, some weak points were specified by the users: • no possibility to assign a label to each block of the structure, • no possibility to exchange structure annotations among user, • no possibility to assign a colour to each block, • a timeline with tempo annotation would be welcome. These user-comments were taken as the basis for the development of the MCIpa interface.

2.3 Requirements

Generic requirements: The tool must be easy to use, to understand and to install. The same tool is used for both annotation and visualization. The automatic extraction is not part of the tool (the content-description is provided by a set of XML files). This allows the user to use its favourite extraction algorithm and allows the application to remain light. The tool must be cross-platform. The interface should read and play directly the most currently used music formats, mp3 included.

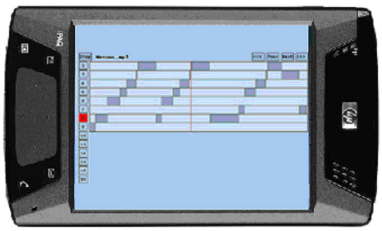


Figure 2. PDA used as a remote-controller for the Semantic-HIFI system: music structure representation.

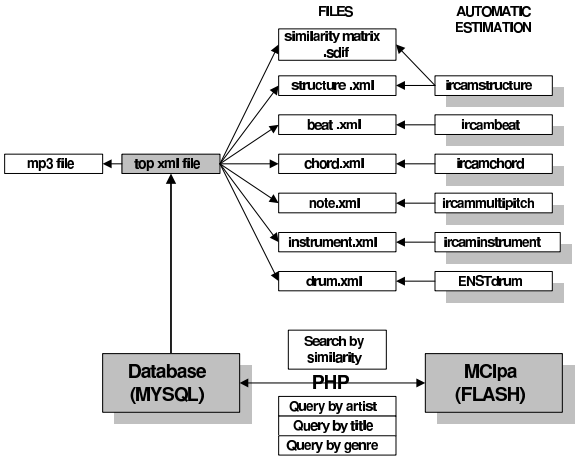


Figure 3. Schema of communication between the MCIpa interface, the database, the audio and content-description files and the corresponding automatic estimation tools.

Interface requirements: The interface must be intuitive. The action must be quick and quickly reachable (keyboard shortcuts are used extensively in order to do that). The main paradigm is "click and listen to what/where you have clicked on". It should be easy for the user to navigate in the representation and to change the type of representations being displayed.

3 DEVELOPED INTERFACE AND FUNCTIONALITIES

The interface is composed of two main parts: the "music database browser" and the "music track browser" (see Fig. 1).

3.1 Music database browser

The music database browser (MDB) allows searching inside a music database by music genre, artist-name, track-name or album-name. The result of the query is displayed as a list of music tracks. Clicking on any item of this list will automatically load the corresponding music track into the "music track browser".

3.2 Music track browser

The music track browser (MTB) represents graphically all Music Content Information of a given track and allows navigation inside it. The top part of the MTB groups the actions for loading directly the file (when not using the "music database browser"), for displaying the editorial metadata (ID3 tags) and for performing the basic playing actions (play, pause, stop, forward/backward).

The middle panel of the MTB represents visually the musical content of the track. Time is represented horizontally. The current playing time is represented as a play-head (red vertical bar) sliding from left to right over the file duration. In this panel are superimposed the various visual objects representing the musical content of the track (the MCI objects). The main paradigm of this panel is "click and listen to what/where you have clicked on". For this, each type of MCI object has a specific behaviour, either as an audio feedback or as a play-head positioning feedback. We have decided to represent all information on the same panel in order to allow comparison between the various MCIs and highlight their musical dependencies.The MCI objects represent the music-content with decreasing temporal scales: similarity matrix, temporal structure, chord progression, music temporal grid (downbeat and beat positions), various notes occurring over time and various sound events occurring over time (musical instruments or percussive sounds). Each type of MCI object has a specific visual representation (shape and colour). They are super-imposed graphically thanks to a masking and transparency system (see Fig. 1 [left] with the transparent structure over the similarity matrix). The user can quickly change the kind of representation during playing using either interface buttons or keyboard shortcuts (showing/masking similarity matrix, structures, chords, beats, notes, sound-events representation).

The bottom part of the MTB represents the time-line of the track. It allows the user to zoom in/ out or to scroll over the track. In this case, the time-line indicates the currently displayed content with a darker rectangle. Auto-scroll functionality is included and can be turned on and off.

3.3 Overall architecture and file formats

The interface directly reads the mp3 format. The descriptions of the various music-contents are stored as a set of XML files. These files are organized in a hierarchical way: a "top" XML file contains a list of pointers to the described mp3 file and to the various content-description files (one file per type of description). The description can • come from manual annotations (either done externally or using the interface itself), • be generated using a set of automatic music-content extraction tools (which are not part of the application). The interface only displays existing data in the XML files (not all of them are mandatory). The XML formats read by MCIpa are described at the following URL [1] . It should be noted that we choose not to use the MPEG-7 Audio XML schema [15] here. This is because, for the kind of music description used here, implementing MPEG-7 is too heavy, morevover MPEG-7 lacks several important music descriptions used here. The proposed XML format has been chosen to be very simple to use and to read. The general schema of the tool is represented in Fig. 3.

[1] http://recherche.ircam.fr/equipes/analyse-synthese/peeters/mcipa/

3.4 Various types of Music Content Information objects

We now describe the various types of Music Content Information (MCI) displayed on the interface, their meaning, their visual representation, their interface-feedback, and the way they can be obtained (by automatic extraction or manual annotation). Table 1 summaries the various MCIs, their representations and corresponding actions.

3.4.1 Similarity matrix

Meaning: The default representation of the music track (representation loaded by default on the background of the MTB) is the similarity matrix [7]. This matrix describes the similarity between the content at the various times of a music track. A point (x,y) in the matrix represents the similarity between the content of the track at time x and time y. *Visual representation:* This matrix is represented as a 2D-image, low similarity values are represented with bright colours, high similarity values with dark colours. Dark lines parallel to the main-diagonal in the matrix represents repetition of segments of times (for example repetition of the same melody). *Extraction and annotations:* This matrix is automatically generated by analyzing the audio signal. It represents the similarities of timbre (instrumentation) and harmony (melodies and chords) as well as their evolution over short periods of time [17]. The similarity matrix cannot be manually generated. *Storage:* The similarity matrix is stored either as an SDIF file or directly as a bitmap file (for faster loading). *Interface feedback:* The user can click anywhere inside the matrix, the playing will starts immediately at the given position. The user can therefore quickly compare several occurrences of a melody by clicking on the various diagonals.

3.4.2 Music structure

Meaning: The music structure represents a music track as a succession of parts (verse, chorus, bridge ...) that can be repeated over time. *Visual representation:* The structure is represented visually as a set of rectangles placed inside various corridors. Each corridor represents a specific type of part; the width of the rectangle inside it represents the time extend of a specific part. The accuracy of the structure (number of corridors used for the decomposition) can be chosen by the user ("structure accuracy" buttons in Fig. 1). The structure can be super-imposed to the similarity matrix in order to allow comparison between the estimated structure and the similarity between the individual times. *Extraction and annotations:* The structure can be estimated automatically from the audio signal using for example the algorithms described in [17] or any algorithm conforming to the proposed XML format definition. The structure can also be manually annotated (or corrected) using the marker system explained in part 3.5. *Storage:* The structure is stored in an XML files describing the decomposition specific to each "structure accuracy" level. *Interface feedback:* Clicking anywhere inside a rectangle starts immediately the playing at the beginning of the corresponding part. The user can therefore quickly compare several parts or several occurrences of the same part. Functionality of forward/ backward by part is also included.

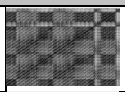
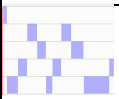
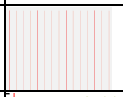
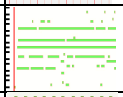
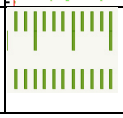
MCI	Graphical representation		User Interaction / Interface feedback
Similarity Matrix	As a 2D image on background		click anywhere inside the image starts playing at the given position
Music Structure	As a part-roll (each type of part is represented on a specific line)		- choose the number of parts used for the subdivision - click inside a part starts playing at the part beginning - forward-backward by parts
Chord progression	As a set of TABs with chord labels		click inside a chord 1) starts playing at the chord beginning 2) plays the corresponding chord prototype
Downbeat/ beat positions	As a set of vertical lines (thick lines for downbeats, thin lines for beats)		Audio click when the play-head crosses a beat marker
Multi-pitch	As a piano-roll (each note-stream is represented by a specific color)		- choose the displayed note channels - click inside a note plays the corresponding note prototype
Sound-events	As a sound-event-roll (each type of sound-event is represented on a specific line)		not yet

Table 1. Music Content Information (MCI) objects.

3.4.3 Chord progression

Meaning: Chord progression denotes the location and duration of each chord of the music track (C Major ... B Major, c minor ... b minor). *Visual representation:* Each chord is represented as a TAB on the top of the middle panel with a specific location and extent. On each TAB is indicated the name of the chord (GM, DM ...). *Extraction and annotations:* The chord progression (chord positions and labels) can be estimated automatically from the audio signal using for example the algorithm described in [16] or any algorithm conforming to the proposed XML format definition. The chords can also be manually annotated (or corrected) using the marker system explained in part 3.5. *Storage:* The chord progression is stored in an XML files. *Interface feedback:* When the user clicks on a specific TAB, the playing starts immediately at the beginning of the corresponding chord. The user can thus quickly skip by chords inside the track. Another feedback, currently under development, is the audio feedback. Clicking on a TAB automatically plays the corresponding prototype chord.

3.4.4 Downbeat and beat positions

Meaning: The downbeats and beats represent the musical temporal grid of a music track. This information is necessary to understand the role of the various musical events existing in the music track. *Visual representation:* Therefore, the downbeats and beats are represented as vertical lines crossing all other types of description. Thick vertical lines are used for downbeats, thin ones for the other beats. *Extraction and annotations:* The beat positions can be estimated automatically from the audio signal using for exam-

ple the algorithm described in [18], the downbeat positions using the one described in [16] or any algorithm conforming to the proposed XML format definition. Beat positions can be corrected manually using the marker system explained in part 3.5. They can also be manually "annotated while listening" by putting "on the fly" markers. This is done using keyboard shortcuts. *Storage:* The downbeat and beat positions are stored on a XML file. *Interface feedback:* The interface provides an audio feedback each time the play-head crosses a beat marker. This allows to "check by listening" the accuracy of the beat marker positions.

3.4.5 Multi-pitch information

Meaning: The multi-pitch information represents the various pitches occurring over time. *Visual representation:* They are represented in a piano-roll way as in most current midi-sequencers. Each line represents a specific pitch; the presence of a note at a given time is indicated by a rectangle at this time and this pitch. Up to 16 different note-channels (based on the 16 midi-channels) can be displayed simultaneously. In order to distinguish the various channels, each note-channel has its own specific colour. The left part of the interface displays the number of existing channels in the description loaded. It allows the user to select which channels to display. *Extraction and annotations:* The multi-pitch information can come from an aligned midi file (converted to the XML format) or can be estimated automatically from the audio signal using for example the algorithm described in [20] or any algorithm conforming to the proposed XML format definition. No multi-pitch annotation or correction is possible in the current version of the interface. *Storage:* The various multi-pitch of the note-channels are stored in an XML files. *Interface feedback:* When the user clicks on a specific note-rectangle, a piano sound at the corresponding pitch is immediately played. This allows, for example, comparing a transcription to the original audio file.

3.4.6 Sound-events information

Meaning: Sound events represent the various sound occurrences over time (instruments, percussive sounds). *Visual representation:* The various occurrences over time of sound events are represented in a "sound-event-roll": each line represents a specific type of sound; the presence of a specific sound is represented by a rectangle at the corresponding line and time. *Extraction and annotations:* This information can be estimated automatically from the audio signal using for example the algorithm described in [8] for drum-event sounds and [14] for instruments or any algorithm conforming to the proposed XML format definition. No sound-events annotation or correction is possible in the current interface. *Storage:* The instrument and drum sound-events are stored in XML files. *Interface feedback:* No interface feedback has been implemented so far.

3.5 Specific actions for annotation

An extension of the music track browser has been realized in order to allow annotation by the user: enter or correct the displayed content description. Indeed, it has been showed

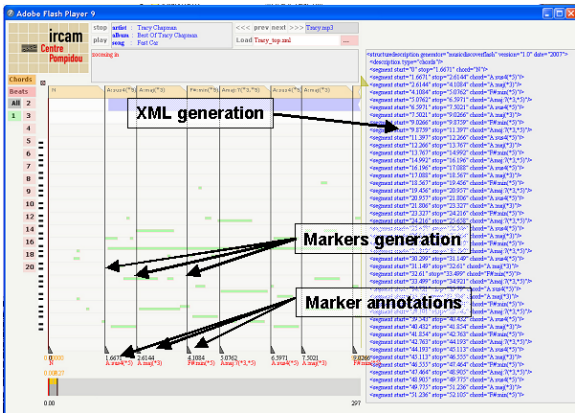


Figure 4. Annotation using the MCIpa interface: markers edition and XML generation

that annotating the content of a music track is greatly facilitated by the knowledge (visualization) of other types of content information. For example, annotating the temporal structure of a track is greatly facilitated by the knowledge (visualization) of the similarity matrix, annotating the chord progression by the knowledge of structure and beat positions ... Annotation is performed using a marker system. The markers can be generated using various methods: • create a marker at the current mouse-cursor position • put "on the fly" markers while listening at the play-head positions (beat annotation) • create a marker at the beginning of the currently selected MCI • generate automatically all the markers corresponding to a specific type of MCI. The markers can then be moved or deleted. Each marker has an associated text label which is editable. If the markers are generated from an existing MCI, the label will be the one of the MCI (for example the one of the chord TAB). An annotation can start from scratch (by generating manually the markers) or by correcting an existing annotation (by generating automatically all the MCI markers and then correcting the markers position and labels). Once finished, the XML code corresponding to the given annotation can be automatically generated and reloaded (see Fig. 4).

4 DEVELOPMENT DETAILS

Graphical interface: The graphical interface has been developed using the FlashDevelop ActionScript 3 environment. The interface is written in the Flash/ ActionScript 3 language, for faster display of numerous items and easy user interaction with the displayed objects. It can easily be run on any platform supporting the Adobe Flash 9 plugin (Linux PC, Windows PC, Mac-OS-X apple computers, portable devices ...). A standalone version can also be run on Mac-OS-X or Windows PC.

Database management: The database management part (music database browser) is calling a PHP script that automatically polls a MySQL database of music titles. This PHP script returns the polled information under the form of a top XML file, which is then read back into the Flash interface.

Any music title has 4 identifiers: title, album, author, music-genre and the path to the top XML file. The top XML file provides the path to the mp3 file and to the various content description XML files (see Fig. 3). When the user asks for a specific title, he just has to click in the list of tracks. The PHP / MySQL / Apache management has been done using the XAMPP environment.

5 CONCLUSION

In this paper, we presented "MCIpa" a Music Content Information player and annotator. The "MCIpa" is a Flash-based (cross-platform) graphical interface that allows intra-document browsing of music track based on content-description. It is also a music-content annotator tool guided by the visualization of other music-contents. The tool represents each type of music-content (similarity matrix, music structure, chord progression, downbeat and beat positions, multi-pitches, sound-events) by a specific Music Content Information object with a specific meaning, visual representation, interface-feedback, extraction-algorithm and XML file format. The XML formats used by MCIpa as well as the software itself are available [2] . We hope this will help people use their favourite content-extraction algorithm with this tool. Because new types of content-description can easily be added to this tool, it could be used as a standard music-content-description player and annotator.

Future works: The paradigms used by MCIpa have been partially tested during the user-testings of the Semantic HIFI intra-document "player" [4]. However they still need to be tested for annotation tasks. Future works will therefore concentrate on that. For this, an experimental protocol for annotation (choice of a set of annotation tasks to be performed and a set of music items) must first be established. Future works will also concentrate on extending the current architecture of MCIpa to a plug-in architecture.

MCIpa usages: We believe that MCIpa can be used for many different purposes since the visual representations used can be easely understand by a large number of people without dedicated signal processing or musical knowledge. MCIpa could be used as a standard media player, for musical education, for comparative musicology (some of the graphical representations of MCIpa are currently used by musicologist at Ircam to compare various interpretations of the same piece; integration into Ircam online mediatheque is also under study), as a musician practicing tools (when playing over music -such as with Aebersold methods- being able to visually locate "theme", "chorus", bars and repetitions is of great help), for research purposes (for quick visualization of results obtained with content-extraction tools) and of course to create annotated training or test-sets.

[2] http://recherche.ircam.fr/equipes/analyse-synthese/peeters/mcipa/

6 ACKNOWLEDGEMENTS

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[4] http://www.quaero.org

A ROBOT SINGER WITH MUSIC RECOGNITION BASED ON REAL-TIME BEAT TRACKING

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ABSTRACT

A robot that can provide an active and enjoyable user interface is one of the most challenging applications for music information processing, because the robot should cope with high-power noises including self voices and motor noises. This paper proposes noise-robust musical beat tracking by using a robot-embedded microphone, and describes its application to a robot singer with music recognition. The proposed beat tracking introduces two key techniques, that is, spectro-temporal pattern matching and echo cancellation. The former realizes robust tempo estimation with a shorter window length, thus, it can quickly adapt to tempo changes. The latter is effective to cancel self periodic noises such as stepping, scatting, and singing. We constructed a robot singer based on the proposed beat tracking for Honda ASIMO. The robot detects a musical beat with its own microphone in a noisy environment. It tries to recognize music based on the detected musical beat. When it successfully recognizes music, it sings while stepping according to the beat. Otherwise, it performs scatting instead of singing because the lyrics are unavailable. Experimental results showed fast adaptation to tempo changes and high robustness in beat tracking even when stepping, scatting and singing.

1 INTRODUCTION

Music information processing draws attention of researchers and industrial people for recent years. Many techniques in music information processing such as music information retrieval are mainly applied to music user interfaces for cellular phones, PDAs and PCs, and various commercial services have been launched[12]. On the other hand, robots like humanoid robots are recently getting popular. They are expected to help us in a daily environment as intelligent physical agents in the future. This means that the robot should not only perform tasks but also make us more enjoyable than PDA or PC based interface. Thus, music is important media for such rich human-robot interaction because music is one of the popular hobbies for humans. This will contribute to MIR society in a sense that robot provides real-world MIR applications. Therefore, we started to apply music information processing to robots. As a first step, we focused on musical beat tracking because it is a basic function to recognize music. However, to be applied to a robot , three issues should be considered for beat tracking as follows:

1. real-time processing by using a robot-embedded microphone,
2. quick adaptation to tempo changes, and
3. high noise-robustness for environmental noises, a robot's own voices and motor noises.

The first issue is crucial to realize natural user interface. A lot of beat-tracking methods have been studied in the field of music information processing [6]. They focus on extraction of complicated beat structures with off-line processing, although there are some exceptions like [5, 8]. Nakadai *et al*. reported the importance of auditory processing by using robots' own ears. They proposed "robot audition" as a new research area[14]. Some robot audition systems which achieved highly noise-robust speech recognition have been reported [7, 18]. However, beat tracking for noisy signals such as robot-noise-contaminated music signals has not been studied so far. The second issue is essential for real-world applications like a robot. For example, in [19], Goto's algorithm was used. It was able to cope with real recording data such as CD music and to apply it to software robot dancer called *Cindy*[3], because it integrates 12 different agents to track musical beats. However, this approach to improve robustness results in insensitivity of tempo changes. This is because a self-correlation-based method requires a longer window to improve noise-robustness, while a short window is necessary to adapt to drastic tempo changes quickly. Thus, they reported that it took around ten seconds to adapt a stepping cycle to tempo changes. Indeed, some probabilistic methods were proposed to cope with tempo changes [10, 2], but these methods tend to require high computational costs and the large amount of memory. Thus, they have difficulty in embedded applications. The last issue is similar to the first one in terms of a noise problem. However, when we consider singing, scatting and stepping functions synchronizing to musical beats, a new problem arises. The noises caused by such functions are periodic because they are generated according to "periodic" beat signals. If the noises and the beats are synchronized, there will be no problem. How-

ever, because scatting/singing is based on estimated beats, entrainment can occur between real and estimated beats in tempo and phase. Thus, it takes a while for them to attain fully synchronization, that is, there is no error between these two beats. This means that the noises affect the performance of beat tracking badly. Scatting and singing cause a much bigger problem than stepping, because the loudspeaker embedded in a robot is usually closer to a robot-embedded microphone than motors and fans. These noises should be suppressed.

In this paper, we proposed a new real-time beat-tracking algorithm by using two techniques to solve the above three issues. One is spectro-temporal pattern matching to realize faster adaptation to tempo changes. The other is noise cancellation based on semi-blind Independent Component Analysis (semi-blind ICA)[16]. We then developed a robot singer with a music recognition function based on proposed real-time beat-tracking for Honda ASIMO. When music is played, the developed robot first detects its beat, secondly recognizes the music based on musical beat information to retrieve the lyrics information from a lyrics database, and finally sings with stepping synchronizing to its musical beat. We evaluated the performance of the proposed beat tracking method in terms of adaptation speed, and noise-robustness through the developed robot system.

2 RELATED WORK IN ROBOTICS

In robotics, music is a hot research topic[1]. Sony exhibited a singing and dancing robot called QRIO. Kosuge *et al*. showed that a robot dancer, MS DanceR, performed social dances with a human partner [17]. Nakazawa *et al*. reported that HRP-2 imitated the spatial trajectories of complex motions of a Japanese traditional folk dance by using a motion capture system [15]. Although these robots performed dances and/or singing, they were programmed in advance without any listening function. Some robots have music listening functions. Kotosaka and Schaal [11] developed a robot that plays drum sessions with a human drummer. Michalowski *et al*. developed a small robot called Keepon which can move its body quickly according to musical beats [13]. Yoshii *et al*. developed a beat tracking robot using Honda ASIMO [19]. This robot was able to detect musical beats by using a real-time beat tracking algorithm [3], and the robot that times its steps to the detected musical beats was demonstrated. These robots worked well only when a music signal is given. However, it is difficult for them to cope with noises such as environmental noises, self voices, and so on. Thus, they have difficulties in singing and scatting that make high power noises.

3 REAL-TIME BEAT TRACKING ALGORITHM

Figure 1 shows an overview of our newly-developed real-time beat tracking algorithm. This algorithm has two input

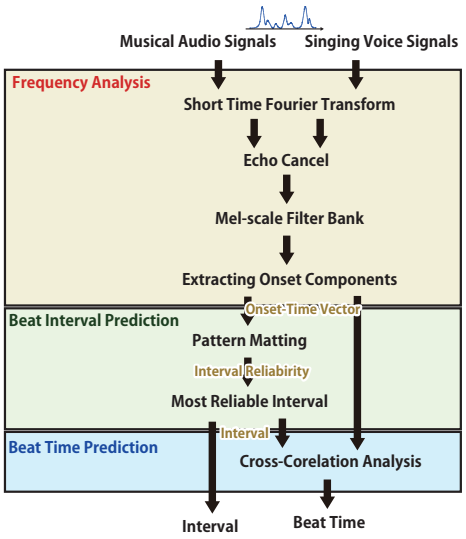

Figure 1. Overview of our real-time beat-tracking

signals. One is a music signal which is usually contaminated by noise sources such as self-noises. The other is a self-noise signal such as a scatting or a singing voice. Because the self-noise is known in advance for the system, pure self-noise can be directly obtained from line-in without using a microphone. The outputs are predicted beat time, and tempo value. It consists of three stages – frequency analysis, beat interval prediction and beat time prediction.

3.1 Frequency Analysis

Spectra are consecutively obtained by applying the short time Fourier transform (STFT) to two input signals sampled at 44.1 kHz. The Hanning window of 4,096 points is used as a window function, and its shift length is 512 points. Echo canceling is, then, applied. It is essential to eliminate self-noises such as singing and scatting voices to improve beat tracking. We introduced semi-blind ICA for echo cancellation[16] which was proposed by our group for self-voice cancellation. We also extended this method to support multi-channel input signals. We used a two-channel version of semi-blind ICA. One channel takes the spectra contaminated by self-noises as an input, and the other channel takes a pure self-noise as an input. The noise-suppressed spectra are sent to Mel-scale Filter Bank. It reduces the number of frequency bins from 2,049 linear frequency bins to 64 mel-scale frequency bins to reduce computational costs in later processes. A frequency bin where a spectral power rapidly increases is detected as an onset candidate at the mel-scale frequency domain. We used the Sobel filter, which is used for visual edge detection, to detect frequency bins only with rapid power increase. Let $d_s(t,f)$ be the spectral power at the t-th time frame and the f-th mel-filter bank bin after the Sobel filtering. An onset belief $d(t,f)$ is estimated by

$$d(t,f) = \begin{cases} d_s(t,f) & \text{if } \mathrm{d_s(t,f)} > 0, \\ 0 & \text{otherwise} \end{cases} \quad (1)$$

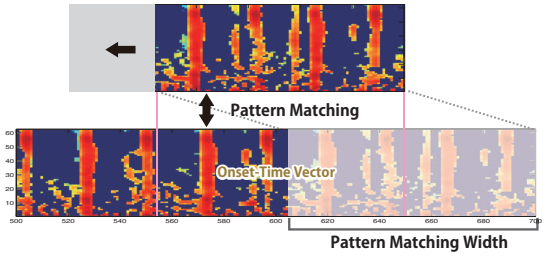

Figure 2. Spectro-Temporal Pattern Matching

where $f = 1, 2, ..., 62$. Thus, a 62-dimensional onset time vector is extracted for each time frame.

3.2 Beat Interval Prediction

To estimate a beat interval defined as the temporal difference between two neighboring beats, spectro-temporal pattern matching is performed by using the onset time vector. As a pattern matching function, we used Normalized Cross-Correlation (NCC) defined by

$$R(t,i) = \frac{\sum_{j=1}^{62}\sum_{k=0}^{P_{width}-1} d(t-k,j)d(t-i-k,j)}{\sqrt{\sum_{j=1}^{62}\sum_{k=0}^{P_{width}-1} d(t-k,j)^2 \cdot \sum_{j=1}^{62}\sum_{k=0}^{P_{width}-1} d(t-i-k,j)^2}} \quad (2)$$

where P_{width} is window length for pattern matching, and i is the shift parameter (Fig. 2).

Frequency-line-based self-correlation is often used for interval estimation. It requires a longer window length for the self-correlation function to improve robustness. This leads to insensitivity to tempo changes. The proposed method uses NCC defined in Eq.(), which corresponds to a kind of *whitening* in signal processing. This improves noise-robustness, even when a window length is as short as 1 sec[1] . Therefore, faster adaptation to tempo changes is achieved. A set of local peaks is, then, extracted by

$$R_p(t,i) = \begin{cases} R(t,i) & \text{if R(t,i}-1) < \text{R(t,i)} < \text{R(t,i}+1), \\ 0 & \text{otherwise.} \end{cases} \quad (3)$$

When two peaks have comparable reliabilities, mis-detection of beat interval occurs. To avoid this mis-detection, beat interval is limited from 61 to 120 M.M.[2] When beat intervals for the first and the second biggest local peaks in $R_p(t,i)$ are I_1 and I_2, beat interval at time t is estimated by

$$I(t) = \begin{cases} 2|I_1 - I_2| & (|I_{n2} - I_1| < \delta \text{ or } |\text{I}_{n2} - \text{I}_2| < \delta) \\ 3|I_1 - I_2| & (|I_{n3} - I_1| < \delta \text{ or } |\text{I}_{n3} - \text{I}_2| < \delta) \\ I_1 & otherwise, \end{cases} \quad (4)$$

$$I_{n2} = 2|I_1 - I_2|, \quad I_{n3} = 3|I_1 - I_2|,$$

[1] This is minimum window length because the lower tempo limit is 60BPM due to a hardware specification of our robot.

[2] Mälzel's Metronome: the number of quarter notes per minute. For example, if the tempo is 60 M.M., the quarter-note length is 1,000 [ms].

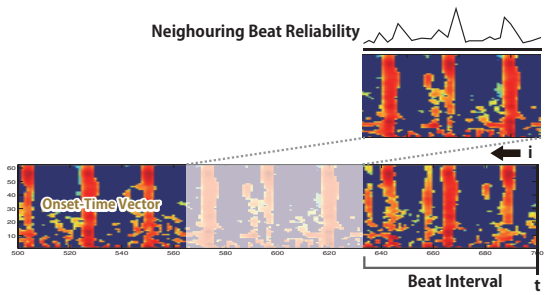

Figure 3. Neighboring Beat Reliability.

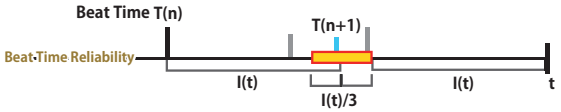

Figure 4. Beat Time Detection

where δ means an error margin parameter. This formulation are defined empirically to avoid mis-estimation such as double and triple tempos.

3.3 Beat Time Prediction

Beat reliability is estimated from two types of reliabilities – neighboring beat reliability and continuous beat reliability. Beat time is predicted according to beat reliability.

Neighboring beat reliability is a reliability on beat existence, and is calculated at the current time and at the previous beat time by using the beat interval shown in Fig. 3. A neighboring beat reliability $S_c(t,i)$ for time $t-i$ at time t is denoted by

$$S_c(t,i) = \begin{cases} \sum_{f=1}^{62} d(t-i,f) + \sum_{f=1}^{62} d(t-i-I(t),f) & (i \le I(t)) \\ 0 & (i > I(t)). \end{cases} \quad (5)$$

Continuous beat reliability is a reliability of a temporal beat sequence. It is calculated from neighboring beat reliabilities.

$$S_r(t,i) = \sum_{m}^{N_{S_r}} S_c(T_p(t,m),i) \quad (6)$$

$$T_p(t,m) = \begin{cases} t - I(t) & (m=0) \\ T_p(t,m-1) - I(T_p(t,m)) & (m \ge 1) \end{cases}$$

where $S_r(t,i)$ denotes continuous beat reliability for time $t-i$ at time t. $T_p(t,m)$ means the m-th previous beat time for time t, and N_{S_r} is the number of beats to calculate continuous beat reliability. This reliability is effective to decide the best beat sequence such as strong beats when multiple beat sequences are detected.

The neighboring beat reliability and the continuous beat reliability are integrated into a beat reliability defined by

$$S(t) = \sum_i (S_c(t-i,i)S_r(t-i,i)). \quad (7)$$

Beat time is then detected. Let the n-th beat time be $T(n)$. When $T(n) \geq t - \frac{3}{4}I(t)$, three-best peaks in $S(t)$ are extracted from $T(n)+\frac{1}{2}I(t)$ to $T(n)+\frac{3}{2}I(t)$. The peak which is closest to $T(n)+I(t)$ is estimated as the next beat time $T(n+1)$ shown in Fig. 4. In case that no peak is found from $T(n)+\frac{2}{3}I(t)$ to $T(n)+\frac{4}{3}I(t)$, $T(n)+I(t)$ is regarded as $T(n+1)$. This beat time detection process was defined empirically. The detected beat time $T(n+1)$ is a past beat, that is, $t > T(n+1)$. To apply beat tracking to scatting or singing, a future beat time T' should be predicted. By using the following extrapolation, a future beat time is predicted.

$$T' = \begin{cases} T_{\text{tmp}} & \text{if } \mathrm{T_{tmp}} \geq \frac{3}{2}\mathrm{I_m(t)} + \mathrm{t} \\ T_{\text{tmp}} + I_m(t) & \text{otherwise.} \end{cases} \quad (8)$$

$$T_{\text{tmp}} = T(m) + I_m(t) + (t - T(m)) - \{(t - T(m)) \mod I_m(t)\}$$

where $I_m(t)$ is a median value of a set of $I(t)$, and $T(m)$ is the latest beat time detected in beat time detection.

4 IMPLEMENTATION OF ROBOT SINGER

Fig. 5 shows the architecture of our robot singer based on the proposed beat-tracking. The system mainly consists of four components – Real-time Beat Tracker, Music Recognizer, Robot Controller, and Humanoid Robot. The Real-time Beat Tracker estimates predicted beat time and a beat interval from a noise-contaminated music signal captured by a robot's microphone as described in Sec. 3. The other three components are described in the following sections. In terms of implementation, Real-time Beat Tracker and Music Recognizer were implemented by C++ on Linux. These components work in real time on a remote PC with Pentium 4. In Robot Controller, scatting and stepping are running the same PC as the above two components, which only singing function is running on Windows PC.

4.1 Specifications of Humanoid Robot

We used Honda ASIMO with a microphone embedded in the head for a singer robot. It has two legs like humans and can stamp its feet on the floor, i.e., perform steps in a stationary location. The step interval is limited to between 1,000 and 2,000 [ms]. If the tempos of musical pieces are between 61 and 120 M.M., The robot records these signals with its own single microphone embedded in the front of the head. It has a loudspeaker for singing at the position of its chest.

4.2 Music Recognizer

Music recognizer consists of two parts – music activity detection and music retrieval. In music activity detection, beat stability is estimated as a ratio of a stable beat period in 3 seconds. When the time difference between the current tempo and the estimated beat interval is within 55 ms, the beat is estimated as stable. When the ratio is higher than 0.8, such a 3-second period is regarded as music. These

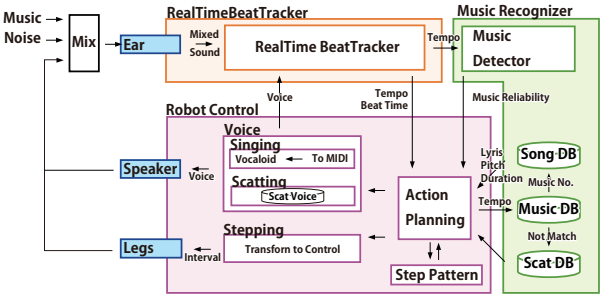

Figure 5. Architecture of a robot singer

thresholds were empirically obtained. Music retrieval returns music ID in the music database by retrieving music which has the closest beat to the estimated one. We simply used tempo information in this retrieval. Practically, when the tempo difference between music and the estimated tempo was within 11 ms, such music was selected. When such music was not found, "unknown music" was returned as a music ID. Music retrieval then obtained the lyrics and notes for the music ID from a song database. In case of unknown music, scatting sounds such as "Zun" and "Cha" were obtained from a scat database in order to utter them instead of singing. Finally, this information was sent to a Robot Control.

4.3 Robot Controller

Robot Controller controls ASIMO to time its steps to musical beats, and to synchronize singing or scatting with the beat. The voices are outputted from a loudspeaker inside ASIMO. The control of stepping is done by using a command via a TCP/IP network.

The stepping function is used to adjust step timings to musical beats only by using a command of specifying a step interval. Because an accurate target value is unavailable, it is theoretically difficult to control a robot even when sophisticated feedback control is used in this case. Thus, we used a simple feedback control to reduce the errors of step timing and interval.

Singing means that a robot sings according to musical beats. Thus, when a music tempo decreases, the robot can sing slowly. As prior information, the melody and lyrics of the music are given to the system as MIDI data. VOCALOID developed by YAMAHA is used as a singing engine. It achieves a singing function with around 200 ms latency. The robot outputs singing voices synchronizing to musical beats by taking such latency into account.

Scatting is used when any appropriate music is not found. Scatting means, here, that a robot outputs sounds according to a beat pattern. In this paper, "zun" was outputted for a strong beat, and "cha" for a weak beat. Since these words have some durations, synchronization between these words and beat time includes some ambiguities. When their correspondence is slightly changed, people easily feel that it is unnatural or the robot is tone deaf. We empirically decided

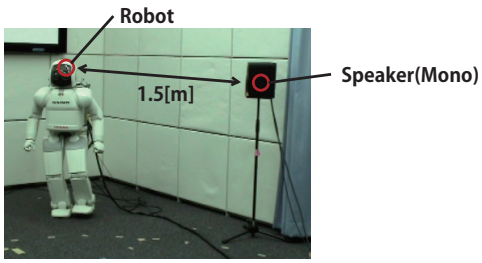

Figure 6. Overview of experimental condition: The system concerning to the robot is completely separated form that concerning to the music playback.

to use onset time of these words, which are detected by onset detection to synchronize with musical beats.

5 EVALUATION

We evaluated our beat tracking using our singer robot in the following three points: 1) adaptation speed to tempo changes, 2) noise-robustness using a beat prediction success rate, 3) music recognition in noisy environments. Three kinds of musical signals were used for these experiments.

T1 *musical signal including tempo changes*
T2 *musical signal with fixed tempo*
T3 *noisy music signals*

For **T1**, we prepared a 4-minute musical signal by selecting three songs (#11, #18, and #62) from the RWC music database (RWC-MDB-P-2001) developed by Goto *et al*. [4]. They include vocals and various instruments as commercial CDs do. Their tempos were 90, 112, and 81 M.M., respectively. We concatenated four 60-s segments that were extracted from the four pieces. For **T2**, we synthesized a musical signal of #62 by using MIDI data. MIDI data provides reference data of beat times. MIDI data is not used as a prior information for tempo and beat time detection. For **T3**, we prepared 10 minute data. The data includes five music signals, i.e., #4, #11, #17, #18 and #29. Each music appears with noises for 20 seconds, and only noise signals are included for the next 20 seconds. For noise data, we used exhibition noise in a booth included in JEIDA-NOISE database. A SNR in T3 was about -4 dB on average.

In every experiment, a loudspeaker was set in a 4 m×7 m room with 0.2 seconds of reverberation time (RT_{20}). The distance between the robot and the speaker was 1.5 m. The musical signals were played from the loudspeaker. This situation is outlined in Fig. 6.

For the first experiment, we used **T1**. The beat tracking delay was measured in five conditions, and was compared with a conventional self correlation based method in [3]. The beat tracking delay was defined as the time difference between when an actual tempo was changed and when the system adapted to the tempo change. Two conditions of the five were the ones with and without scatting when ASIMO was turned off. The other three conditions were the ones without scatting, with scatting and with singing when ASIMO was turned on and performed stepping.

For the second experiment, we used **T2**, and the beat prediction success rate was measured in five conditions. The beat prediction success rate r is defined by

$$r = \frac{100 \cdot N_{\text{success}}}{N_{\text{total}}}. \tag{9}$$

where N_{success} is the number of successfully predicted beats, and N_{total} is the number of total beats. When the error of a predicted beat time is within $\pm 0.35 I(t)$ as defined in [3], it is regarded as successfully predicted. Three conditions of the five are the ones when ASIMO was turned off. One was the condition without scatting and with echo canceling. Another two were the ones with and without canceling while scatting. The other two conditions of the five are the ones with and without echo canceling when ASIMO was turned on with stepping while scatting. For the last experiment, we used **T3**. As metrics for music activity detection, we used precision(P), recall(R), and F-measure(F) defined by

$$P = \frac{C}{N}, \quad R = \frac{C}{A}, \quad F = \frac{2 \cdot P \cdot R}{P + R} \tag{10}$$

where C is a period when music is successfully detected, N is the total period estimated as music, and A is the total music length. As a metric for music retrieval, we used music recognition rate (M) defined by

$$M = \frac{C'}{N} \tag{11}$$

where C' is a period when music was retrieved correctly.

5.1 Results

Table 1 shows the results for the first experiment. This shows that our proposed method adapted to the tempo changes 20 times faster than the conventional one when no voice exists, and it is still 10 times faster than when scatting voices exist. The self-correlation based system failed in beat tracking when singing voices existed, while the proposed was still robust. Table 2 shows the results of the second experiment. "Correct" means the beat tracking system correctly predicted beats, that is, strong beats. "Half-shifted" means that it predicted beats, but weak beats were predicted. This shows self-noises affected beat tracking due to its periodicity, and echo cancel drastically reduced the effect of such self-noises. However, other noises generated by robot's motors and fans were not suppressed explicitly in this paper. Such noise suppression will be attained by using microphone array techniques [18]. Table 3 shows the results of the last experiment. The average precision was around 10 points higher than the average recall. This is caused by the fact that music activity detection is unstable for 2.4 seconds (3 × 0.8) from the beginning of the music due to the

Table 1. Tracking Delay for Tempo Changes (in second)

	ASIMO power off		ASIMO with step		
scatting	off	on	off	on	off
singing	off	off	off	off	on
self-correlation	11.24	29.91	14.66	20.43	N/A
proposed	1.31	1.31	1.29	1.29	1.29

Table 2. Beat Prediction Success Rate

	ASIMO power off			ASIMO power on (with step)	
scatting	off	on		on	
echo cancel	off	on	off	on	off
Correct	95%	97%	68%	95%	64%
Half shifted	5%	1%	40%	4%	40%

Table 3. Music Recognition Result (P: precision, R: recall rate, F: f-measure)

		with noise			clean		
ID	bpm	P (%)	R (%)	F	P (%)	R (%)	F
#4	86	94.7	84.9	0.90	94.8	81.2	0.87
#11	90	74.3	67.3	0.71	96.1	72.1	0.82
#17	97	88.0	83.1	0.85	95.3	81.6	0.88
#29	103	93.4	81.5	0.87	95.9	82.2	0.88
#18	112	89.6	82.8	0.86	95.9	83.2	0.89

3-second window. In #11 and #17, precision was affected by noises. This is because the noise includes a periodic signal between 90 and 97 bpms. *M* was 95.8% for clean data, and 88.5% for noisy data. We can say that music recognition worked well for a small number of songs although using only tempo information. To improve the scalability of music recognition, we will use higher information such as rhythmic features such as [9].

6 CONCLUSIONS

We presented a real-time beat-tracking method for robots which is noise-robust and quickly-adaptable to musical beat changes. The method uses spectro-temporal pattern matching to improve the adaptation speed against tempo changes, and echo canceling based on semi-blind independent component analysis to suppress self periodic noises such as scatting and singing. We showed a singer robot using Honda ASIMO as an application of the proposed beat-tracking. It sings or scats while stepping synchronized to musical beats detected by using robot-embedded microphones, and also it has a simple function to recognize music based on musical beat information. Performance evaluation of the proposed beat tracking method showed high noise-robustness, quick adaptation to tempo changes, high music recognition performance. We believe that the proposed method and its extension will help to realize more active and enjoyable user interface through music, although further evaluation with benchmark datasets is necessary to know its performance precisely. More sophisticated robot motions such as dancing, improvements of robustness of beat tracking, introduction of other music information processing are remaining future work.

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SOCIAL PLAYLISTS AND BOTTLENECK MEASUREMENTS : EXPLOITING MUSICIAN SOCIAL GRAPHS USING CONTENT-BASED DISSIMILARITY AND PAIRWISE MAXIMUM FLOW VALUES

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ABSTRACT

We have sampled the artist social network of Myspace and to it applied the pairwise relational connectivity measure Minimum cut/Maximum flow. These values are then compared to a pairwise acoustic Earth Mover's Distance measure and the relationship is discussed. Further, a means of constructing playlists using the maximum flow value to exploit both the social and acoustic distances is realized.

1 INTRODUCTION

As freely–available audio content continues to become more accessible, listeners require more sophisticated tools to aid them in the discovery and organization of new music that they find enjoyable. This need, along with the recent advent of Internet based social networks and the steady progress of signal based Music Information Retrieval have created an opportunity to exploit both social relationships and acoustic similarity in recommender systems.

Motivated by this, we examine the Myspace artist network. Though there are a number of music oriented social networking websites, Myspace [1] has become the *de facto* standard for web-based music artist promotion. Although exact figures are not made public, recent estimates suggest there are well over 7 million artist pages [2] on Myspace. For the purpose of this paper, *artist* and *artist page* are used interchangeably to refer to the collection of media and social relationships found at a specific Myspace page residing in Myspace's artist subnetwork, where this subnetwork is defined as those Myspace user pages containing the audio player application.

The Myspace social network, like most social networks, is based upon relational links between *friends* designating some kind of association. Further, a Myspace user has a subset of between 8 and 40 *top friends*. While all friends are mutually confirmed, individual users unilaterally select top friends. Additionally, pages by *artists* will usually contain streaming and downloadable media of some kind either audio, video or both.

Social networks of this sort present a way for nearly anyone to distribute their own media and as a direct result, there is an ever larger amount of available music from an ever increasing array of artists. Given this environment of content, how can we best use all of the available information to discover new music? Can both social metadata and content based comparisons be exploited to improve navigation?

To work towards answers to these and related questions, we explore the relationship between the connectivity of pairs of artists on the Myspace top friends network and a measure of acoustic dissimilarity of these artists.

We begin this paper by briefly reviewing graph theoretic network flow analysis and previous work in related topics including musician networks, content-based artist similarity. We go on to explain our methodology including our network sampling method in Section 3.1 and our connectivity analysis techniques in Section 3.2. These connectivity measures are then compared to acoustic artist similarity for the structured network in Section 4 and they are used to construct a social playlist in Section 5. We finish with a discussion of the results and what these results may mean for future work in this space.

2 BACKGROUND

This work uses a combination of complex network theory, network flow analysis and signal-based music analysis. Both disciplines apply intuitively to Music Information Retrieval; however, the two have only recently been applied simultaneously to a single data set [9].

2.1 Complex Networks

Complex network theory deals with the structure of relationships in complex systems. Using the tools of graph theory and statistical mechanics, physicists have developed models

[1] http://myspace.com/

[2] http://scottelkin.com/archive/2007/05/11/MySpace-Statistics.aspx reports as of April 2007 ∼25 million songs, our estimates approximate 3.5 songs/artist, giving ∼7 million artists

and metrics for describing a diverse set of real-world networks – including social networks, academic citation networks, biological protein networks, and the World-Wide Web. In contrast to *simple networks*, all these networks exhibit several unifying characteristics such as small worldness, scale-free degree distributions, and community structure [19]. We briefly introduce below some definitions and concepts that will be used in this work.

A given network G is described by a set of *nodes* N connected by a set of *edges* E. Each edge is defined by the pair of nodes (i, j) it connects. This pair of nodes are *neighbors* via edge $E(i, j)$. If the edges imply directionality, $(i, j) \neq (j, i)$, the network is a *directed network*. Otherwise, it is an *undirected network*. In this paper, all edges are directed unless otherwise stated. In some graphs each edge $E(i, j)$ will have an associated label $w(i, j)$ called the *weight*. This weight is sometimes thought of as the cost of traversing an edge, or an edge's resistance. The number of edges incident to a node i is the *degree* k_i. In a directed network there will be an *indegree* k_i^{in} and an *outdegree* k_i^{out} corresponding to the number of edges pointing into the node and away from the node respectively.

The *degree distribution* $P(k)$ of a graph is the proportion of nodes that have a degree k. The shape of the degree distribution is an important metric for classifying a network – *scale-free* networks have a power-law distribution $P(k) \propto k^{-\gamma}$, while *random* networks have a Poisson distribution [19]. Many real-world networks are approximately scale-free over a wide range of scales. Conceptually, a scale-free distribution indicates the presence of a few very-popular *hubs* that tend to attract more links as the network evolves [19].

2.2 Network Flow Analysis

The basic premise in network flow analysis is to examine a network's nodes as sources and sinks of some kind of *traffic*[2]. Typically, though not exclusively, flow networks are directed, weighted graphs. A simple flow network can be seen in Figure 1. Many useful strategies for determining the density of edge connectivity between sources and sinks can be found in this space[18]. One of the most common among them is the Maximum Flow/Minimum Cut Theorem[8], which is a means of measuring the maximum capacity for fluid to flow between a source node to a sink node or, equivalently, the smallest sum of edge weights that must be *cut* from the network to create exactly two subgraphs, one containing the source node and one containing the sink node. This will be discussed in more detail in Section 3.2. The few examples of network flow type analysis in music deal primarily with constructing playlists using partial solutions to the Traveling Salesman Problem [12] or use exhaustive and explicit metadata[3].

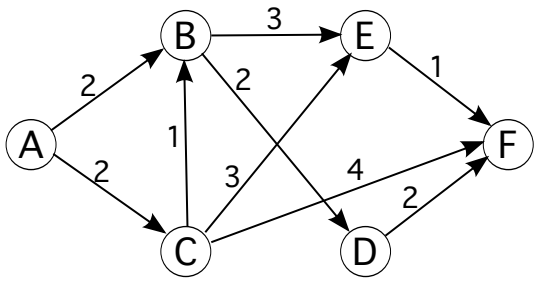

Figure 1. A simple flow network with directed weighted edges. Here the source is node A and the sink is node F.

2.3 Musician Networks

Quite naturally, networks of musicians have been studied in the context of complex network theory – typically viewing the artists as nodes in the network and using either collaboration, influence, or similarity to define network edges. These networks of musicians exhibit many of the properties expected in social networks [7, 10, 21]. However, these studies all examine networks created by experts (*e.g.* All Music Guide [3]) or via algorithmic means (*e.g.* Last.fm [4]) as opposed to the artists themselves, as is seen in Myspace and other similar networks. Networks of music listeners and bipartite networks of listeners and artists have also been studied [4, 14].

2.4 Content-Based Music Analysis

Many methods have been explored for content-based music analysis, attempting to characterizing a music signal by its timbre, harmony, rhythm, or structure. One of the most widely used methods is the application of Mel-frequency cepstral coefficients (MFCC) to the modeling of timbre [16]. In combination with various statistical techniques, MFCCs have been successfully applied to music similarity and genre classification tasks [6, 17, 20]. A common approach for computing timbre-based similarity between two songs or collections of songs creates Gaussian Mixture Models (GMM) describing the MFCCs and comparing the GMMs using a statistical distance measure. Often the Earth Mover's Distance (EMD), a technique first used in computer vision [22], is the distance measure used for this purpose [5, 20]. The EMD algorithm finds the minimum work required to transform one distribution into another.

[3] http://www.allmusic.com/
[4] http://www.lastfm.com/

2.5 Bringing It Together

There has been some recent work attempting to bridge the divide between content-based analysis and human generated metadata. Most of this work [12, 23] focuses on various ways of exploiting the human-generated metadata to filter content prior to, or instead of, conducting content-based analysis, similar to the techniques discussed in Section 2.4, in order to reduce computational load.

3 METHODOLOGY

3.1 Sampling the Social Web

The Myspace social network presents a variety of challenges. For one, the size of the network prohibits analyzing the graph in its entirety, even when considering only the artist pages. Therefore in the work we deal with a sample (of large absolute size) of the network. Also, the Myspace social network is filled with noisy data – plagued by spammers and orphaned accounts. We limit the scope of our sampling in a way that minimizes this noise. And finally, there currently is no interface for easily collecting the network data from Myspace [5]. Our data is collected using web crawling and HTML scraping techniques [6].

3.1.1 Artist Pages

It is important to note we are only concerned with a subset of the Myspace social network – the Myspace *artist* network. Myspace artist pages are different from standard Myspace pages in that they include a distinct audio player application. We use the presence or absence of this player to determine whether or not a given page is an artist page.

A Myspace page will most often include a top friends list. This is a hyperlinked list of other Myspace accounts explicitly specified by the user. The top friends list is limited in length with a maximum length of 40 friends (the default length is 16 friends). In constructing our sampled artist network, we use the top friends list to create a set of directed edges between artists. Only top friends who also have artist pages are added to the sampled network; standard Myspace pages are ignored. We also ignore the remainder of the friends list (*i.e.* friends that are not specified by the user as top friends), assuming these relationships are not as relevant. This reduces the amount of noise in the sampled network but also artificially limits the outdegree of each node. Our sampling is based on the assumption that artists specified as top friends have some meaningful musical connection for the user – whether through collaboration, stylistic similarity, friendship, or artistic influence.

The audio files associated with each artist page in the sampled network are also collected for feature extraction. Cached versions of the audio files are downloaded and audio features are extracted.

3.1.2 Snowball Sampling

There are several network sampling methods; however, for the Myspace artist network, snowball sampling is the most appropriate method [1, 15]. In this method, the sample begins with a seed node (artist page), then the seed node's neighbors (top friends), then the neighbors' neighbors, are added to the sample. This breadth-first sampling is continued until a particular sampling ratio is achieved. Here, we randomly select a seed artist [7] and collect all artist nodes within 6 edges to collect 15,478 nodes. This produces a dataset where no more than six directed top friends links need to be followed to get from the seed artist to any other artist in the dataset. If the size of the Myspace artist network is around 7 million, then this dataset approximates the 0.25% sampling ratio suggested for accurate degree distribution estimation in sampled networks. However, it is insufficient for estimating other topological metrics such as the clustering coefficient and assortativity [13]. Of course, a complete network topology is not our primary concern here.

With snowball sampling there is a tendency to over sample hubs because they have high indegree connectivity and are therefore picked up disproportionately frequently in the breadth-first sampling. This property would reduce the degree distribution exponent γ and produce a heavier tail but preserve the power-law nature of the network [15].

3.2 Minimum Cut/Maximum Flow

We use the Maximum Flow value as a means of determining the number of independent paths from a source node to a sink node. Formally the Maximum Flow/Minimum Cut theorem[8], it is used to calculate the highest weight in the narrowest part of the path from source to sink. The theorem's name comes from the equivalence in the smallest weight of edges that must be removed in order to create two subgraphs which disconnect the source and sink nodes. Further, if the edges in the graph are unweighted, this value is also equivalent to the number of paths from the source to the sink which share no common edges. As this is a mature algorithm there are a number of optimization strategies that have been examined [2, 11].

An example of Maximum Flow can be seen on the network in figure 1. It can been seen that the narrowest point

[5] At time of writing Myspace has recently published a public API, this may allow future work to occur without the need for html scraping, which would greatly decrease the compute time required for graph generation.

[6] Myspace scraping is done using tools from the MyPySpace project available at `http://mypyspace.sorceforge.net`

[7] The artist is *Karna Zoo*, Myspace url: `http://www.myspace.com/index.cfm?fuseaction=user.viewProfile&friendID=134901208`

from node A to node F is $E(a,b)$ and $E(a,c)$. The maximum flow can simply be found via Equation 1.

$$M = \sum m(i,j) \quad (1)$$

Where $m(i,j)$ is the magnitude of each edge in the minimum cut set.

In our Myspace top friends graph, the maximum flow is measured on the unweighted directed graph from the source artist node to the sink artist node.

4 CONNECTED SOUND

4.1 Experiment

We calculate the maximum flow value, using the snowball sample entry point as the fixed source against every other node in turn as a sink, yielding the number of edges connecting each sink node to the entry point node at the narrowest point of connection. The acoustic distances can then be compared to these maximum flow values.

4.1.1 Signal-based analysis

Cepstral coefficients are extracted from each audio signal using a Hamming window on 8192 sample FFT windows with 4096 sample overlap. For each artist node a GMM is built from the concatenation of MFCC frames for all songs found on each artist's Myspace page (the mean number of songs per artist is 3.5). We calculate the Earth Mover's Distance between the GMMs corresponding to each source sink pair in the sample. All MFCCs are created with the fftExtract tool [8] .

4.1.2 Random Networks

In order to better understand a result from analysis of our Myspace sample, a baseline for comparison must be used. To that end, random permutations of the node locations are examined. In order to preserve the overall topology present in the network, this randomization is performed by randomizing the artist label and associated music attached to a given node on the network. This is done ten fold, creating a solid baseline to test the null hypothesis that the underlining community structure is not responsible for any correlation between maximum flow values and Earth Mover's Distance.

4.2 Result

The results of the first experiment show no simple relationship between the sampled netwrok and the randomized network. This can be seen in Table 1 and in Figures 2 and 3. There is an increase in the median EMD for the less well connected (*i.e.* lower maximum flow value) node pairs in the Myspace sample graph, though this is not significant enough to indicate a correlation, while the randomized permutations are near flat. While the median EMD of the artist pairs with a maximum flow of 10 is appreciably higher than all other in the randomized graph, this is likely related to the relatively large size of this group. Perhaps the easiest way to examine the relationship between the sampled graph and randomized one is through the deltas of each group's median from the entire dataset median. This data is shown in the second and forth colum in Table 1 and Figure 4. Further, the Kruskal-Wallis one-way ANOVA results for both the sample graph and averaged across the 10 fold permutations are shown in Table 2.

Max Flow	median	deviation	randomized	deviation
1	40.80	1.26	39.10	−0.43
2	45.30	5.76	38.34	−1.19
3	38.18	−1.35	38.87	−0.66
4	38.21	−1.32	38.64	−0.89
5	40.00	0.47	39.11	−0.42
6	41.77	2.25	39.02	−0.51
7	39.94	0.41	39.24	−0.29
8	39.38	−0.15	38.76	−0.77
9	38.50	−1.03	38.87	−0.66
10	39.07	−0.46	40.85	1.32

Table 1. Node pairs average EMD values grouped by actual minimum cut values and randomized minimum cut values, shown with deviations from the global median of 39.53.

	H-value	P-value
From sample	12.46	0.19
Random permutations	9.11	0.43

Table 2. The Kruskal-Wallis one-way ANOVA test results of EMD against maximum flow for both the sampled graph and it's random permutations. The H-values are drawn from a chi-square distribution with 10 degrees of freedom.

5 THE MAX FLOW PLAYLIST

In order to build playlists using both acoustic and social network data, we use the Earth Mover's Distance between each pair of neighbors as weights on the Myspace sample network. Two artists are then selected, a starting artist as the source node and a final artist as the sink node. One or more paths are then found through the graph via the maximum flow value, generating the list and order of artists for the playlist. The song used is the most popular at the time of the page scrape. In this way playlists are constructed that are both influenced by timbre similarity and bound by so-

[8] source code at *http://omras2.doc.gold.ac.uk/software/fftextract/*

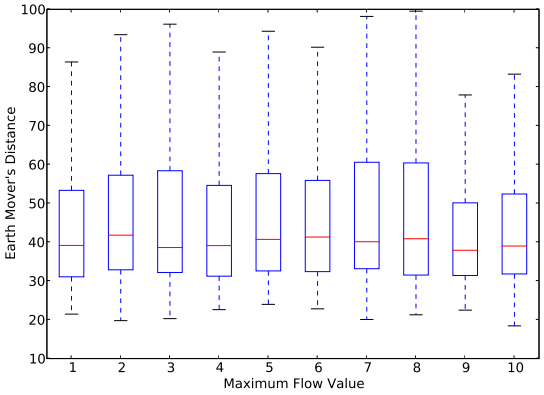

Figure 2. The box and whisker plot showing the distribution of EMD grouped by maximum flow value between artists on the Myspace social graph.

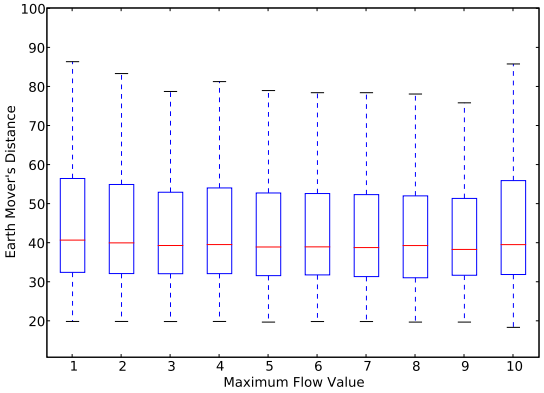

Figure 3. The box and whisker plot showing the distribution of EMD grouped by maximum flow value between artists on the randomized graph, maintaining the original edge structure.

cial context, regardless of any relationship found between these two spaces found via the work discussed in Section 4. Playlists generated using this technique were informally auditioned, and were found to be reasonable on that basis.

6 DISCUSSION AND FUTURE WORK

While an inverse relationship between Earth Mover's Distance and the maximum flow value might be expected on the basis of the conventional wisdom that a community of artists tend to be somehow aurally similar, this does not appear to be strictly the case. The evidence, at least in this sample set, does not support this relationship, though it doesn't disprove it either. However, based upon the difference in result from the Kruskal-Wallis one-way ANOVA test and simple observation of the deviation from the global median the maximum flow values and Earth Mover's Distances do seem affected by the artist created social links, though it is not a simple relationship and describing it precisely is difficult. This seems to suggest that there may be no noticeable correlation between density of connection (maximum flow values) and acoustic similarity (GMMs compared with EMD), at least across an entire sample.

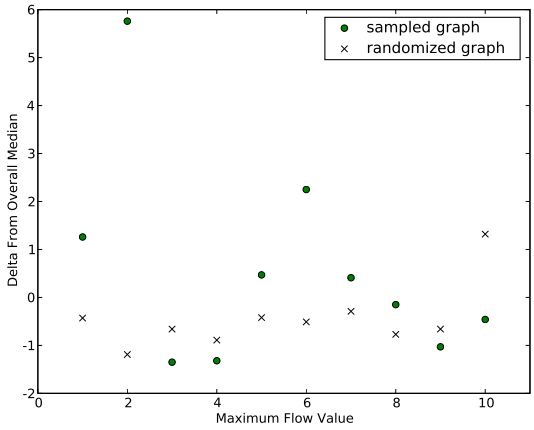

Figure 4. The deltas from the global median for each maximum flow value group of EMD values, from the sampled graph and the randomized graph.

There does seem to be some potential in the idea of the maximum flow playlist. When using the EMD as a weight the results appear to be quite good, at least from a qualitative perspective. The imposed constraint of the social network alleviates to some extent short comings of a playlist built purely through the analysis of acoustic similarity by moving more toward the balance between completely similar works and completely random movement.

Having shown the lack of a strong relationship between the maximum flow values and acoustic artist similarity, where do we go from here?

The most promise lies in the exploration of the maximum flow based playlist. A network could be built which was song to song exhaustive, having duplicate edges link each song individually to an artist's friends' songs. These edges would be weighted according to their acoustic similarity and a more complete playlist generation system would be created. A serious hurdle to the implementation of such a system lies in the computational complexity of the maximum flow value. Its compute time is typically dependent on both the number of nodes and the number of edges making it very slow to run on a network as dense as the one just described. This is less of a concern if some form of localized subgraphs were used, *e.g.* maximum flow is found against only friends (the *greedy* approach) or friends of friends. That said, there

may be strategies to get around these problems of complexity leading to novel and interesting playlist generation.

7 ACKNOWLEDGEMENTS

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USING EXPRESSIVE TRENDS FOR IDENTIFYING VIOLIN PERFORMERS

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ABSTRACT

This paper presents a new approach for identifying professional performers in commercial recordings. We propose a Trend-based model that, analyzing the way Narmour's Implication-Realization patterns are played, is able to characterize performers. Concretely, starting from automatically extracted descriptors provided by state-of-the-art extraction tools, the system performs a mapping to a set of qualitative behavior shapes and constructs a collection of frequency distributions for each descriptor. Experiments were conducted in a data-set of violin recordings from 23 different performers. Reported results show that our approach is able to achieve high identification rates.

1 INTRODUCTION

Expressive performance analysis and representation is currently a key challenge in the sound and music computing area. Previous research has addressed expressive music performance using machine learning techniques. For example, Juslin *et al.* [6] studied how expressivity could be computationally modeled. Ramirez *et al.* [12] have proposed an approach for identifying saxophone performers by their playing styles. Lopez de Mantaras *et al.* [9] proposed a case based reasoning approach to deal with expressiveness. Hong [7] investigated expressive timing and dynamics in recorded cello. Dovey [2] analyzed Rachmaninoff's piano performances using inductive logic programming. Work on automatic piano performer identification has been done by the group led by Gerhard Widmer. To cite some, in [14] they represent pianists' performances as strings; in [16] they study how to measure performance aspects applying machine learning techniques; and in [15], Stamatatos and Widmer propose a set of simple features for representing stylistic characteristics of piano music performers. Sapp [13] work should also be cited as an interesting proposal for representing musical performances by means of scape plots based on tempo and loudness correlation. Goebl [3] is focused on finding a 'standard performance' by exploring the consensus among different performers.

In this paper, we focus on the task of identifying violinists from their playing style using descriptors automatically extracted from commercial audio recordings by means of state-of-the-art feature extraction tools. First, since we consider recordings from quite different sources, we assume a high heterogeneity in the recording conditions. Second, as state-of-the-art audio transcription and feature extraction tools are not 100% precise, we assume a partial accuracy in the extraction of audio features.

Taking into account these characteristics of the data, our proposal therefore identifies violin performers through the following three stage process: (1) using a higher-level abstraction of the automatic transcription focusing on the melodic contour; (2) tagging melodic segments according to the E. Narmour's Implication-Realization (IR) model [10]; and (3) characterizing the way melodic patterns are played as probabilistic distributions.

The rest of the paper is organized as follows: In Section 2 we present the data collection being used. In Section 3, we describe the proposed *Trend-Based model* and the developed system, including data gathering, representation of recordings and distance measurement. In Section 4, experiments for the case study are explained and results are presented. The paper concludes with final considerations and pointing out to future work in Section 5.

2 MUSICAL DATA

We have chosen to work with Sonatas and Partitas for solo violin from J.S. Bach [8]. Sonatas and Partitas for solo Violin by J.S. Bach is a six work collection (three Sonatas and three Partitas) composed by the German musician. It is a well-known collection that almost every violinist plays during its artistic life. All of them have been recorded many times by several players. The reason of using this work collection is twofold: 1) we have the opportunity of testing our model with existing commercial recordings of the best known violin performers, and 2) we can constrain our research on monophonic music.

In our experiments, we have extended the musical collection presented in [11]. We analyzed music recordings from 23 different professional performers. Because these audio files were not recorded for our study, we have not interfered at all with players' style at the performance [5]. The scores

of the analyzed pieces are not provided to the system.

3 TREND-BASED MODELING

Our approach for dealing with the identification of violin performers is based on the acquisition of *trend models* that characterize each particular performer to be identified. Specifically, a trend model characterizes, for a specific audio descriptor, the relationships a given performer is establishing among groups of neighbor musical events. We perform a qualitative analysis of the variations of the audio descriptors. Moreover, as we will describe in the next subsection, we analyze these qualitative variations with a local perspective. We will be using two trend models in this paper: energy and duration. The trend model for the energy descriptor relates, qualitatively, the energy variation for a given set of consecutive notes, while the trend model for duration indicates, also qualitatively, how note durations change for note sequences. Notice that trend models are not trying to characterize the audio descriptors with respect to an expected global behavior.

Given an input musical recording of a piece, the trend analysis is performed by aggregating the qualitative variations on their small melody segments. Thus, in advance of building trend models, input streams are broken down into segments. As most of automatic melody segmentation approaches, we will perform note grouping according to a human perception model.

Our system has been designed in a modular way with the intention of creating an easy extendable framework. We have three different types of modules in the system: 1) the audio feature extraction modules; 2) the trend analysis modules; and 3) the identification modules. Moreover, the system may work in two different modes: in a training mode or in a testing mode. Modules from (1) and (2) are used in both modes. Modules from (3) are only used in the testing mode. Figure 1 shows a diagram of the system modules. On top, audio files in *.wav* format as input.

At the training stage, the goal of the system is to characterize performers by extracting expressive features and constructing trend models. Next, at the identification stage, the system analyzes the input performance and looks for the most similar previously learnt model. The training process is composed of three main steps: 1) the extraction of audio descriptors and the division of a performance in segments; 2) the tagging of segments according to IR patterns; and 3) the calculus of probabilistic distributions for each IR pattern and descriptor (trend generation).

3.1 Feature Extraction and Segmentation

The first step consists on extracting audio features. Our research is not focused on developing new methods for extracting audio features, so that we employ existing techniques. At the moment, we consider fundamental frequency and energy, as these are the main low-level audio features related to melody. These features are then used to identify note boundaries and to generate melodic segments. The current version of our system uses the fundamental frequency estimation algorithm proposed by Arturo Camacho [1]. This module provides a vector with instantaneous fundamental frequency and energy values.

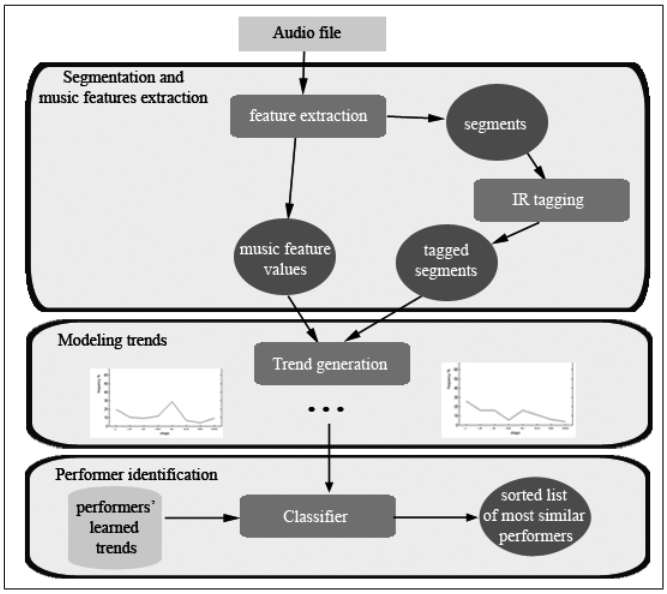

Figure 1. System's dataflow

We have developed a post-processing module for determining the possible notes played by the performers. This module first converts fundamental frequencies into quantized pitch values, and then a pitch correction procedure is applied in order to eliminate noise and sudden changes. This correction is made by assigning to each sample the value given by the mode of their neighbors around a certain window of size σ. With this process, a smooth vector of pitches is obtained. By knowing on which frames pitches are changing, a note-by-note segmentation of the whole recording is performed. For each note we collect its pitch, duration and energy.

We assume that there might be some errors in this automatic segmentation, given the hetereogenity of recording conditions. Our approach for dealing with this problem consists of using a more abstract representation that the real notes, but still close to the melody. That is, instead of focusing on the absolute notes, we are interested in modeling the melodic surface.

We use the IR model by E. Narmour [10] to perform melodic segmentation. This model tries to explicitly describe the patterns of expectations generated in the listener with respect to the continuation of the melody. The IR model describes both the continuation implied by particular melodic

intervals, and whether or not this expected continuation is fulfilled by the following interval. Taking this cognitive model as the basis for the melodic segmentation, each IR pattern determine a different segment.

The IR model has been shown to be suitable for assessing melodic similarity (see MIREX'05 [4]). Since our goal is to characterize expressive trends, we analyze the way different audio descriptors change in the different IR patterns. See [10] for a detailed description of IR patterns.

3.2 Modeling Trends

A trend model is represented by a set of discrete probability distributions for a given audio descriptor (e.g. energy). Each of these probability distributions represents the way a given IR pattern is played against that certain audio descriptor.

To generate trend models for a particular performer and audio descriptor, we use the sequences of values extracted from the notes identified in each segment. From these sequences, a qualitative transformation is first performed to the sequences in the following way: each value is compared to the mean value of the fragment and is transformed into a qualitative value where + means 'the descriptor value is higher than the mean', and - means 'the descriptor value is lower than the mean'. Being s the size of the segment and n the number of different qualitative values, there are n^s possible resulting shapes. In the current approach, since we are segmenting the melodies in groups of three notes and using two qualitative values, eight (2^3) different shapes may arise. We note these possibilities as: —, –+, -+-, -++, +–, +-+, ++- and +++.

Next, a histogram per IR pattern with these eight qualitative shapes is constructed by calculating the percentage of occurrence of each shape. These histograms can be understood as discrete probability distributions. Thus, trend models capture statistical information of how a certain performer tends to play. Combining trend models from different audio descriptors, we improve each performer characterization.

Since our goal is the identification of violin performers, the collection of trend models acquired for each performer is used as the patterns to compare with when a new audio recording is presented to the system.

3.2.1 Current Trends

We have generated trend models for both duration and energy descriptors. Note durations are calculated as the number of samples between pitch changes. Qualitative deviations are calculated by comparing the real and the average duration for each note. Then, the trend model for the duration descriptor was calculated from the frequency distributions of each IR pattern. Figure 2 shows, for the duration descriptor, the frequency distributions of the eight shapes in P patterns (ascending or descending sequences of similar intervals) and for four violin performers (Ara Malikian, Arthur Grumiaux, Brian Brooks, and Christian Tetzlaff).

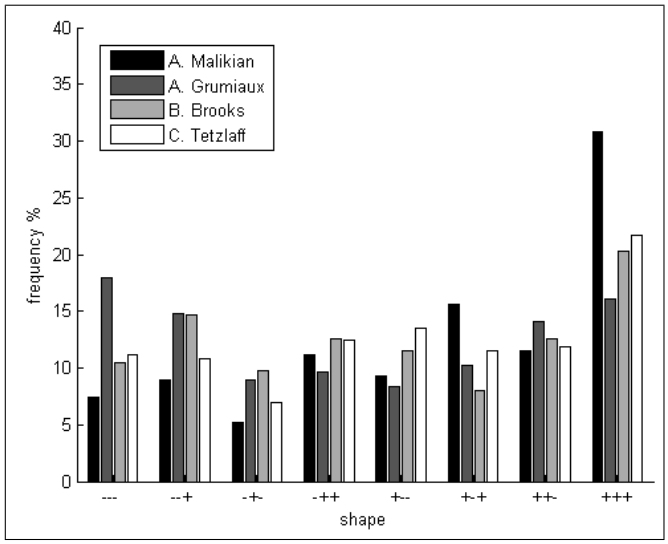

Figure 2. Frequency distribution of duration deviations for the P pattern in the Sixth movement of Partita N.1. Only four performers are shown

We can observe that the way different professional performers are playing is not equally distributed. For instance, A. Malikian has a higher propensity to extend the durations while an opposite behavior can be observed for A. Grumiaux (see his values for the two left qualitative shapes). It should be noticed that more exact ways of obtaining this measure could be used, as well as taking into account the attack and release times, as other researchers do [12].

We have also acquired trend models for the energy descriptor in an analogous way. The energy average for each fragment is calculated and, given the energy of each note, the qualitative deviations are computed. Next, from these qualitative values, the trend models are constructed by calculating the frequencies of the eight shapes for each IR pattern.

3.3 Classifying new performances

A nearest neighbor (NN) classifier is used to predict the performer of new recordings. Trend models acquired in the training stage, as described in the previous section, are used as class patterns, i.e. each trained performer is considered a different solution class. When a new recording is presented to the system, the feature extraction process is performed and its trend model is created. This trend model is compared with the previously acquired models. The classifier outputs a ranked list of performer candidates where distances determine the order, with 1 being the most likely performer relative to the results of the training phase.

3.3.1 Distance measure

The distance d_{ij} between two trend models i and j, is defined as the weighted sum of distances between the respective IR patterns:

$$d_{ij} = \sum_{n \in N} w_{ij}^{n} dist(n_i, n_j) \quad (1)$$

where N is the set of the different IR patterns considered; $dist(n_i, n_j)$ measures the distance between two probability distributions (see (3) below); and w_{ij}^{n} are the weights assigned to each IR pattern. Weights have been introduced for balancing the importance of the IR patterns with respect to the number of times they appear. Frequent patterns are considered more informative due to the fact that they come from more representative samples. Weights are defined as the mean of cardinalities of respective histograms for a given pattern n:

$$w_{ij}^{n} = (N_i^n + N_j^n)/2 \quad (2)$$

Mean value is used instead of just one of the cardinalities to assure a symmetric distance measure in which w_{ij}^{n} is equal to w_{ji}^{n}. Cardinalities could be different because recognized notes can vary from a performance to another, even though the score is supposed to be the same.

Finally, distance between two probability distributions is calculated by measuring the absolute distances between the respective patterns:

$$dist(s, r) = \sum_{k \in K} |s_k - r_k| \quad (3)$$

where s and r are two probability distributions for the same IR pattern; and K is the set of all possible values they can take (in our case $|K| = 8$).

When both audio descriptors are considered, we simply aggregate the individual corresponding distances.

4 RESULTS AND DISCUSSION

We tested our system by performing experiments using different information coming from the acquired trend models. Specifically, we evaluated each experiment setting with only duration-based trend models, only energy-based trend models, and with both trend models.

Each experiment consisted in training the system with one movement and then testing the trend models acquired presenting to the system the recordings from another movement. For experimentation, we used a collection of audio recordings from 23 different professional violinists. We performed two different types of experiments. The first experiment was focused on assessing the performance of the system by using two movements from the same piece. Specifically, we used the Second and the Sixth movements of Partita No. 1. These fragments are quite interesting for early testing because most of the notes are eighth notes, leading us to acquire a model based on many homogeneous segments. In the following, we will call **set-1** the experiment where the second movement was used for training and the sixth for testing. Analogously, we will call **set-2** the experiment where the sixth movement was used for training and the second for testing.

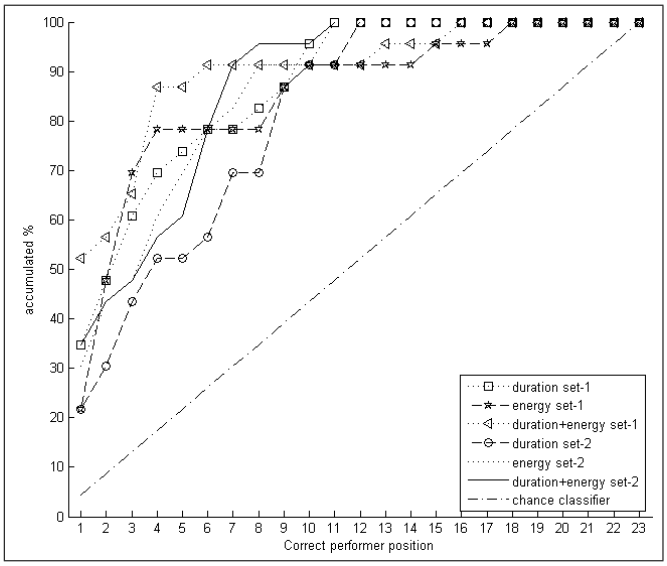

Figure 3. Accumulated success rate by position of the correct performer. **Set-1** and **set-2** are shown

The second type of experiments was focused on assessing the performance of the system by using two movements from different pieces. Specifically, we used the sixth movement of Partita No. 1 for training and the fifth movement of Partita No. 3 for testing. We will refer to this test as **set-3**.

For each input recording, the system outputs a ranked list of performers sorted from the most similar to the farthest one to the input. In experiments **set-1** and **set-2**, the correct performer was mostly identified in the first half of the list, i.e. at most in the 12th position. In **set-3**, the most difficult scenario, the 90% of identification accuracy was overcame at position 15. Regarding the different trend models, the energy model was the one that achieved a highest accuracy. This result is not surprising since the duration descriptor presents a high sensitivity with respect to the analysis precision. The combination of both trend models improves the results when focusing only on the first proposed player.

A complete view of the performance of the system is shown in Figure 3 and summarized in Table 1. Figure 3 shows the percentage of input recordings identified at each position. It provides a picture of the accuracy of the system using as a threshold the length of the proposed ranking. The results are promising, especially comparing with a random classification where the success rate is clearly outperformed. We can observe that the system achieved a 50% of success

	set-1			**set-2**			**set-3**		
	1st	3rd	10th	1st	3rd	10th	1st	3rd	10th
duration	34.8	60.9	95.7	21.7	43.5	91.3	10.5	26.3	68.4
energy	21.7	69.6	91.3	30.4	47.8	91.3	15.8	31.6	73.7
both	52.2	65.2	91.3	34.8	47.8	95.7	15.8	26.3	68.4

Table 1. Success rate (%) in all experiments taking into account three different ranking positions proposed for the correct performer: 1st, 3rd, and 10th

using the four top candidates in **set-1** and **set-2**.

Table 1 summarizes the performance of the system for the three experimental sets and the three trend models. The three columns of each experiment show, respectively, the percentage of performers identified in the first position, at least in the third position, and at least in the tenth position. We can observe that for settings **set-1** and **set-2** the correct performer is predicted, in the worst case, 20% of times as the first candidate, clearly outperforming the random classifier (whose success rate is 4.3%).

Figure 4 presents a matrix that summarizes the classifier output for **set-2** using both duration and energy trend models. The figure shows, for each input recording (row), the sorted list of predicted performers as squares. The gray scale maps to the ranking values. The black color indicates the first performer proposed and the gray degradation is used to draw all the performers predicted until the correct one. Notice that the success in the first position means a black square in the diagonal. The matrix is not supposed to be symmetric and each column can have the same color several times because a predicted performer can occur in the same position for several inputs. For instance, we can see that Garret Fischbach's performance (`gar`) for Sixth Movement is very different from the rest of performers' Second Movement performances: all values correspond to last position (i.e. the furthest). On the other hand, Christian Tetzlaff's (`chr`) and Rachel Podger's (`rac`) performances are quite similar to most of Second Movement performances since there are many squares in their columns.

Finally, Figure 5 shows in which position the correct performer is ranked for each performer in the test set. This Figure complements the former two. The results came from **set-1** using both trend models ('duration+energy set-1' curve in Figure 3). Twelve right identifications were achieved at first position (52% of success rate). The rest was correctly identified in positions 2 to 4 except three performers. Nathan Milstein was identified at position 6. Finally, Sergiu Luca and Shlomo Mintz were not clearly identified. After a detailed analysis of the distances among all performers, we observed that these two musicians are not clearly distinguished. We mean, small variations in the trend models confuse the identification process.

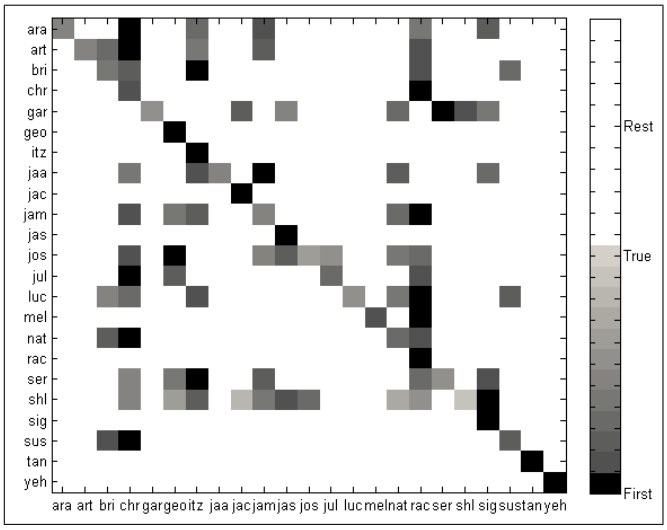

Figure 4. Classifier output in matrix form for **set-2** where both trend models were used

5 CONCLUSIONS

This work focuses on the task of identifying violinists from their playing style by building trend-based models that capture expressive tendencies. Trend models are acquired by using state-of-the-art audio feature extraction tools and automatically segmenting the obtained melodies using IR patterns. Performers were characterized by a set of probability distributions, capturing their personal style with respect to a collection of melodic patterns (IR patterns). We have shown that, without a great analysis accuracy, our proposal is quite robust.

The experiments were concentrated on identifying violinists and using note durations and energies as descriptors. We tested the system with 23 different professional performers and different recordings. Results obtained show that the proposed model is capable of learning performance patterns that are useful for distinguishing performers. The results clearly outperform a random classifier and, probably, it would be quite hard for human listeners to achieve such recognition rates. In order to assure the robustness of the system, other sets of works should be used for learning and testing.

We have presented a qualitative analysis using only two qualitative values. We want to keep our model at this qualitative level but we plan to extend the model with the use of fuzzy sets. This improvement will allow us to use the capabilities of fuzzy theory for a better assessment in the similarity measure.

Combining information from different music features have been demonstrated to improve results. We are currently working in increasing the number of descriptors. Since the predictability of a given descriptor varies depending on the

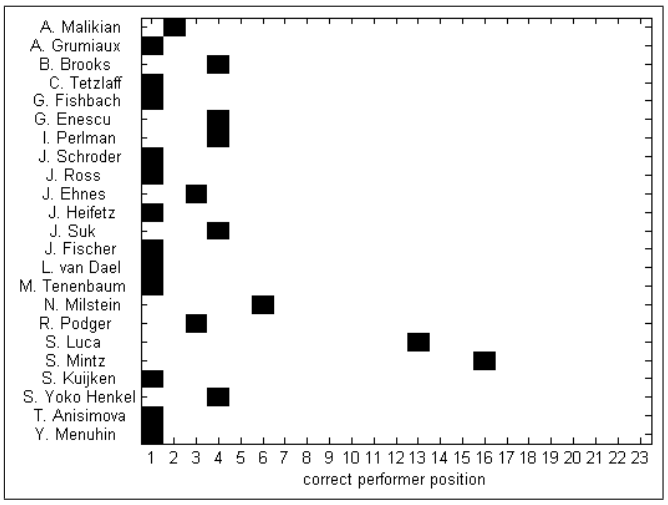

Figure 5. Correct performer position for each performance in **set-1**. Both trend models are used

performers, we are also interested in discovering relations among the descriptors. Finally, the use of hierarchical classifiers or ensemble methods is a possible way of improving the identification.

6 ACKNOWLEDGEMENTS

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CONNECTING THE DOTS: MUSIC METADATA GENERATION, SCHEMAS AND APPLICATIONS

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ABSTRACT

With the ever-increasing amount of digitized music becoming available, metadata is a key driver for different music related application domains. A service that combines different metadata sources should be aware of the existence of different schemas to store and exchange music metadata. The user of a metadata provider could benefit from knowledge about the metadata needs for different music application domains. In this paper, we present how we can compare the expressiveness and richness of a metadata schema for an application. To cope with different levels of granularity in metadata fields we defined clusters of semantically related metadata fields. Similarly, application domains were defined to tackle the fine-grained functionality space in music applications. Next is shown to what extent music application domains and metadata schemas make use of the metadata field clusters. Finally, we link the metadata schemas with the application domains. A decision table is presented that assists the user of a metadata provider in choosing the right metadata schema for his application.

1. INTRODUCTION

Metadata plays an important role in MIR research. In 2007, we proposed a semi-automatic approach for the generation of music metadata [2] in the rockanango project [1]. In this approach, we bundled the power of computational techniques, the wisdom of the crowds and the experience of the music experts of Aristo Music (www.aristomusic.com), the company with whom we collaborate. This was needed because of the increasing volume of music that needs annotation within a reasonable amount of time and cost.

A diversity of available metadata web services can be used to facilitate the annotation process. Amazon.com has a substantial web API to retrieve metadata about their products. Available music metadata for CD's includes reviews from editors and customers, track listings, release dates, genres, labels, popularity and similar items. Last.fm (http://last.fm), the well-known social recommendation music service, provides a web service to retrieve user, artist, group and tag data. The All Music Guide is also a very rich source of music metadata. MusicBrainz (http://www.musicbrainz.org) is a community based music metadata database used for CD recognition. It contains artists, tracks, labels and releases. Discogs (http://discogs.com) offers a neat web service with good pointers to different production houses and publishers.

Another kind of web service is available that, given an audio file, returns the results of different signal processing algorithms. Echo Nest (http://analyze.echonest.com/) recently released a web service that returns a set of musical features, like timbre, pitch, and rhythm. It is aiming at different applications, such as visualizations, games and DJ-software. Furthermore, Echo Nest has been developing software that 'listens' to music and tries to predict which songs will become hits. The MIR Group of the Vienna University of Technology (http://www.ifs.tuwien.ac.at/mir/webservice/) also made a web service available that returns a set of musical features for a given song (e.g. rhythm patterns, statistical spectrum descriptors and rhythm histograms) and allows the training of self-organizing music maps. In conclusion, these systems provide diverse and useful metadata, however not without a substantial amount of overlap.

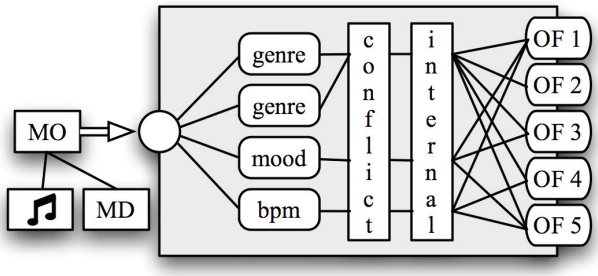

Figure 1. Internals of the metadata framework.

We want to build a framework that makes use of the strength of the different systems (see Figure 1). The input is a musical object (MO) that consists of an audio file (the signal) and possibly some initial metadata (MD). When it enters the system, a federated request for metadata among available generators is done. Generators can use available a priori metadata to enhance predictions. To cope with potentially conflicting results, [3] suggests different conflict resolution approaches. The resulting metadata is

stored in an internal format. The generated metadata can cover the full range from low-level features, over factual data e.g. the arrangement of the band, to affective metadata e.g. the mood, similar songs or popularity.

By offering the choice between different existing music metadata formats (OF), we enable the reuse of existing tools and parsers to handle the generated metadata. To be able to store all the generated metadata we are looking for a suitable metadata schema for internal use in the metadata framework. In this paper we will select a set of metadata standards relevant for the task, investigate in which application domains they are useful and evaluate their descriptive richness.

The description of large music works, e.g. the complete opus of a composer, is beyond the scope of this paper. Likewise metadata schemas for metadata exchange (e.g. METS [18]) or rights (e.g. ODRL [19]) are also out of the scope. We focus on descriptive metadata schemas.

2. COMPARISON OF METADATA SCHEMAS

The goal of collecting metadata is to enable functionalities in an application. To understand the usefulness of a metadata schema in different cases, we will investigate which metadata is involved in different use cases. This will be the foundation for defining music application domains and selecting a number of relevant metadata schemas. For the sake of easier comparison, we introduce a level of abstraction to the metadata fields by means of field clustering. The three elements: the selected metadata standards, the different application domains and categories of metadata fields will be introduced in this section. The question which metadata schemas to use when building an application that offers a certain functionality can then be answered through the comparison of the three elements.

2.1. Application domains

Music metadata can be anywhere in the production cycle of music, whether it be copyright information about a drum loop while composing or the number of searches for a musical piece at an online store. For the use of metadata schemas, we limit our application domains to software.

Clustering software applications into application domains allows easier comparison between different applications. The actual clustering is based on the actions people perform when handling music [4] [14]. The eight clusters cover the production cycle from conception (composition) over consumption to transactions.

- *Music library/encyclopedia:* software systems for physical libraries, encyclopediae or companies that license music, describing factual knowledge for large collections of music, e.g. AllMusic Guide.
- *Personal collection management:* software to organize your music collection, e.g. iTunes, Winamp media library, Collectorz.com Music Collector.
- *Commerce and transactions:* applications involved in the act of shopping for music, this includes presenting songs, searching and trading, e.g. iTunes Music Store, Amazon, Magnatune.
- *Music editing/production:* the tools deployed in the creation and adaptation of music, e.g. Logic Pro.
- *Music playback:* applications that render music files to its audible form, e.g. your favorite music player.
- *Music recommendation:* services for discovery of new and similar music, e.g. Pandora, Last.fm.
- *Music retrieval:* search and identification tools with different query interfaces in all their forms, e.g. query by humming.
- *Musical notation:* creation and manipulation tools for musical scores, e.g. Sibelius, WAV2Midi.

A real-life application will most likely use a number of application domains, e.g. playlist generation can be classified as music recommendation and library functionality, some music players also offer management of the personal music collection.

2.2. Metadata standards

Metadata standards originate from different sources: it can evolve out of the design of an application and through wide adoption become a de facto standard, it can be focused on interoperability or it can be designed as a standard from the start. No single metadata standard is at the moment available for music covering all the possible requirements. Based on industry standards and the use in ongoing research we selected eight music metadata standards. The methodology presented in this paper is applicable to the many other available standards (e.g. MARC [20], MODS [21]). Future work includes extending the comparison with other relevant schemas.

- *ID3:* An ID3-tag is a data container of a prescribed format embedded within an audio file. The stored data can contain the artist name, song title and genre of the audio file. ID3 has a wide spread use in music players and devices like iTunes, iPod and Winamp.
- *FreeDB:* is an online database to look up CD information by calculating a unique ID for a CD to query the database. FreeDB is a community version of the commercial Gracenote service; both are used in a variety of playback and MP3-ripping software.
- *MusicBrainz:* the scope of MusicBrainz [6] is the same as FreeDB, but they have a moderated database and use identification on track level. MusicBrainz is an RDF-based [7] web service.
- *Dublin Core:* is a standard for cross-domain information resource description, which provides a simple and standardized set of conventions for

describing things online [11]. It is widely used to describe digital multimedia materials.

- *Music Vocabulary:* describes classic musical works and performances in an RDF-ontology [8], defining classes for musical works, events, instruments, performers and relationships between them.
- *Music Ontology:* the goal is to link information about artists, albums and tracks together and express relationships between musical information for the Semantic Web [10]. The Music Ontology is based on Functional Requirements for Bibliographic Records (FRBR) [13], but removes the Expression entity to be able to model the creational process behind music and is available as an RDF-ontology.
- *MPEG-7:* is a multimedia content description standard. MPEG-7 [9] provides standardized tools for describing different aspects of multimedia at different levels of abstraction. The XML-based syntax enables interchange across applications, but without precise semantics metadata interoperability can be a problem.

2.3. Metadata field clusters

Different metadata standards describe different things. To facilitate the comparison of the standards we clustered the metadata fields in an iterative process. First we clustered the elements of ID3 in semantically related clusters. Next, we tried to map Music Ontology onto the ID3 clustering. This resulted in a small adjustment in the clusters. The same procedure was used to add the remaining metadata standards. The final clustering defines semantically related elements, independent of metadata schemas or metadata fields. The following clusters are defined:

- *Musical info:* musical properties of the audio signal, e.g. bpm, duration, key.
- *Classifiers:* categorizers for music organization, e.g. genre.
- *Performance:* descriptions of the people that are involved in a musical performance, e.g. artists, conductor, engineers.
- *Versioning:* descriptions for different versions of the musical work and relationships between the involved people and the related works, e.g. original artists, contains sample from, remix of.
- *Descriptor:* describing the musical work, e.g. title.
- *Rights & ownership:* information about intellectual property management, e.g. license, owner.
- *Playback rendition:* information useful for rendering of the music file, e.g. relative volume adjustment.
- *Lyrics:* text of the musical work and related information, e.g. translated lyrics, synched lyrics.
- *Grouping & referencing:* data clustering structures and linking to web sites and other works, e.g. wikipedia link, album listing structure.
- *Identifiers:* identification keys, e.g. ISRC-code.
- *Record-info:* information about the recording and the involved people, e.g. recording type, album name.
- *Instrumentation & arrangement:* about the used instruments and orchestration, e.g. instrument type.
- *Sound & carrier:* information about the available media of a musical work, e.g. media type.
- *Event:* information about public performances of the musical work, e.g. festival program, place.
- *Time-modeling:* metadata structures to express points in time, e.g. interval, at duration.
- *Musical notation:* everything for a symbolic representation of music, e.g. clefs, measures, notes.
- *Attention-metadata & usage:* information about the user's attention to music, e.g. times played.
- *Publishing:* information about the publishing process, e.g. record label.
- *Composition:* information about the composition process, e.g. movement, composer.
- *Production:* information about the creative process of the production and its actors, e.g. producer.
- *Meta-metadata:* information about the metadata, e.g. who annotated the metadata.

2.4. Comparison

In this subsection we determine how the clusters, the schemas and the application domains relate to one another. As said in 2.1, we are mainly interested in determining which metadata schema to use for some desired functionality. We use the level of abstraction of the metadata clusters to join metadata schemas and application domains.

2.4.1. Application domains vs. metadata field clusters

In table 1 we compared the music metadata fields with the application domains. When assigning music metadata field clusters to application domains, the different domains are considered from the perspective of the provider of the functionality. We used the results of survey [4] on how users search for the library and encyclopedia domain. For the personal music collection management domain, we used a combination of the survey presented in [5] and we looked at the applications in the

domain. For the other domains, we investigated the metadata usage of different applications in these domains to determine to what degree which clusters are used.

	libr./enc.	pers. coll.	comm.	edit./prod.	playback	recomm.	retrieval	notation
musical info	-	+/-	-	+	+/-	+	+	+
classifiers	+	+	+	-	-	+	-	-
performance	+	+	+	+	+/-	+	+	-
versioning	+	+	+	+/-	-	+	+	-
descriptor	+	+	+	+	+	+	+	+
rights & ownersh.	+	+	+	+	+/-	-	-	-
playback rendition	+/-	+/-	-	-	+/-	-	-	-
lyrics	+	+	-	+	+/-	+	+	+
group. & refs.	+	+	+	+/-	-	+	-	-
identifiers	+	+	+	-	-	+/-	+/-	-
record-info	+	+	+	-	-	+	+/-	-
instr. & arrang.	+	+/-	+/-	-	-	+	-	+
sound & carrier	+	+	+	-	+/-	-	-	-
event	+	+/-	+/-	-	-	+/-	-	-
time-modelling	-	-	-	+	-	-	+/-	+
notation	-	-	-	+/-	-	-	+	+
attention-metadata	-	+	-	-	+	+	+/-	-
publishing	+	+	+	-	-	+/-	-	-
composition	+	+	+	+	-	+	-	+
production	+	+	+	+	-	+	-	-
meta-metadata	+	+	+	+	-	-	-	-

Table 1. Metadata field clusters vs. application domains

Based on Table 1, we can conclude that library and encyclopedia, personal collection management, commercial applications and recommendation make use of the largest set of metadata clusters. The bibliographical nature of libraries strives for completeness and accounts for a wider range of clusters. For collections, recommendation and commercial use, [16] and [4] suggest wide ranges of user goals, often with a focus on enjoyment that require a broad range of information. In the latter case, rights management and information about the carrier is relevant. Information closely describing the actual audio signal found in musical info is not so relevant here as opposed to e.g. playback where descriptive metadata typically is not needed to render an audio file.

Recommendation and retrieval rely both on signal and metadata in order to function. Pachet [16] defines 3 music metadata categories: editorial, cultural and acoustic. Music recommendation techniques can make use of elements out of these 3 categories [17].

Music notation is useful, either during composition, early in the music creation cycle, or while describing already existing music by means of transcription. Information about publishing, producing, recording, etc. are not immediately relevant.

The table can be used as a reference for the metadata needed when building an application.

2.4.2. Metadata standards vs. metadata field clusters

	ID3	freeDB	MusicBrainz	Dublin Core	Music Vocabulary	Music Ontology	MPEG-7	MusicXML
musical info	+	+/-	+/-	-	+/-	+	+	+
classifiers	+/-	+/-	-	+/-	+/-	+/-	+	+/-
performance	+	+/-	+/-	+	+	+	+	+/-
versioning	+	-	-	+	+/-	+	+	-
descriptor	+	+	+	+	+	+	+	+
rights & ownersh.	+	-	-	+	-	+	+	+
playback rendition	+	-	-	-	-	+	+	+
lyrics	+	-	-	-	+	+	+	+
group. & refs.	+	-	-	+	+/-	+	+	+
identifiers	+	+	+	+	-	+	+	-
record-info	+	+	+	-	-	+	+	-
instr. & arrang.	+/-	-	-	-	+	+	+	+
sound & carrier	-	-	-	+	-	+	+	+/-
event	-	-	-	-	+	+	+	-
time-modelling	-	-	-	-	-	+	+	+
notation	-	-	-	-	+/-	+/-	-	+
attention-metadata	+	-	-	-	-	+	+	-
publishing	+	-	-	-	-	+	+	-
composition	+	-	-	+/-	+	+	+	+
production	-	-	-	-	-	+	+	-
meta-metadata	-	+	-	-	-	-	+	-

Table 2. Metadata field clusters vs. metadata standards

By doing the clustering of the metadata fields for each metadata standard (see 2.3), we obtain a table with the fields of each metadata standard in the corresponding clusters. This table enables us to determine how good the metadata clusters are represented in the different metadata standards, as can be seen in table 2.

From table 2, we can clearly see that the scope of Dublin Core is more general than music alone, because the clusters typically related to music are not present. MPEG-7 and Music Ontology are the most versatile schemas that are present in almost all clusters. MusicBrainz and freeDB have similar fields and scope.

Some clusters are better represented in some standards, like the performance cluster in Music Vocabulary, Music Ontology and MPEG-7.

2.4.3. Application domains vs. metadata standards

	libr./enc.	pers. coll.	comm.	edit./prod.	playback	recomm.	retrieval	notation
ID3	70	75	70	67	90	81	81	64
freeDB	30	31	36	29	30	30	38	21
MusicBrainz	21	22	25	19	30	26	38	21
Dublin Core	48	47	57	43	50	41	44	21
Music Vocabulary	45	40	39	50	45	56	56	71
Music Ontology	91	91	89	88	100	96	94	93
MPEG-7	100	100	100	95	100	100	88	86
MusicXML	48	47	43	71	70	52	63	100

Table 3. Application domains vs. metadata standards in %

We use the first two tables to determine which metadata schema is most apt, given the application domain.

To be able to compute the decision table we formalized table 1 and 2. The +, +/- and - signs are converted to 1, 0.5 and 0 respectively. This results in 2 matrices: MCAD (table 1) of dimension (#metadata field clusters (MC) x #application domains) and a matrix MCMS (table 2) of dimension (#metadata field clusters x #metadata standards). The sum of the columns of MCAD results in the maximum score any metadata standard can reach for an application domain, a vector mssum.

$$mssum_j = \sum_{i=1}^{\#MC} MCAD_{i,j} \quad (1)$$

We now calculate the product of the transposed MCAD and MCMS in (2). This measures how well a metadata standard covers the clusters for each application domain.

$$R_{i,j} = \sum_{x=1}^{\#MC} MCAD_{i,x}^{T} . MCMS_{x,j} \quad (2)$$

The mssum vector is used to normalize the values for the different schemas (3) and this results in a matrix MFAD, comparing metadata schemas with application domains, see table 3 for the actual values.

$$MFAD_{i,j} = \frac{R_{i,j}}{mssum_j} \quad (3)$$

Note that we do not penalize excess information, because most of the metadata fields are not obligatory. If they are not needed, they can be omitted.

3. ANALYSIS OF THE DECISION TABLE

It is clear that MPEG-7 scores well for all application domains. Does this mean that this is the über music metadata standard that should be the only output format for a metadata generation framework? No, though it is possible to store any kind of information in MPEG-7 through the use of description schemas and the description definition language. It might be the case that the more limited set of e.g. Dublin Core is sufficient. The same is valid for MusicOntology.

Some interoperability issues with MPEG-7 have been signaled [15]. Semantically identical data can be represented in multiple ways, e.g by keywords, free text and a structured annotation with labels. Another issue is that the intended semantics of description structures are not always clear in MPEG-7.

FreeDB and MusicBrainz seem to score low overall, but seem best suited for music retrieval. Both are deployed in online services focus on identification of songs/albums. The available information linked to the identifier consists mainly about artist, track and album related metadata. Applications will use that data to fill in other metadata schemas, for example an ID3 tag.

As expected, MusicXML is extremely relevant for notation. This is the main goal of the standard. With the existence of tools that convert MusicXML to MIDI, it can also be used in playback.

Music Vocabulary's goal is to define a vocabulary to describe classical music masterworks and performances/concerts [6]. This limited scope has severe consequences for the general expressiveness of the schema. The fact that it scores relatively high on music notation is mainly because of its elaborate description of composition and orchestration/instrumentation. It could be used together with Music Ontology, to enrich its classical work descriptions.

Dublin core is designed to offer simple cross-domain metadata fields. This leaves very little margin for music domain specific information. However, often a simple, interoperable description is desired for an application. MusicXML, Music Ontology and Music Vocabulary use parts of Dublin Core, for example dc:author.

The creators of ID3 use 'The audience is informed' as their motto. This is visible in the good performance in the playback application domain. It scores equally well for recommendation applications, due to its expressiveness in classifiers, descriptors, performance and versioning.

4. FUTURE WORK

The implementation of a music metadata framework as described in Figure 1 is currently under development. It will be integrated in the SAmgI framework [3], where metadata for learning content is gathered, combined and presented to the end user in multiple formats, e.g. LOM, Dublin Core and human readable. The fundamentals for conflict resolution and management of different generators are already present. Furthermore SAmgI runs as a Java web service that embraces distributed metadata generation. The main challenge is adapting the internal structure for music specific metadata fields.

5. CONCLUSION

In conclusion of this study on the expressiveness and richness of different music metadata schemas and their relation with application domains, we can state that the formalization of these relations offers two practical uses in the context of music metadata generation. Firstly, once an application developer has decided on the functionality, he can determine what parameters (clusters) he should retrieve from a metadata generation framework based on the desired application domains. Secondly, the metadata framework developer can decide on the metadata formats to be returned given the set of requested parameters.

The scope of this work is not to provide a metadata schema decision table to the detail of individual fields of the metadata schemas. However it provides useful insights in whether schemas are relevant for certain application domains. The granularity of functionality can be refined after a first selection, but this requires deep knowledge of the considered application.

6. ACKNOWLEDGEMENT

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CREATING AND EVALUATING MULTI-PHRASE MUSIC SUMMARIES

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ABSTRACT

Music summarization involves the process of identifying and presenting melody snippets carrying sufficient information for highlighting and remembering a song. In many commercial applications, the problem of finding those snippets is addressed by having humans select the most salient parts of the song or by extracting a few seconds from the song's introduction. Research in the automatic creation of music summaries has focused mainly on the extraction of one or more highly repetitive phrases to represent the whole song. This paper explores whether the composition of multiple "characteristic" phrases that are selected to be highly dissimilar to one another will increase the summary's effectiveness. This paper presents three variations of this multi-phrase music summarization approach and a human-centered evaluation comparing these algorithms. Results showed that the resulting multi-phrase summaries performed well in describing the songs. People preferred the multi-phrase summaries over presentations of the introductions of the songs.

1. INTRODUCTION

People use summaries to concisely describe or highlight the major points of the genuine object. In text, for example, the authors of a scientific paper summarize the key points of their presentation in an abstract, a paragraph briefly describing the topic and their achievements or ideas. Accordingly, in music, vendors of CDs and mp3s (like Amazon.com) provide small snippets of songs to help potential customers become familiar with the contents of an album or to find songs they can only recall by melody. Similarly, radio stations remind listeners of the top-ten hits of the week by playing the refrains of the respective songs.

As the examples above indicate, a music summary consists of the part(s) of the song that are short in duration but rich enough in information to describe and identify the total for the given current task. Such a conclusion implies that the location or the duration of those parts is not fixed for all music since their selection depends on factors like the song, the user's perception of music and the task at hand (e.g. selecting from known music or deciding whether to buy unknown music). Finding a summarization approach that takes into account all three factors requires a model of the music, a model of the user, and a model of the task. Most commercial on-line music stores preview their songs by either the introduction of the song, a randomly selected phrase, or the (often manually selected) refrain. The simplicity of these approaches has two main problems. On one hand, there is no guarantee that the selected phrase is sufficient for becoming familiar with or recognizing the song. On the other hand, using human resources to find the refrain for thousands of songs is costly in terms of time, effort and money.

Much of the existing work in music summarization focuses on the selection of the most repeated phrase(s). When more than one phrase is selected, it is generally because the desired summary length is longer than the identified refrain and the added segment is the phrase identified as the next most frequent regardless of its similarity to the already selected refrain. As a step beyond refrain selection, we explore summaries designed to include more parts than the most salient phrase or the introduction of the song. To examine the design space for such algorithms, we compare algorithms that compose a summary from a fixed number of components (three) but vary the selection of those components between preferring phrases that are sonically different and phrases that are repeated more often.

In this paper we present three summarization algorithms that follow the principles above and an evaluation comparing their performance in the context of pop and rock music. Section 2 discusses the related work in automatic summarization and a comparison of techniques. Section 3 describes the summarization algorithms. Section 4 introduces the study and the procedures followed. An analysis of the results is presented in section 5. Finally, section 6 summarizes the results and presents directions for possible improvements of the current algorithms and follow-on studies.

2. RELATED WORK

Research into techniques for the extraction of sound / music features (i.e. tempo, brightness, fundamental frequencies, bounds of phrases) is quite fertile. This work has expanded into research for developing music summaries that tend to focus on the problem of identifying musical phrases and, in particular, the refrain. Hence, the success of summarization algorithms has been typically

evaluated based on how accurately they can determine the most repeated phrase. There are a variety of approaches to identifying the refrain.

A number of algorithms [1, 2] use a pattern matching approach where the structure of the content, and more specifically the most salient phrase, is determined by comparing candidate segments (a fixed sequence of frames) with the whole song. Cooper and Foote [3], after the parameterization of the signal with the calculation of the Mel Frequency Cepstrum Coefficients (MFCCs), find the distance in the parameter vectors of all frame combinations and store the results in a two-dimensional self-similarity matrix. To select the segment (sequence of frames) that best represents the entire song, they calculate the similarity of each segment to the whole and choose the one with the maximum value. If the phrase is not as long as the desired summary, they add the next highest ranking phrase(s).

Other algorithms develop more domain-specific models of the music in order to identify the most repeated phrase. Logan and Chu [5] use a three step process for extracting the key phrase. After segmenting the song, they cluster the resulting segments using a modified cross-entropy or Kullback Leibler (KL) distance to infer the structure of the song and label its different parts. The key phrase is then selected based on the frequency of those labels. Lu and Zhang [6] use the frequency, energy and position to detect the boundaries of musical phrases by analyzing each frame's estimated tempo and computing a confidence value of the frame being a phrase boundary. Depending on the type of music (instrumental or including vocals), Xu and Maddage [12] first extract the features that better catch the attributes of the segmented signal (e.g. MFCCs and amplitude envelope for instrumental music; linear prediction coefficients (LPCs) and derived cepstrum coefficients (LLPCs) [10] for vocal music). Those features are then used for content based clustering, and the output is used for the extraction of the most representative theme. Kim et al. [4] take changes in tempo as a primary indicator for summarization. They first segment the signal based on changes in tempo and then cluster segments based on their MFCCs. Shao et al. [11] analyze a song's structure based on the rhythm and note the onset of the signal and then cluster the segments according to their melody-based (chord contours) and content-based (chord contours and vocal content) similarity. The earliest segments containing the chorus together with some directly preceding and succeeding phrases are used for the creation of the final summary. Mardirossian and Chew [7] generate music thumbnails using the sequence of the keys in time and the average time in each key to detect the most prominent melody.

Peeters et al. [9] generate a state representation of the song to discover its structural components. After discovering the potential states of the signal, they apply k-means clustering to associate each frame to one of the discovered states and a Hidden Markov Model (HMM) to identify the state sequence. The state representation is then used for the creation of the summary by choosing states and transitions according to user needs. They describe four different possible ways to generate a multi-phrase summary based on the signal analysis.

As described, most of the work on music summarization has focused on the identification of music phrases. Our work is complementary in that it explores the design of multi-phrase summaries once phrases have been identified.

3. ALGORITHMS

Augmenting the refrain by compositing music phrases that are repeated in the music yet significantly different from one another can enhance the value of a summary. There are many examples where frequently occurring phrases other than the refrain are effective for recognizing a song. A highly repeated instrumental motif or a dominant verse can be as characteristic as the most salient phrase of a melody. In Lynyrd Skynyrd's "Sweet Home Alabama", for example, the introductory theme, which appears several times in the song, is almost as recognizable as the refrain itself. To explore how choices in the selection of additional phrases affects users perceptions of the summary, our summaries consist of three parts: the most salient phrase (usually the refrain) and two additional phrases. We compare three algorithms for selecting the two additional phrases. These algorithms vary the bias between phrases that are repeated and phrases that are sonically distinct.

3.1 Most Salient Phrase Detection

Phrase detection is not the focus of our work and, indeed, many of the algorithms found in related work could be used instead to identify phrase boundaries and determine repetitions. All three of our algorithms follow a common approach for the detection of the most salient (or key) phrase. In the preprocessing phase, the signal, after the removal of its first and last 10 seconds (that often carry non-useful information), is segmented into fixed, non overlapping blocks of 0.75 second each. A Hamming window is applied on each block to prepare the signal for the Fast Fourier Transform (FFT) which in turn returns the frequency components of the signal. Afterwards, the algorithm calculates the MFCCs of each block as they provide a better estimation of how humans perceive frequencies.

In the next phase, groups of eight successive blocks are formed where successive groups have a 50% overlap (i.e., 4 blocks). For each group we determine its MFCCs by taking the average of the MFCCs of its blocks. The Euclidean distance between the MFCCs of each pair of groups is then calculated and normalized. Starting with a strict (restrictive) distance threshold, clusters are computed using each group as a centroid. The largest resulting cluster is then selected. Clusters that include only

contiguous segments of the music are not considered. If the threshold is too strict to generate any non-contiguous clusters, the process repeats with a more relaxed threshold.

Once the largest cluster is identified, the key-phrase is selected by identifying the block with the smallest amplitude (lowest sound level) within a range of eight blocks (6 seconds) before the starting block of each group in the cluster. The group with the smallest corresponding amplitude is selected due to the likelihood that the block is near the start of a music phrase. The start of the key-phrase is chosen to be 3 seconds prior to the selected group and the key-phrase lasts for 8 seconds.

The next subsection presents a high-level description of the three algorithms for selecting the complementary parts of the summary. This is followed by a more detailed description of how the algorithms are instantiated.

3.2 Complementary Parts Selection (Overview)

The three algorithms presented here vary the selection of the two complementary parts (segments or clusters) of the summary based on a combination of the segments' musical similarity (distance between MFCCs), the number of identified repetitions (size of cluster), and the temporal location in the musical piece. Conceptually, the first algorithm follows an approach oriented more in finding complementary parts according to their frequency of occurrence in the song. In comparison, the second algorithm increases the importance of the sonic distance in the selection process while the third algorithm places most of the emphasis on the sonic distance.

The first algorithm (Repetition Emphasis Algorithm - REA) selects the complementary phrases by placing an upper bound on the similarity between the three phrases but otherwise picks the most repeated phrases prior to and after the identified key phrase. The second algorithm (Intermediate Algorithm - IA) again selects the first complementary phrase by selecting the most repeated phrase prior to the key phrase that differs by more than a threshold. It selects the second complementary phrase to maximize the minimum of 1) the similarity between the second phrase and the refrain and 2) the similarity between the second phrase and the first selected phrase. In this way, the IA puts a higher precedence on ensuring difference between all three of the selected musical segments than it puts on the second phrase's repetition. The third algorithm (Sonic Difference Emphasis Algorithm - SDEA) goes a step further by selecting the two complementary segments that minimize the musical similarity between the three segments without considering whether the complementary phrases were repeated or not.

3.3 Complementary Parts Selection (Details)

The first two algorithms share their approach to selecting the first complementary part. They also steer the selection of the first and second complementary parts towards earlier and later potions of the song, respectively.

After the selection of the key-phrase from the largest cluster of blocks, the first complementary part is selected from the next largest cluster that resides, if possible, in the interval between the start of the song and the key-phrase and differs by more than a minimum threshold. The difference between the two clusters is the mean distance between the MFCCs of its groups. To be a candidate for selection of a complementary part, the mean distance must be greater than a predefined threshold and the variance of the distances must be lower than a specific limit. These thresholds reduce the likelihood that the algorithm will choose a cluster of phrases that sound very similar to the key phrase (i.e. the refrain without the voice or a variation of it). Once the next-largest cluster that meets the MFCC distance requirements has been found, the group of blocks that occurs prior to the key-phrase and is temporally most distant from the key-phrase is chosen as the first complementary part. If no groups of blocks are prior to the key-phrase, then the first complementary part is chosen to be the one closest to the end of the song (furthest from the key-phrase).

The selection of the second complementary part in the REA proceeds similarly. Again, the algorithm selects the next largest cluster with significant differences in MFCC means from both the key-phrase and the first complementary part. Once the cluster is identified, the group of blocks closest to the end of song is selected (assuming the first complementary part was chosen from before the key-phrase, otherwise it will select the group of blocks closest to the start of the song).

In the IA, the selection of the first complementary part uses the technique described in the first algorithm. However, the selection process of the second complementary part deviates significantly except that the search is still focused on the interval between the key-phrase and the end of the song. The second complementary part is chosen as the group that maximizes the minimum of the value of F(i) in formula (1):

$$F(i) = \min (DK(i), DP(i)) \text{ where} \quad (1)$$
$$DK(i) = \Delta (V(group_i), V(\text{key-phrase})) \quad (2)$$
$$DP(i) = \Delta (V(group_i), V(\text{first complementary part})) \quad (3)$$

where $\Delta(V_1,V_2)$ is the Euclidean distance of the MFCC vectors V_1,V_2 and i is the number of the groups for the portion of the song being examined.

The SDEA differs substantially. After the extraction of the key-phrase, the ten least similar groups of blocks (in terms of MFCC vector similarity) to the key-phrase group of blocks are used as candidates for the selection of the phrases. From the ten candidates, the pair with the minimum similarity (maximum Euclidean distance) is used for the extraction of the two complementary parts. Thus, this algorithm places greater emphasis on sonic difference.

3.4 Summary Creation

To create the final summary, we take an eight-second slice from the key phrase and a six-second slice from each of the two complementary parts (total twenty seconds) and order them temporally. A one-second silence is introduced between the segments to diminish the effects of the abrupt switches. Fading in and out is not currently used since it "steals" potentially valuable time from the summary. However, the evaluation indicated that smoothing the transitions between phrases is important to users so crossfades will be part of our future efforts. The final summary has length twenty two seconds which is similar to many of the commercial summaries found.

4. STUDY DESIGN

We designed an experimental study to evaluate and compare our three summarization approaches and to test their performance over a widely used technique. The study was conducted in the Center for the Study of Digital Libraries at Texas A&M University. Fifteen participants over 18 years old, mainly students, were recruited to take part including 12 men and 3 women. The majority (67%) had some kind of music education and more than half of them (67%) had a personal music collection of at least 50 songs (8 participants had more than 200 songs).

Participants were asked to listen carefully to the summaries of twenty popular rock and pop songs and choose the summary that best represents each song. In contrast to the study conducted by Ong [8], our evaluation criteria focused on user preference and summary completeness and not on metrics measuring the ability of subjects to assign song titles. There were four 22 second-summaries per song, three generated with our algorithms and one that was merely the first 22 seconds of the song. Participants answered a series of multiple-choice questions about the quality of the selected summary and their familiarity with the song before proceeding to the next one. The summaries, as well as the songs (in their full version), were accessible through a web-based interface. Participants were able to navigate through the songs and listen to the summaries as many times they wanted. There was no time limit for the completion of the task. The order in which the songs and the summaries were presented to the participants was balanced across participants.

Demographic data about the participants was collected via a pre-task questionnaire. Post-task, semi-structured interviews were used to gather information about the participants' perceptions of the task, their experience with the algorithms and their ideas for future improvements.

5. STUDY RESULTS

To get an idea of what users really appreciate in a music summary we asked them to name (pre-task questionnaire) the parts or features of songs they consider fundamental for becoming familiar with and recalling music.

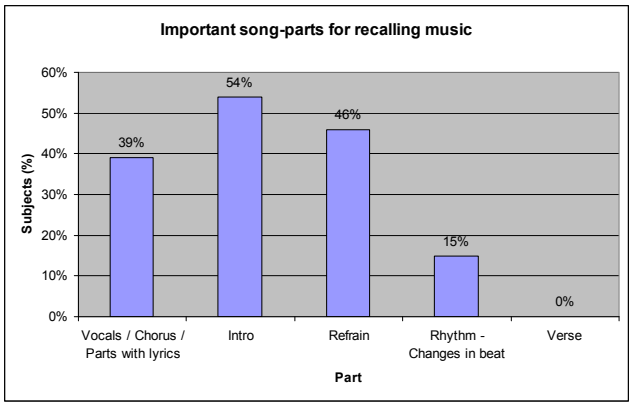

Figure 1. Important parts for recalling music

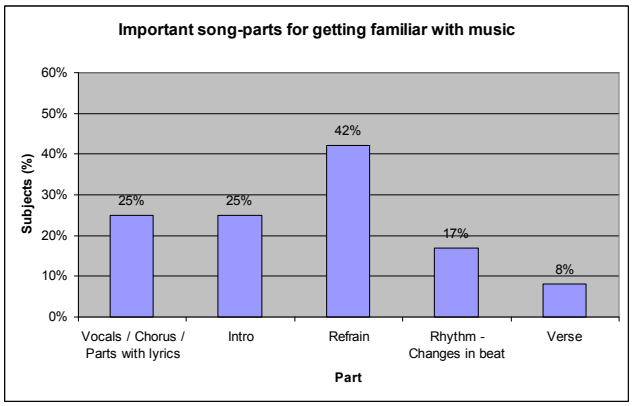

Figure 2. Important parts to become familiar with music

The results confirmed that both the introduction and the refrain are believed to have an important role in the process of understanding and recognizing music (see Figures 1 & 2). However, there was a distinction between the two cases. The introduction of the song was indicated most important for remembering (although with small difference from the refrain that was second) while the refrain was ranked best for becoming acquainted with the music. The higher score of the introduction and the vocals / chorus / lyrics in recalling music matches the fact that a few words or notes can be sufficient for identifying a song we know but provide little information for a song we do not know. Finally, other musical parts like bridges and verses scored very low in the preference ranking or were not mentioned at all by the participants.

Figure 3 shows the distribution of participants' selections for their favored summary. Participants chose the introduction summary in only 13% of the cases and the REA, the most popular, in 35% of the cases. Analysis of the data shows the difference in the selection of the four

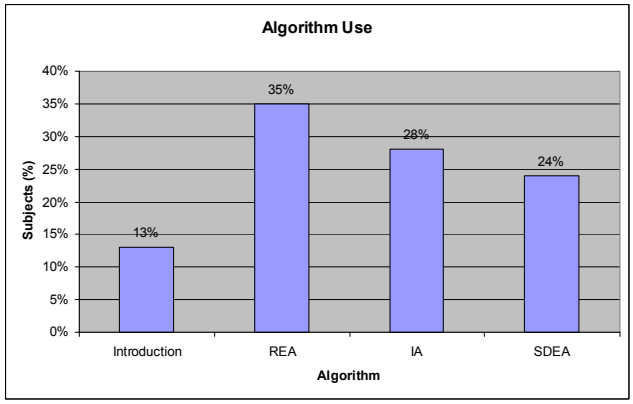

Figure 3. Users' algorithm choices

(I) Algorithm	(J) Algorithm	Mean Diff. (I-J)	Std. Error	Sig.
Introduction	REA	-4.4286	1.03067	**.001**
	IA	-2.8571	1.03067	**.041**
	SDEA	-2.1429	1.03067	.178
REA	Introduction	4.4286	1.03067	**.001**
	IA	1.5714	1.03067	.433
	SDEA	2.2857	1.03067	.136
IA	Introduction	2.8571	1.03067	**.041**
	REA	-1.5714	1.03067	.433
	SDEA	.7143	1.03067	.899
SDEA	Introduction	2.1429	1.03067	.178
	REA	-2.2857	1.03067	.136
	IA	-.7143	1.03067	.899

Table 1. Pairwise comparison of the four algorithms

algorithms was statistically significant (F-test, P=0.0013, α=0.05). Table 1 presents the results from the pairwise comparison of the algorithms with the Tukey HSD test. The numbers show a statistically significant difference between the introduction-based algorithm and the REA and IA (P=0.001 and P=0.041 respectively, α=0.05).

While there was no statistically significant difference between the three multi-phrase algorithms, the trends in the data show that identifying repeated phrases is likely to add value to the resulting multi-phrase summary.

The correlation between the algorithm choice and how good the summaries were (see Figure 4) wasn't statistically significant. However, the numbers look promising considering that 13 of the participants evaluated their choice as at least a good representation of the song, and this choice was one of our algorithms in 87% of the songs. However, the most interesting point about the weakness of the song introduction as summary came out from the post-task interviews. The analysis showed that, while a respective number of the participants (10) listen to the song previews provided by the on-line music stores, only 3 of them are confident that these previews describe the songs sufficiently. Since most online stores preview music using a single contiguous snippet, this indicates a need for an alternative to current summaries.

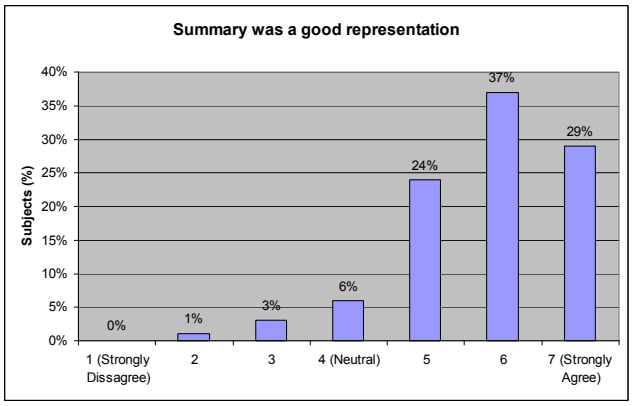

Figure 4. Evaluation of summaries performance

Algorithm \ Knowledge of Songs	I know it well	I don't know it well	I haven't heard the song
Introduction	32 (16%)	4 (11%)	1 (2%)
REA	66 (33%)	10 (28%)	23 (54%)
IA	53 (26%)	14 (39%)	10 (23%)
SDEA	50 (25%)	8 (22%)	9 (21%)

Table 2. Algorithm selection and familiarity with music

Participants reported knowing 71% of the songs well, not knowing 16% of the songs, and having limited knowledge of 13% of the songs. Analysis of the data showed that there is no statistically-significant correlation between the choice of the algorithm and how familiar the participants were with the music. Table 2 shows the number of times participants selected each algorithm based on their knowledge of the song. The proposed techniques are superior in effectiveness over the traditional introduction-based approach no matter whether the user is a customer browsing a new album or a person trying to retrieve an already known mp3 from a personal collection.

Strongly related to participants' knowledge of the songs is how recently they had heard them. Participants had listened on average to about 59% of the songs within a year (see Figure 5). However, the statistics again did not show a significant correlation with the algorithm choice.

One of our concerns when we were designing the proposed summarization techniques was that the segments in each summary would be too short to sufficiently describe the section of the song that had been extracted. A music phrase (especially in classical music) can have duration much longer than the six seconds selected as the length for complementary parts. However, for this collection of pop and rock songs, the results showed that only 20% of the selected summaries were considered "too short" while 53% evaluated were considered "good" and 27% were viewed as "too long".

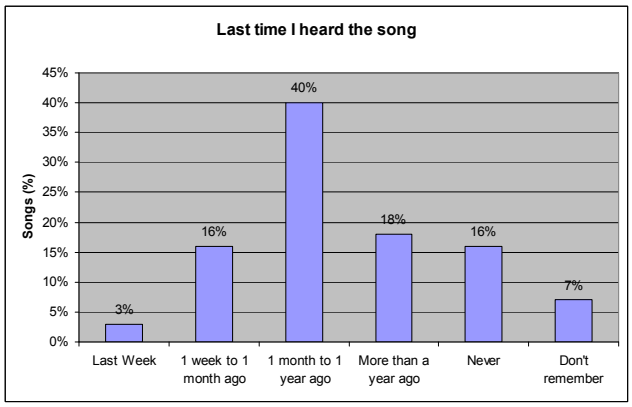

Figure 5. Participants' last time of listening to the song

6. DISCUSSION

Three algorithms for creating multi-phrase music summaries in are presented and evaluated. The design of the algorithms reflects a range of approaches that vary between emphasizing the selection of repeated phrases and the selection of sonically different phrases. The study showed that participants believed that the multi-phrase summaries better represented the song than the introduction to the song. While the difference between the three algorithms was not significant, the results indicate a likely preference for algorithms that emphasize the selection of repeated phrases, at least in the genre of pop and rock where the structural components of the melody are more standardized and identifiable.

There are several things we want to test and improve about our algorithms in the future. One of the complaints participants had during the task was that the switch from part to part in the summaries was too abrupt and hence distractive or even annoying. Use of phrase bounds detection for selecting the start of phrases could help as could the use of fade-in and fade-out effects. Based on participants' feedback about which parts / features of the songs are important, it would be interesting to examine if the integration of the introduction in our summaries can improve or accelerate the process of becoming familiar with new music. A comparison of the best of our techniques with summaries containing only the most salient phrase of the song is also in our future plans. Finally, we would like to improve the accuracy of our summarization approach to be applicable in genres like classical music and jazz where identification of the various themes and important components is more challenging.

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HYBRID NUMERIC/RANK SIMILARITY METRICS FOR MUSICAL PERFORMANCE ANALYSIS

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ABSTRACT

This paper describes a numerical method for examining similarities among tempo and loudness features extracted from recordings of the same musical work and evaluates its effectiveness compared to Pearson correlation. Starting with correlation at multiple timescales, other concepts such as a performance "noise-floor" are used to generate measurements which are more refined than correlation alone. The measurements are evaluated and compared to plain correlation in their ability to identify performances of the same Chopin mazurka played by the same pianist out of a collection of recordings by various pianists.

1 INTRODUCTION

As part of the Mazurka Project at the AHRC Centre for the History and Analysis of Recorded Music (CHARM), almost 3,000 recordings of Chopin mazurkas were collected to analyze the stylistic evolution of piano playing over the past 100 years of recording history, which equates to about 60 performances of each mazurka. The earliest collected performance was recorded on wax cylinders in 1902 and the most recent posted as homemade videos on YouTube. Table 1 lists 300 performances of five mazurkas which will be used for evaluation later in this paper since they include a substantial number of recordings with extracted tempo and loudness features.

Mazurka		*Performances*	
Opus	*Key*	*Collected*	*Processed*
17/4	A minor	93	63
24/2	C major	63	29
30/2	B minor	60	33
63/3	C♯ minor	87	57
68/3	F major	51	50

Table 1. Collection of musical works used for analysis.

For each of the processed recordings, beat timings in the performance are determined using the *Sonic Visualiser* audio editor [1] for markup and manual correction with the assistance of several vamp plugins. [2] Dynamics are then extracted as smoothed loudness values sampled at the beat positions.[3] Feature data will eventually be extracted from all collected mazurkas in the above list, but comparisons made in Section 3 are based on the processed performance counts in Table 1. Raw data used for analysis in this paper is available on the web. [3]

Figure 1 illustrates extracted performance feature data as a set of curves. Curve 1a plots the beat-level tempo which is calculated from the duration between adjacent beat timings in the recording. For analysis comparisons, the tempo curve is also split into high- and low-frequency components with linear filtering. [4] Curve 1b represents smoothed tempo which captures large-scale phrasing architecture in the performance (note there are eight phrases in this example). Curve 1c represents the difference between Curves 1a and 1b which is called here the desmoothed tempo curve, or the residual tempo. This high-frequency tempo component encodes temporal accentuation in the music used by the performer to emphasize particular notes or beats. Mazurka performances contain significant high-frequency tempo information, since part of the performance style depends on a non-uniform tempo throughout the measure—the first beat usually shortened, while the second and/or third beat are lengthened. Curve 1d represents the extracted dynamics curve which is a sampling of the audio loudness at each beat location.

Other musical features are currently ignored here, yet are important in characterizing a performance. In particular, pianists do not always play left- and right-hand notes together, according to aural traditions, although they are written as simultaneities in the printed score. Articulations such as legato and staccato are also important performance features but are equally difficult to extract reliably from audio data. Nonetheless, tempo and dynamic features are useful for developing navigational tools which allow listeners to focus their attention on specific areas for further analysis.

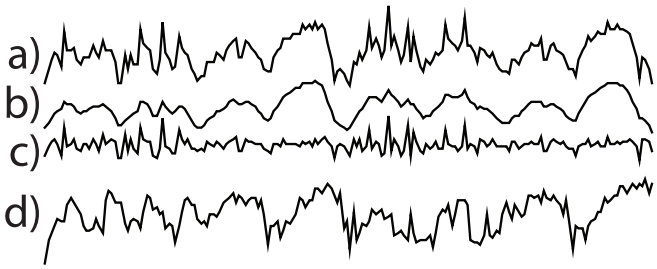

Figure 1. Extracted musical features from a recording of Chopin's mazurka in B minor, 30/2: a) tempo between beats; b) smoothed tempo; c) residual tempo ($c = a - b$); and d) beat-level dynamics.

[1] http://www.sonicvisualiser.org
[2] http://sv.mazurka.org.uk/download
[3] http://mazurka.org.uk/info/excel
[4] The filtering method is available online at http://mazurka.org.uk/software/online/smoother .

2 DERIVATIONS AND DEFINITIONS

Starting with the underlying comparison method of correlation (called S_0 below), a series of intermediate similarity measurements (S_1, S_2, and S_3) are used to derive a final measurement technique (S_4). Section 3 then compares the effectiveness of S_0 and S_4 measurements in identifying recordings of the same performer out of a database of recordings of the same mazurka.

2.1 Type-0 Score

As a starting point for comparison between performance features, Pearson correlation, often called an r-value in statistics, is used:

$$\text{Pearson}(x, y) = \frac{\sum_n (x_n - \bar{x})(y_n - \bar{y})}{\sqrt{\sum_n (x_n - \bar{x})^2 \sum_n (y_n - \bar{y})^2}} \quad (1)$$

This type of correlation is related to dot-product correlation used in Fourier analysis, for example, to measure similarities between an audio signal and a set of harmonically related sinusoids. The value range for Pearson correlation is -1.0 to $+1.0$, with 1.0 indicating an identical match between two sequences (exclusive of scaling and shifting), and the value 0.0 indicating no predictable linear relation between the two sequences x and y.

Correlation values between extracted musical features typically have a range between 0.20 and 0.97 for different performances of mazurkas. Figure 2 illustrates the range of correlations between performances in two mazurkas. Mazurka 17/4 is a more complex composition with a more varied interpretation range, so the *mode*, or most-expected value, of the correlation distribution is 0.67. Mazurka 68/3 is a simpler composition with fewer options for individual interpretations so the mode is much higher at 0.87.

These differences in expected correlation values between two randomly selected performances illustrate a difficulty in interpreting similarity directly from correlation values. The correlation values are consistent only in relation to a particular composition, and these absolute values cannot be compared directly between different mazurkas. For example, a pair of performances which correlate at 0.80 in mazurka 17/4 indicates a better than average match, while the same correlation value in mazurka 68/3 would be a relatively poor match. In addition, correlations at different timescales in the same piece will have a similar problem, since some regions of music may allow for a freer interpretation while other regions may have a more static interpretation.

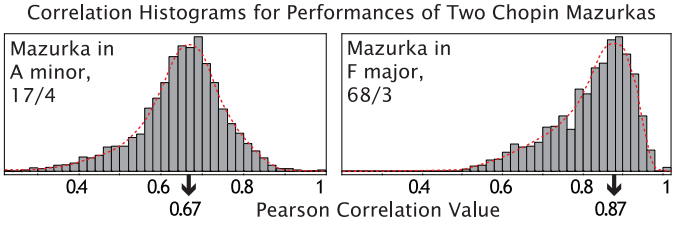

Figure 2. Different compositions will have different expected correlation distributions between performances.

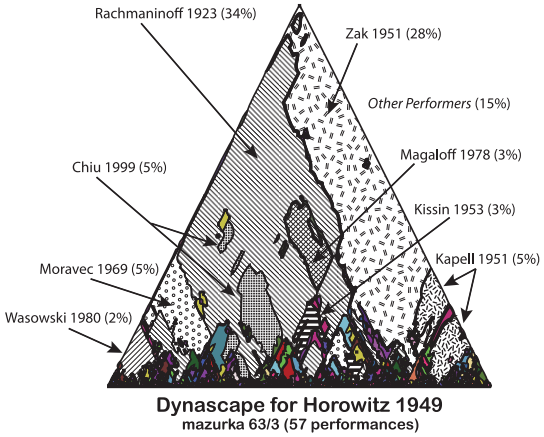

Figure 3. Scapeplot for the dynamics in Horowitz's 1949 performance of mazurka 63/3 with the top eight matching performances labeled.

2.2 Type-1 Score

In order to compensate partially for this variability in correlation distributions, scapeplots were developed which only display nearest-neighbor performances in terms of correlation at all timescales for a particular reference performance.[3] Examples of such plots created for the Mazurka Project can be viewed online. [5]

The S_1 score is defined as the fraction of area each target performance covers in a reference performer's scapeplot. Figure 3 demonstrates one of these plots, comparing the dynamics of a performance by Vladimir Horowitz to 56 other recordings. In this case, Rachmaninoff's performance of the same piece matches better than any other performance, since his performance covers 34% of the scape's plotting domain. At second best, Zak's performance matches well towards the end of the music, but covers only 28% of the total plotting domain. Note that Zak's performance has the best correlation for the entire feature sequence (S_0 score) which is represented by the point at the top of the triangle. The S_1 scores for the top eight matches in Figure 3 are listed in Table 2. There is general agreement between S_1 and S_0 scores since five top-level correlation matches also appear in the list.

2.3 Type-2 Score

Scape displays are sensitive to the *Hatto effect*: if an identical performance to the reference, or query, performance is present in the target set of performances, then correlation values at all time resolutions will be close to the maximum value for the identical performance, and the comparative scapeplot will show a solid color. All other performances would have an S_1 score of approximately 0 regardless of how similar they might otherwise seem to the reference performance. This property of S_1 scores is useful for identifying two identical recordings, but not useful for viewing similarities to other performances which are hidden behind such closely neighboring performances.

One way to compensate for this problem is to remove the best match from the scape plot in order to calculate the next best match. For example, Figure 4 gives a rough schematic for

[5] http://mazurka.org.uk/ana/pcor-perf

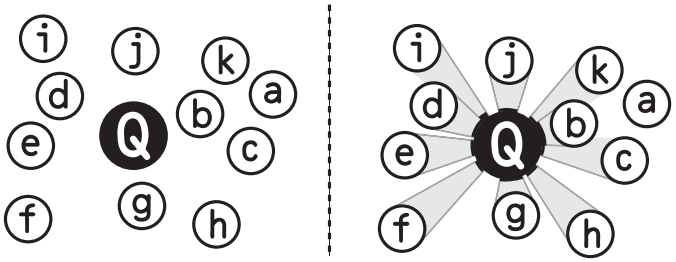

Figure 4. Schematic of nearest-neighbor matching method used in comparative timescapes.

how scapeplots are generated. Capital 'Q' represents the query, or reference, performance, and lower-case lettered points represent other performances. The scapeplot basically looks around in the local neighborhood of the feature space and displays closest matches as indicated by lines drawn towards the query on the right side of Figure 4. Closer performances will tend to have larger shadows on the query, and some performances can be completely blocked by others, as is the case for point "*a*" in the illustration.

An S_2 score measures the coverage area of the most dominant performance which is assumed to be most similar to the reference performance. This nearest of the neighbors is then removed from the search database, and a new scapeplot is generated with the remaining performances. Gradually, more and more performances will be removed which allows for previously hidden performances to appear in the plot. For example, point "*a*" in Figure 4 will start to become visible once point "*b*" is removed. S_2 scores and ranks are independent. As fewer and fewer target matches remain, the S_2 scores will increase towards 1.0 while the S_2 ranks decrease towards the bottom match.

In Figure 3, Rachmaninoff is initially the best match in terms of S_1 scores, so his performance will be removed and a new scapeplot generated. When this is done, Zak's performance then represents the best match, covering 34% of the scapeplot. Zak's performance is then removed, a new scapeplot is calculated, and Moravec's performance will have the best coverage at 13%, and so on. Some of the top S_2 scores for Horowitz's performance are listed in Table 2.

2.4 Type-3 Score

Continuing on, the next best S_2 rank is for Chiu, who has 20% coverage. Notice that this is greater than Moravec's score of 13%. This demonstrates the occurrence of what might be called the lesser Hatto effect: much of Moravec's performance overlapped onto Chiu's region, so when looking only at the nearest neighbors in this manner, there are still overlap problems. Undoubtedly, Rachmaninoff's and Zak's performances mutually overlap each other in Figure 3 as well. Both of them are good matches to Horowitz, so it is difficult to determine accurately which performance matches "best" according to S_2 scores since they are interfering with each others scores and are both tied at 34% coverage.

In order to define a more equitable similarity metric and remove the Hatto effect completely, all performances are first ranked approximately by similarity to Horowitz using either

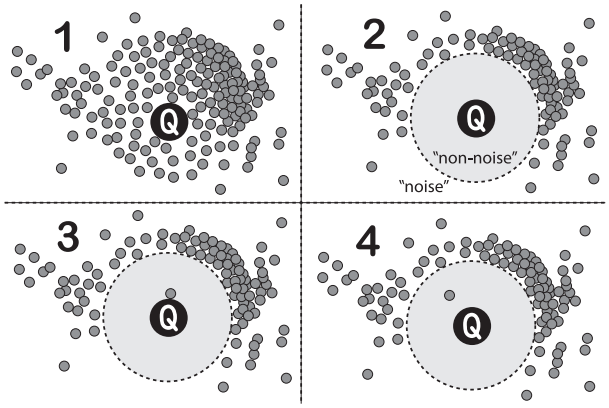

Figure 5. Schematic of the steps for measuring an S_3 score: (1) sort performances from similar to dissimilar, (2) remove most similar performance to leave noise floor, (3) & (4) insert more similar performances one-by-one to observe how well they can occlude the noise-floor.

S_0 values or the rankings produced during S_2 score calculations. Performances are then divided into two groups, with the poorly-matching half defined as the performance "noise-floor" over which better matches will be individually placed.

To generate an S_3 score, the non-noise performances are removed from the search database as illustrated in step 2 of Figure 5, leaving only background-noise performances. Next, non-noise performances are re-introduced separately along with all of the noise-floor performances and scapeplot is generated. The coverage area of the single non-noise performance represented in the plot is defined as its S_3 similarity measurement with respect to the query performance.

This definition of a performance noise-floor is somewhat ar-

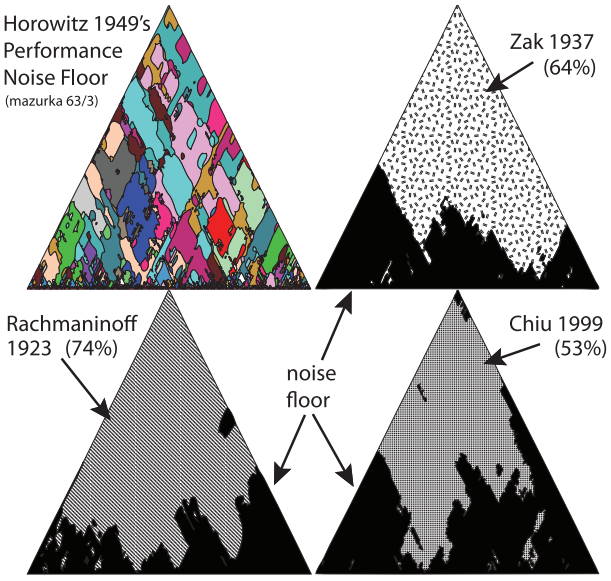

Figure 6. Dynascapes for Horowitz's performance of mazurka 63/3. Top left is a plot of the noise-floor performances, and the other three plots separate include one of the top matching performances which can cover most of the noise-floor.

Target	S_0	R_0	S_1	S_2	S_3	S_{3r}	S_4	R_4
Rac23	0.60	3	0.34	0.34	0.74	0.82	0.78	1
Zak37	0.64	1	0.28	0.34	0.64	0.60	0.62	2
Mor69	0.59	4	0.05	0.13	0.57	0.54	0.55	3
Chi99	0.49	20	0.05	0.20	0.53	0.54	0.53	4
Kap51	0.51	17	0.05	0.08	0.24	0.17	0.20	22
Mag78	0.62	2	0.03	0.27	0.59	0.37	0.47	7
Kis93	0.52	15	0.03	0.09	0.44	0.23	0.32	11
Was80	0.58	5	0.02	0.11	0.41	0.55	0.47	6

Table 2. Scores and rankings for sample targets to Horowitz's 1949 performance of mazurka 63/3.

Query	*Target*	*Tempo*			T_S			T_d			*Dynamics*			*TD*		
		R_0	R_3	R_4	R_0	R_3	R_4	R_0	R_3	R_4	R_0	R_3	R_4	R_0	R_3	R_4
Rub39	Rub52	2	3	2	3	8	8	3	2	1	6	8	8	3	3	2
Rub39	Rub66	1	1	1	1	1	1	2	3	2	4	4	2	1	2	1
Rub52	Rub39	6	9	2	27	31	12	3	3	2	28	23	6	12	9	2
Rub52	Rub66	2	2	1	3	2	1	2	2	1	2	2	1	2	2	1
Rub66	Rub39	3	4	2	2	3	1	3	6	5	15	8	6	5	5	3
Rub66	Rub52	1	2	1	3	2	2	2	2	1	1	3	2	1	2	1
Ranking Averages (by feature)	R_0	2.5			6.5			2.5			9.3			4.0		
	R_3	3.5			7.8			3.0			8.0			3.8		
	R_4	1.5			4.2			2.0			4.2			1.7		
Overall Averages:		$R_0 = 4.97$			$R_3 = 5.32$			$R_4 = 2.70$								

Table 3. Rankings for 17/4 Rubinstein performances. Shaded numbers indicate perfect performance of a similarity metric.

bitrary but splitting the performance database into two equal halves seems the most flexible rule to use, and is used for the evaluation section later in this paper. But the cut-off point could be a different percentage, such as the bottom 75% of ranked scores, or an absolute cut-off number. In any case, it is preferable that the noise floor does not appear to have any favored matches, and should consist of uniform small blotches at all timescales in the scapeplot representing many different performers as is the example shown in Figure 6 (top left part of the figure). While Rachmaninoff and Zak have equivalent S_2 scores, Rachmaninoff's performance is able to cover 74% of the noise-floor, while Zak's is only able to cover 64%.

2.5 Type-4 Score

Type-3 scores require one additional refinement in order to be useful since performances are not necessarily evenly distributed in the feature space. The plots used to calculate the S_3 scores are still nearest-neighbor rank plots, so the absolute numeric distances between performances are not directly displayed. Unlike correlation values between two performances, S_3 scores are not symmetric: the score from *A* to *B* is not the same value as from *B* to *A*. It is possible for an outlier performance to match well to another performance closer to the average performance just because it happens to be facing towards the outlier, with the similarity just being a random coincidence.

Therefore, the geometric mean is used to mix the S_3 score with the reverse-query score (S_{3r}) as shown in Equation 4.

$$S_3 = A \Rightarrow B \text{ measurement} \quad (2)$$

$$S_{3r} = A \Leftarrow B \text{ measurement} \quad (3)$$

$$S_4 = \sqrt{S_3\, S_{3r}} \quad (4)$$

The arithmetic mean could also be used, but the geometric mean is useful since it penalizes the final score if the type-3 and its reverse scores are not close to each other. For example, the arithmetic mean between 0.75 and 0.25 is 0.50, while the geometric mean is lower at 0.43. Greatly differing S_3 and S_{3r} scores invariably indicate a poor match between two performances, with one of them acting as an outlier to a more central group of performances.

Table 2 shows several of the better matches to Horowitz's performance in mazurka 63/3, along with the various types of scores that they generate. S_0 is the top-level correlation between the dynamics curves, and R_0 is the corresponding similarity rankings generated by sorting S_0 values. Likewise, S_4 and R_4 indicate the final proposed similarity metric, and the resulting rankings generated by sorting these scores.

3 EVALUATION

When evaluating similarity measurement effectiveness, a useful technique with a clear ground-truth is to identify recordings by the same performer mingled among a larger collection of recordings.[5] Presumably pianists will tend to play more like their previous performances over time rather than like other pianists. If this is true, then better similarity metrics should match two performances by the same pianist more closely to each other than to other performances by different pianists.

3.1 Rubinstein performance matching

Arthur Rubinstein is perhaps the most prominent interpreter of Chopin's compositions in the 20th century, and luckily he has recorded the entire mazurka cycle three times during his career: (1) in 1938–9, aged 51; (2) in 1952–4, aged 66, and (3) in 1966, aged 79.

Table 3 lists the results of ranking his performances to each other in mazurka 17/4 where the search database contains an additional 60 performances besides the three by Rubinstein. The first column in the table indicates which performance was used as the query (Rubinstein's 1939 performance, for example, at the top of the first row). The *target* column indicates a particular target performance which is one of the other two performances by Rubinstein in the search database. Next, five columns list three types of rankings for comparison. The five columns represent four different extracted features as illustrated in Figure 1, plus the *TD* column which represents a 50/50 admixture of the tempo and dynamics features.

For each musical feature, three columns of rankings are reported. R_0 represents the rankings from the S_0 scores; R_3 being the type-3 scoring ranks, and R_4 resulting from sorting the S_4 similarity values. In these columns, a "1" indicates that the target performance was ranked best in overall similarity to the query performance, "2" indicates that it is the second best match, and so on (see the search database sizes in Table 1). In the ranking table for mazurka 17/4 performances of Rubinstein, the shaded cells indicate perfect performance matches by a particular similarity metric where the top two matches are both Rubinstein. Note that there is one perfect pair of matches in all of the R_0 columns which is found in the full-tempo feature

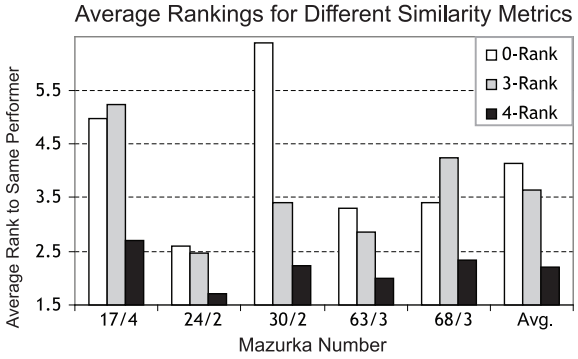

Figure 7. Ability of metrics to identify the same performer in a larger set of performances, using 3 performances of Rubinstein for each mazurka. (Lower rankings indicate better results.)

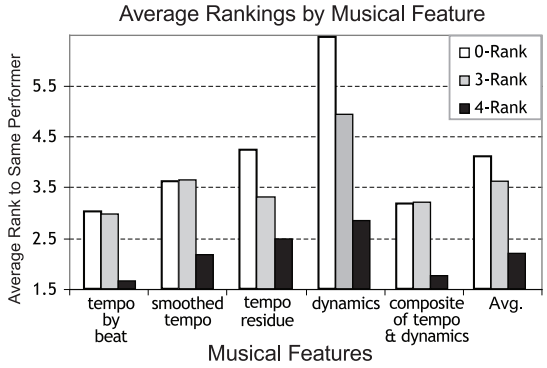

Figure 8. Ranking effectiveness by extracted musical features, using three performances of Rubinstein for each mazurka. (Lower values indicate better results.)

when Rubinstein 1939 is the target performance. No columns for R_3 contain perfect matching pairs, but about 1/2 of the R_4 columns contain perfect matches: all of the full-tempo R_4 rankings are perfect, and a majority of the desmoothed tempo and joint tempo/dynamics rankings are perfect. None of the metrics contain perfect matching pairs for the dynamics features. This is perhaps due to either (1) the dynamics data containing measurement noise (due to the difficultly of extracting dynamics data from audio data), or (2) Rubinstein varying his dynamics more over time than his tempo features, or a combination of these two possibilities.

Figure 7 shows the average rankings of Rubinstein performances for all extracted features averaged by mazurka. The figure shows S_4 scores are best at identifying the other two Rubinstein performances for all of the five mazurkas which were used in the evaluation. Typically S_4 gives three to four times better rankings than the S_0 values according to this figure. S_3 scores (used to calculate S_4 scores), are slightly better than plain correlation, but sometimes perform worse than correlation in some mazurkas.

Figure 8 evaluates the average ranking effectiveness by musical feature, averaged over all five mazurkas. Again S_4 scores are always three to four times more effective than plain correlation. S_3 scores are approximately as effective as S_0 rankings for full and smoothed tempo, but perform somewhat better on residual tempo and dynamics features, probably by minimizing the effects of sudden extreme differences between compared feature sequences caused by noisy feature data.

Mazurka	*Query*	*Target*	T R_0	T R_4	T_s R_0	T_s R_4	T_d R_0	T_d R_4	D R_0	D R_4	TD R_0	TD R_4
17/4	Cze49	Cze49b	1	1	1	1	1	1	3	1	1	1
17/4	Cze49b	Cze49	1	1	1	1	1	1	7	1	1	1
63/3	Fri23	Fri30	1	1	1	1	1	1	1	1	1	1
63/3	Fri30	Fri23	1	1	1	1	1	1	1	1	1	1
17/4	Hor71	Hor85	2	1	13	1	2	1	1	1	2	1
17/4	Hor85	Hor71	1	1	1	1	1	1	1	1	1	1
30/2	Fou78	Fou05	2	1	1	1	13	17	2	2	2	1
30/2	Fou05	Fou78	1	1	1	1	2	6	3	2	2	1
63/3	Uni32	Uni71	1	1	1	1	5	1	1	1	1	1
63/3	Uni71	Uni32	1	1	1	1	1	1	1	1	1	1

Table 4. Performer self-matching statistics.

3.2 Other performers

Rubinstein tends to vary his performance interpretation more than most other pianists. Also, other performers may tend to emulate his performances, since he is one of the more prominent interpreters of Chopin's piano music. Thus, he is a difficult case to match and is a good challenge for similarity metric evaluations.

This section summarizes the effectiveness of the S_0 and S_4 similarity metrics in identifying other pianists found in the five selected mazurkas for which two recordings by the same pianist are represented in the data collection (only Rubinstein is represented by three performances for all mazurkas).

Table 4 presents ranking evaluations for performance pairs in a similar layout to those of Rubinstein found in Table 3. In all except two cases (for Fou) the S_4 metrics perform perfectly in identifying the other performance by the same pianist. Top-level correlation was able to generate correct matches in 75% of the cases. An interesting difference between the two metrics occurs when Hor71 is the query performance. In this case S_0 yields a rank of 13 (with 12 other performance matching better than his 1985 performance), while S_4 identifies his 1985 performance as the closest match.

Fou's performance pair for mazurka 30/2 is also an interesting case. For his performances, the phrasing portion of the full- and smoothed-tempo features match well to each other, but the tempo residue does not. This is due to a significant change in his metric interpretation: the earlier performance has a strong mazurka metric pattern which consists of a short first beat, followed by a lengthened second or third beat in each measure. His 2005 performance greatly reduces this effect, and beat durations are more uniform throughout the measure in comparison to his 1978 performance.

Finally, it is interesting to note the close similarity between Uninsky's pair of performances listed in Table 4. These two performances were recorded almost 40 years apart, one in Russia and the other in Texas. Also, the first was recorded onto 78 RPM monophonic records, while the later was recorded onto 33-1/3 RPM stereo records. Nonetheless, his two performances indicate a continuity of performance interpretation over a long career.

feature	*Cortot ConA*		*Cortot Sony*	
	R_4	S_4	R_4	S_4
T	1. Rangell 2001	0.65	1. Luisada 1990	0.40
	2. Uninsky 1971	0.47	2. Ferenczy 1956	0.31
	3. Yaroshinsky 2005	0.44	3. Rubinstein 1966	0.27
	...		...	
	31. Cortot Sony	0.04	33. Cortot ConA	0.04
T_s	1. Pobłocka 1999	0.60	1. Lushtak 2004	0.27
	2. Luisada 1990	0.60	2. Milkina 1970	0.18
	3. Mohovich 1999	0.56	3. Fou 2005	0.18
	...		...	
	26. Cortot Sony	0.07	24. Cortot ConA	0.07
T_d	1. Rangell 2001	0.79	1. Luisada 1990	0.53
	2. Uninsky 1971	0.72	2. Rubinstein 1966	0.49
	3. Yaroshinsky 2005	0.57	3. Rubinstein 1939	0.47
	...		...	
	32. Cortot Sony	0.03	35. Cortot ConA	0.03
D	1. Brailowsky 1960	0.38	1. Fliere 1977	0.46
	2. Biret 1990	0.25	2. Sofronitsky 1960	0.42
	3. Milkina 1970	0.24	3. Ashkenazy 1981	0.40
	...		...	
	33. Cortot Sony	0.06	32. Cortot ConA	0.06
TD	1. Rangell 2001	0.40	1. Avg. performance	0.16
	2. Pobłocka 1999	0.37	2. Indjic 1988	0.16
	3. Yaroshinsky 2005	0.35	3. Rubinstein 1952	0.15
	...		...	
	35. Cortot Sony	0.03	34. Cortot ConA	0.03

Table 5. Comparison of Cortot performances of mazurka 30/2 (m. 1–48).

4 APPLICATION

As an example application of the derived similarity metrics, two performances of mazurka 30/2 performed by Alfred Cortot are examined in this section. One complete set of his mazurka performances can be found on commercially release recordings from a 1980's-era issue on cassette tape "recorded at diverse locations and dates presumably in the period of 1950–1952."[1] These recordings happen to be issued by the same record label as the recordings of Joyce Hatto, which casts suspicion on other recordings produced on that label.[4]. A peculiar problem is that no other commercial recordings exist of Cortot playing any mazurka, let alone the entire mazurka cycle.

In 2005, however, Sony Classical (S3K89698) released a 3-CD set of recordings by Cortot played during master classes he conducted during the late 1950's, and in this set, there are six partial performances of mazurkas by Cortot where he demonstrates how to play mazurkas to students during the class. His recording of mazurka 30/2 on these CDs is the largest continuous fragment, including 75% of the entire composition, stopping two phrases before the end of the composition.

Table 5 lists the S_4 scores and rankings for these two recordings of mazurka 30/2, with the Concert Artist's rankings on the left, and the Sony Classical rankings to the right. The five different musical features listed by column in previous tables for Rubinstein and other pianists are listed here by row. For each recording/feature combination, the top three matches are listed, along with the ranking for the complimentary Cortot recording.

Note that in all cases, the two Cortot recordings match very poorly to each other. In two cases, the worst possible ranking of 35 is achieved (since 36 performances are being compared in total). Perhaps Cortot greatly changes his performances style in the span of 6 years between these two recordings late in his life, although data from Tables 3 and 4 would not support this view since no other pianists has significantly alter all musical features at once, and only Fou significantly changes one musical feature between performances.

Therefore, it is likely that this particular mazurka recording on the Concert Artist label was not actually performed by Cortot. Results from further investigation of the other five partial mazurka performances on the Sony Classical recordings would help to confirm or refute this hypothesis, but the other examples are more fragmentary, making it difficult to extract reasonable amounts of recital-grade performance material. In addition, no performer in the top matches for the Concert Artist Cortot performance match well enough to have likely recorded this performance, so it is unlikely that any of the other 30 or so performers being compared to this spurious Cortot performance is the actual source for this particular mazurka recording.

5 FUTURE WORK

Different methods of combining S_3 and S_{3r} scores, such as measuring the intersection between plot areas rather than measuring the geometric mean to calculate S_4 should be examined. The concept of a noise floor when comparing multiple performances is useful for identifying features which are common or rare, and allows similarity measurements to be more consistent across different compositions, which may aid in the identification of pianists across *different* musical works.[6]

Further analysis of the layout of the noise-floor as seen in Figure 6 might be useful in differentiating between directly and indirectly generated similarities between performances. For example in this figure, Rachmaninoff's performance shows more consistent similarity towards smaller-scale features, which may indicate a direct influence on Horowitz's performance style. Zak's noise-floor boundary in Figure 6 may demonstrate an indirect similarity, such as a general school of performance.

Since the similarity measurement described in this paper works well for matching the same performer in different recordings, and examination of student/teacher similarities may be done. The analysis techniques described here should be applicable to other types of features, and may be useful with other underlying similarity metrics besides correlation. For example, It would be interesting to extract musical features first from the data with other techniques such as Principle Component Analysis[2] and use this derived feature data for characterizing the similarities between performances in place of correlation.

6 REFERENCES

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GAMERA *versus* ARUSPIX TWO OPTICAL MUSIC RECOGNITION APPROACHES

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ABSTRACT

Optical music recognition (OMR) applications are predominantly designed for common music notation and as such, are inherently incapable of adapting to specialized notation forms within early music. Two OMR systems, namely Gamut (a Gamera application) and Aruspix, have been proposed for early music. In this paper, we present a novel comparison of the two systems, which use markedly different approaches to solve the same problem, and pay close attention to the performance and learning rates of both applications. In order to obtain a complete comparison of Gamut and Aruspix, we evaluated the core recognition systems and the pitch determination processes separately. With our experiments, we were able to highlight the advantages of both approaches as well as causes of problems and possibilities for future improvements.

1 INTRODUCTION

Optical music recognition (OMR) enables document images to be encoded into digital symbolic music representations. The encoded music can then be used and processed within a wide range of applications, such as music analysis, editing and music information retrieval. Over the years, multiple approaches have addressed this difficult task, in some cases focusing exclusively on one type of music document, such as keyboard music, orchestral scores, or music manuscripts [1]. Most commercial tools today are non-adaptive, acting as black-boxes that do not improve their performance through usage: when a symbol is misread, it continues to be misread in the subsequent pages, even if the user corrects the results by hand on every page. Attempts have been made to remedy this situation by merging multiple OMR systems, yet this remains a challenge [4].

Two research projects have taken an innovative approach to OMR by adopting adaptive strategies, namely Gamut, built based on the Gamera framework, and Aruspix. The main idea behind the adaptive approach in OMR is to enable the tools to improve themselves through usage by taking benefits from the training data that becomes available as the user corrects recognition errors. Adaptive approaches have been proven to be very promising with historical documents, because the very high variability within the data requires the tools to constantly adapt and retarget themselves. This is particularly true for early music prints from the sixteenth and seventeenth centuries, for which Aruspix was designed, as they contain an unpredictable variability of font shape, page noise, brightness and contrast [14].

This study is a first attempt at a comparison between Gamut and Aruspix. We specifically address the accuracy of the tools and their capability to adapt themselves to a new dataset. This paper is structured as follows. In section 2 we present a brief overview of the two OMR applications in comparison. In section 3 we present the experiments undertaken to enable such a comparison at different stages of the recognition process. Results are presented in section 4, and conclusions and future work are discussed in section 5.

2 INFRASTRUCTURE

2.1 Gamera and Gamut

Gamera is an open-source framework for the creation of structured document analysis applications by domain experts [8]. This system is designed for use with any type of image documents, and provides tools for preprocessing, segmentation, and classification of symbols within a document. The environment, structured as a composite of C++ and Python, provides facilities for developers to integrate plugins or toolkits suited for the specific type of documents being analyzed. Its framework is accessible enough to provide the tools for the development of an application for a specific domain without requiring a strong programming background. Within a Gamera application, individual symbols may be extracted by connected-component (CC) analysis and recognized using the *k-nearest neighbour* (kNN) classifier. If the symbols themselves are split during CC analysis, the kNN classifier is capable of automatic regrouping and recognition, which has been shown to be a necessary feature when working with degraded documents [7].

Gamut is an OMR application built with the Gamera framework [10]. It is comprised of several plugins and toolkits that provide procedures specific to the task. As in most OMR systems, Gamut requires additional preprocessing to remove staff lines prior to symbol recognition, a task that is especially important for subsequent pitch detection from classified symbols [5]. Within Gamut, staff detection and removal is performed by the MusicStaves toolkit, a collection staff-removal algorithms specialized for specific types of musical documents.

2.2 Aruspix

Aruspix is a cross-platform software program for OMR of early music prints. Aruspix performs the entire OMR process, from preprocessing to score encoding. The preprocessing stage includes document deskewing, border removal, binarization, and a heuristic approach to preclassify each region of the image as text, music, or an ornate letter. Once preprocessing is complete, Aruspix reads the musical content of each page and converts it into an editable digital music format. Early music has been a challenge for traditional approaches to OMR [11], and so Aruspix has taken a unique approach based on hidden Markov models (HMMs) [12, 13]. The two key features of the method are that it does not require staff removal and that pitch detection is integrated within the recognition process. Aruspix also provides a built-in music editor to assist in correcting recognition mistakes. Features like this one make Aruspix usable not only as a research tool but also as an end-user application.

3 EXPERIMENTS

To perform our comparison, we evaluated both systems on the same set of pages to measure their accuracy. In particular, we were interested in comparing individual symbol recognition rates in addition to the overall performance of both applications. For these experiments, we used a data set built from four books of madrigals for four voices, composed by Luca Marenzio (1533–99). The books were printed in Venice and Antwerp between 1587 and 1607 (RISM [15] M-0579, M-0580, M-0583 and M-0585). As they are all re-editions of the same collection of madrigals, they contain the same musical content, with very few exceptions. The uniformity of content ensures that performance variations across books within our experiments are mainly related to the graphical appearance of the music fonts and document degradation, rather than the musical content itself.

From each book, we selected 40 pages for the training sets and 30 pages for the testing sets, resulting in total of 280 pages. For the training sets, we chose the first 20 pages of both the *Canto* and the *Alto* parts. This was not an arbitrary decision, as it is in this order that the pages would logically appear if either Gamut or Aruspix were used in a real-life project, as one would most likely start from the beginning of the books and continue onward. For this reason, and as the intent of the experiment was to reflect a comparison of tools rather than their absolute performance, we opted not to cross-validate our results. Further, it has been shown that cross-validated results do not vary significantly in such experiments [14].

(a) RISM M-0579 (R. Amadino, Venice, 1587)

(b) RISM M-0580 (G. Vincenti, Venice, 1587)

(c) RISM M-0583 (A. Gardano, Venice, 1603)

(d) RISM M-0585 (P. Phalèse, Antwerp, 1607)

Figure 1: Prints used for the experiments

Within Aruspix, each image underwent preprocessing with skew-removal and binarization, and all borders and non-musical symbols (e.g., titles, lyrics, and ornate letters) were removed. Because some image features used for classification are size dependent, the image size was then normalized to a staff height of 100 pixels. Ground-truth data were generated by transcribing the entire dataset using the Aruspix integrated editor.

3.1 Preprocessing in Gamut: staff removal

Staff-line removal is known to be very difficult within early documents due to numerous printing irregularities and frequent document degradations [9]. Dalitz et al. have recently presented an extensive comparison of several staff removal algorithms using self-generated degraded images and three different error metrics to gauge their performance [5]. This study clearly demonstrates that of the six algorithms tested, no single algorithm outperforms the others across all the data. As this evaluation does not focus on early music documents in particular, it was not possible to assume which algorithm would be most suitable for our printed sources.

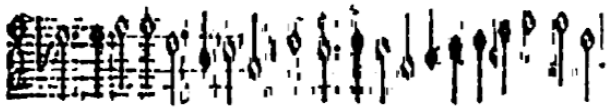

Figure 2: Typical staff removal error in M-0583

Moreover, as the study was based on artificial data, we could not be certain that a best candidate would be ideal for *real* data. For this reason, a preliminary experiment was undertaken to determine the most suitable algorithm for our purposes within the MusicStaves toolkit. A test database was created comprised of 70 hand-selected images chosen to provide a reasonable sampling of possible pages that one would encounter within this historical period.

For our purposes, it was easy to determine the three best algorithms amongst those tested (*simple*, *carter*, *roach-tatem*, *linetracking*, *fujinaga*, and *skeleton*) through a subjective evaluation of the results, as some of them failed dramatically. We have subdivided the errors accumulated over the data into minor and major errors. Minor errors are determined to be local to an individual or pair of symbols; effects to symbols beyond those affected were negligible. Alternatively, major errors encompass more destructive errors, usually including several symbols [1] . *Carter* was invariably unable to detect most staves, while *simple* and *roach-tatem* proved too destructive to symbols. These methods were significantly outperformed by the *linetracking*, *fujinaga*, and *skeleton* methods. In most images, these three algorithms were capable of the identification and removal of a majority of staves, including cases in which the others algorithms encountered major problems. We found the *linetracking* method to be most effective while avoiding major symbol deterioration, yet on occasion, noticeable problems were still encountered (figure 2).

3.2 Training and optimization

With all the data in hand, we were able to train HMMs for Aruspix and kNN classifiers for Gamut towards a comparison of the two systems. To measure the learning rates of both systems, we trained different HMMs and kNN classifiers using different numbers of pages of our training sets, from 1 to 40. Each model or classifier was then evaluated against the same testing set, such that we were able to determine the change in both systems once trained on 1 to 40 pages. We chose not to mix pages from the different books as we wanted the models and classifiers to be highly book specific in order to evaluate the graphical characteristics that make either system perform better or worse.

[1] A comprehensive discussion of the results of this evaluation, as well as supplementary images are available online at `coltrane.music.mcgill.ca/DDMAL/`

Both Gamut and Aruspix core recognition systems can be optimized with specific approaches that have been evaluated as well. The performance of the Gamut classifier can be improved by adjusting and optimizing the features used to represent symbols. Selecting the best combination of features is not a trivial task, and it has been demonstrated that a high number of features does not necessarily increase the recognition rate [6]. For this reason, we used the same limited set of features as in [6], i.e., *aspect ratio*, *moments*, *nrows*, and *volume64regions*. The genetic algorithm (GA) for feature weight optimization in Gamut was then tested in our experiments. As optimizing a classifier is a fairly computationally expensive task (approximately 24 hours for a minimal optimization of one 40-page classifier), we were not able to test this approach on all generated classifiers.

Similarly, Aruspix HMMs may be improved with the incorporation of *n-gram*-based music models. These models are built by performing statistical analysis on training data, and used during decoding to evaluate the likelihood of symbol classification according to $n-1$ previously recognized symbols. In our experiments, we chose $n = 3$, which conveys that the likelihood of symbols to be recognized is evaluated according to the two preceding symbols. The music models were generated from the same data as from training HMMs (using 1 to 40 pages for each book).

3.3 Post-processing in Gamut: pitch detection

In Gamut, a post-processing step is required to retrieve pitch points from the recognized symbols. This task can be done heuristically, for example, by localizing the center of a note head according to the staff line positions. With well-printed and non-degraded documents, this is usually a straightforward task. Early music sources, however, introduce many printing irregularities and document degradations, which often result in minor failures during staff removal (i.e., small staff elements remaining), making this post-processing step much more challenging. Thus, as the solution to the problem is highly dependent on the type of music documents being processed, we had to design an entire set of specialized heuristic methods to retrieve the pitch points. The heuristic methods are based on basic image analysis techniques such as projection analysis and centroid detection. This was done using the plugins and toolkit facilities in Gamut which enable custom functionalities to be implemented easily and added to the framework.

3.4 Evaluation procedure

The recognition results were evaluated by calculating both recall and precision on the best-aligned sequences of recognized symbols. Other existing evaluation methods such as Bellini et al. [2] were reviewed, but they were not found to be relevant to the present study at it was primarily de-

signed for common music notation. Our first evaluation was to measure the performance of the core recognition engines by determining symbol recognition rates for both the Gamut kNN and Aruspix HMM methods, without pitch detection. This was a trivial alteration within Gamut, as we only needed to consider the output of the classifier, prior to pitch point detection. For Aruspix, however, it was necessary to short-circuit the system, as symbol and pitch recognition is normally performed concurrently. The HMMs were modified in such a way that pitches were ignored, reducing the number of different symbols to be recognized from more than 130 to less than 30, hence enabling a direct comparison with Gamut classifiers. We also evaluated and compared improvement rates of both systems through incorporation of their specific optimization schemes, i.e., the genetic algorithm for Gamut and *n-gram*-based music models for Aruspix. Finally, we assessed the complete recognition process including pitch detection, yielding a full comparison of both OMR approaches for early music prints.

4 RESULTS

4.1 Evaluation of the recognition systems

Training the recognition systems of Gamut and Aruspix neither yields similar results nor generates the same learning curves. After 40 pages of training, Aruspix HMMs outperform the Gamut kNN classifiers for all four books, as shown in table 1. Only within the fourth book (M-0585) does the kNN classifier produce similar results to those of the HMMs, with more than 93% recall and more than 94% precision. Overall, the HMMs appears to be more robust and reliable, as the variability across the results is far smaller than within the kNN classifiers. While learning rates for Aruspix HMMs are fairly consistent across the four books, the Gamut kNN classifiers varies significantly from one book to another.

Figure 3 shows the learning curves for Aruspix HMMs and Gamut kNN classifiers for the book M-0579 and for the book M-0585. When the Gamut kNN classifiers perform well, they learn quicker than the HMMs, as demonstrated within the fourth book M-0585 (figure 3, bottom). In all cases, the kNN classifiers reach a plateau after 10 to 20 pages, whereas the HMMs continue to improve gradually even after 30 pages.

Book	Recall		Precision	
	kNN	HMMs	kNN	HMMs
M-0579	86.90	95.99	86.63	95.63
M-0580	86.76	93.51	89.36	97.32
M-0583	77.30	87.58	81.04	93.95
M-0585	93.35	94.72	93.47	96.80

Table 1: Recall and precision for the core recognition systems after 40 pages

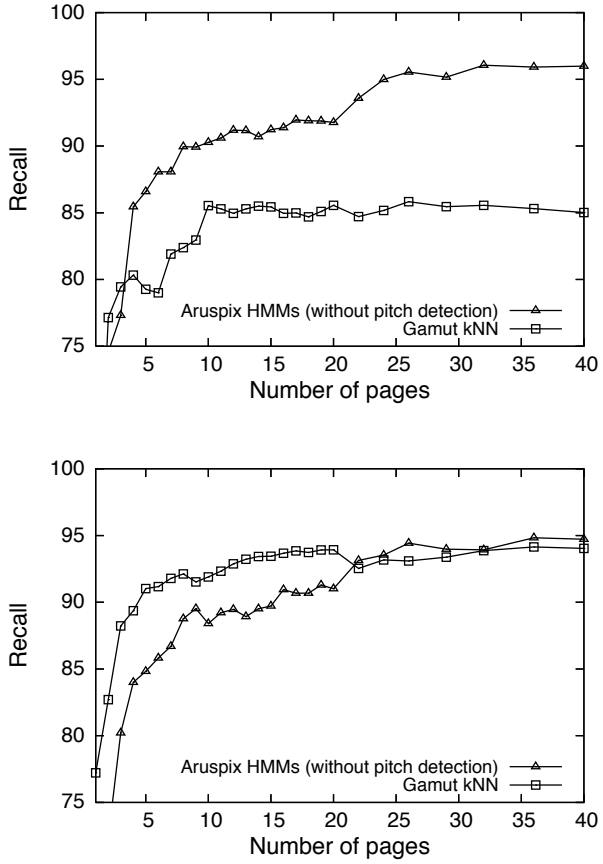

Figure 3: Learning curves (recall) of the core recognition systems on M-0579 and M-0585

For both Gamut kNN classifiers and Aruspix HMMs, the third book (M-0583) appears to be the most difficult. It is the most degraded book in our set, with strong bleed-through showing printed characters of the verso through the page. This book was printed with a very similar music font to that in book M-0585 (see figure 1), upon which both systems performed quite well: the best results for Gamut were found within this book. We may therefore conclude that document degradation, and not font shape, is the main cause of difficulty for both systems encountered with this book. Such degradations make the binarization step, which is critical in such applications, much more challenging [3]. Figure 4 clearly illustrates the effects of binarization errors on subsequent staff removal and recognition.

4.2 Evaluation of the optimization techniques

GA optimization of Gamut kNN classifiers improved the results for each of the four books in our experiments. Further, it appeared to be extremely influential when the number of

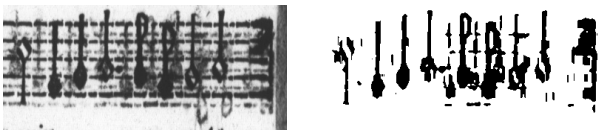

Figure 4: Original greyscale image and the same image after binarization and staff removal in one page of M-0583

pages was very small. Even within kNN classifiers already demonstrating steep (fast) learning curves, the GA optimization led to markedly faster learning, as shown in figure 5. Yet, in most cases, results were not significantly improved when using 20 pages or more.

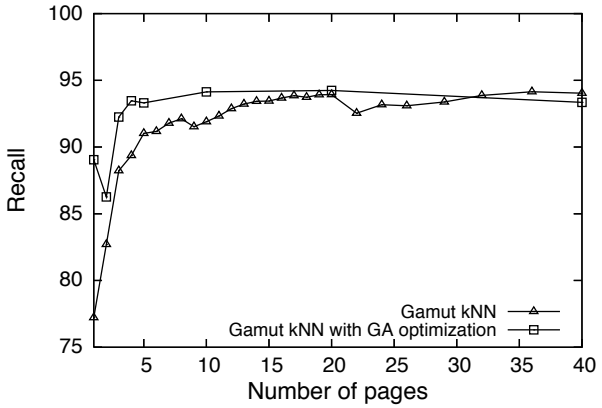

Figure 5: Improvement (recall) with GA optimization in Gamut on M-0585

For Aruspix, the integration of the *n-gram*-based models also improved results for all books, but in a notably different way than GA optimization within Gamut. First, the *n-gram*-based models improve precision rates much more so than recall rates. This verifies the logical intuition that the introduction of the model will reduce the likelihood of non-expected symbols during decoding, and thus reduce the number of insertions. Secondly, the improvement curve is also quite different than that of GA optimization within Gamut, as *n-gram* integration requires more pages to be significant, but displays a longer stable growth period, as shown in figure 6.

4.3 Evaluation of the OMR approaches

Table 2 presents the overall OMR recall and precision rates of both applications after 40 pages. In Gamut, pitches were determined heuristically in a post-processing step, whereas Aruspix uses an all-inclusive approach directly at the HMM level. On average, the differences for the complete OMR process from one book to the next are very similar to the those observed for the core recognition systems (see table 1), with a more significant loss in Gamut than in Aruspix.

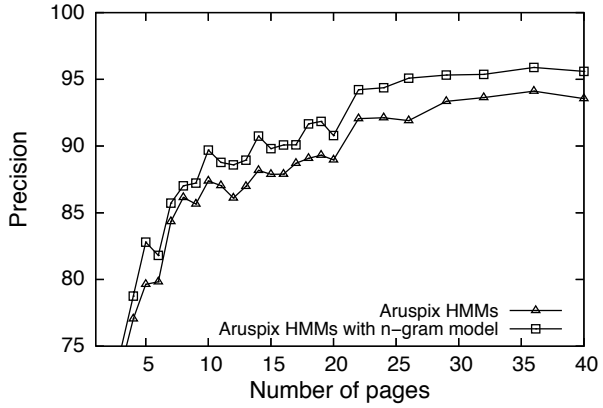

Figure 6: Improvement (precision) with the *n-gram*-based music models in Aruspix on M-0579

Book	Recall		Precision	
	Gamut	Aruspix	Gamut	Aruspix
M-0579	79.34	92.44	84.13	95.60
M-0580	76.18	92.52	84.55	96.44
M-0583	68.13	86.75	76.54	92.76
M-0585	81.61	92.77	88.34	96.23

Table 2: Recall and precision for the complete OMR results after 40 pages

These results distinctly show that retrieval of pitch points after symbol recognition is far from trivial with early music documents. In this task, the flexibility of Gamut could have been leveraged more, as the heuristic approach may be adjusted for a particular dataset.

In our experiment, Gamut failed to perform as well as Aruspix, despite the fact that the all-inclusive approach of Aruspix offer very few entry points for adjustment. Within the fourth book (M-0585), for which the core recognition systems performance is nearly equivalent, the final OMR output decreases significantly with the addition of pitch detection in Gamut (-11.71% for the recall rate) whereas the decline is only minimal in Aruspix (-1.95%). This slight reduction at the incorporation of pitch detection in Aruspix is an intriguing aspect of these results, as the all-inclusive approach of Aruspix has drastically inflated the number of symbols to be recognized (from less than 30 to more than 130 with our datasets) without affecting the recognition rates significantly.

5 CONCLUSIONS AND FUTURE WORK

The results of these experiments confirm the need for adaptability within tools for optical early music recognition. Even if the systems learn and perform differently, they both under-

performed on the same book (M-0585), which was by far the most degraded in our experimental set. This clearly indicates that binarization of degraded music documents is still an area in which improvements must be made. The difference between the results of M-0582 and M-0585, two books with the same content and quasi-identical music font, illustrates well the extent to which degradation may affect the performance of the systems. This is a quite probable explanation as to why our results obtained with Gamut are significantly lower than those obtained by Dalitz et al. [6] from a lute tablature application also built within the Gamut framework, which was evaluated with retouched facsimile images.

Our experiments with Gamut also demonstrate that staff removal within early music sources cannot be considered a solved problem. None of the algorithms tested worked extraordinarily well, and even minor failures, such as remaining small staff elements, certainly affect performance throughout the entire OMR process. Retrieving the pitch from the recognized symbols appears to be a challenging step within such documents, and the heuristic method currently used in Gamut could certainly be improved further.

The flexibility and the extensibility of the Gamut infrastructure proved its utility, as the system can easily be modified to improve its performance for a particular set of sources. Additionally, the learning speed of Gamut is an asset to the approach. The kNN classifiers learn quite rapidly, enabling a specific application to be built upon only a handful of pages. With the addition of the GA optimization, the amount of data required to achieve the best performance is reduced even further. Aruspix distinguishes itself by its performance and robustness. Together, HMMs and integrated pitch detection are an excellent approach with which to handle noise introduced by printing irregularities and degradations in early printed sources. Aruspix also provides a built-in editor specifically designed to assist in the correction of recognition errors. To enhance performance, Aruspix implements dynamic adaptation, which enables the amount of training data to be significantly reduced through optimization of previously trained HMMs.

Although both applications present distinct advantages, Aruspix clearly benefits from having been optimized for early music prints and may be used directly out-of-the-box, while the adaptability of Gamut may be advantageous for projects with a wider scope.

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THE *PERLHUMDRUM* AND *PERLLILYPOND* TOOLKITS FOR SYMBOLIC MUSIC INFORMATION RETRIEVAL

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ABSTRACT

PerlHumdrum is an alternative toolkit for working with large numbers of Humdrum scores. While based on the original Humdrum toolkit, it is a completely new, self-contained implementation that can serve as a replacement, and may be a better choice for some computing systems. *PerlHumdrum* is fully object-oriented, is designed to easily facilitate analysis and processing of multiple humdrum files, and to answer common musicological questions across entire sets, collections of music, or even the entire output of single or multiple composers. Several extended capabilities that are not available in the original toolkit are also provided, such as translation of MIDI scores to Humdrum, provisions for constructing graphs, a graphical user interface for non-programmers, and the ability to generate complete scores or partial musical examples as standard musical notation using *PerlLilypond*. These tools are intended primarily for use by music theorists, computational musicologists, and Music Information Retrieval (MIR) researchers.

1 INTRODUCTION

Humdrum [5, 3, 4] is a computer-based system pioneered by David Huron for manipulating and querying symbolic representations of music. Unlike notation-based systems such as GUIDO or MusicXML, Humdrum is primarily intended as a set of analytical tools to aid music theorists, musicologists, acousticians, cognitive scientists, and MIR researchers, among others. Also, Humdrum data files are differentiated from stored-performance formats such as MIDI, in that almost all have been encoded by hand from musical scores; more than 40,000 Humdrum files have been encoded to date, the majority of which are freely available online [1, 6].

The original motivation for this research was the desire to use the Humdrum resources in a series of music analysis projects, coupled with a certain frustration with the lack of functionality of some of the original tools which were reliant on older Unix environments and were difficult to make work under current versions of Linux. Other Humdrum tools, such as those for entering MIDI information, are associated with hardware that currently unavailable in most operating systems. Also, many of the problems that the project explored required writing additional programs that worked directly with the Humdrum data, such as data collection and extraction across large numbers of scores, and it was simply easier to write them fresh in Perl than to try to accomplish the same thing based on one of the older Awk tools. Over time a collection of alternate tools, extensions, and replacements began to take shape, and at some point it simply made sense to collect everything under a single code base that shared methods for common tasks in a object oriented framework. In the process, it became possible to rethink some aspects of the original Humdrum Toolkit, and introduce some new possibilities that are useful when working on these sorts of problems.

Additionally, from working with various musicologists, it became clear that a system was needed for automatically producing musical excerpts in common musical notation, as a method for displaying search results. This led to the creation of the *PerlLilypond* programs, that provide a scriptable environment for generating notation. The two systems are designed to complement one another.

The author refers to the two systems as *PerlHumdrum* and *PerlLilypond*. This is primarily to differentiate their names the original programs. However, within the Perl environment, the module names `Humdrum` and `Lilypond` are used. This is partially because of traditional naming conventions in Perl, but also because the author doesn't see the necessity of repeatedly inflicting an extra syllable on developers and users during tasks such as multiple object creation.

2 OVERVIEW OF *HUMDRUM*

The overall *Humdrum* system is comprised of two main parts: a set of file-based storage formats for time-aligned symbolic data, and the accompanying collection of UNIX-based tools for manipulating and querying this data. The *PerlHumdrum* system discussed in this paper is designed to simplify operations on the former, while providing an alternative to the latter.

Humdrum data files consist of symbols representing musical or other symbolic information, arranged in time-aligned columns known as *spines*. As an example, the first measures

of a Bach chorale and the representative Humdrum excerpt are shown in Figure 1 and Table 1 respectively.

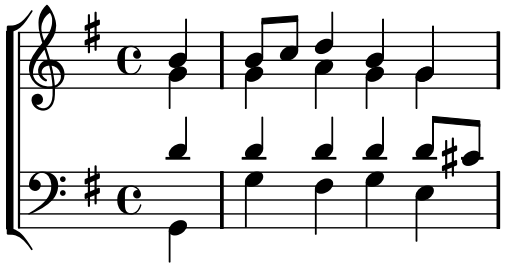

Figure 1. *Uns ist ein Kindlein heut' gebor'n*, J. S. Bach

**kern	**kern	**kern	**kern
*bass	*tenor	*alto	*soprn
*k[f#]	*k[f#]	*k[f#]	*k[f#]
4GG	4d	4g	4b#
=1	=1	=1	=1
4G	4d	4g	8b#
.	.	.	8cc
4F#	4d	4a	4dd
4G	4d	4g	4b
4E	8d	4g	4g
.	8c#	.	.

Table 1. A Humdrum `Kern` encoding of Figure 1

Humdrum files consist only of ASCII characters, including tabs and spaces, and are easy to create, view and edit in any standard text editor. Each vertical spine represents a single stave of music from the original score. Where new events in the original score occur horizontally, moving from left to right. In the Humdrum version, new events are added to the bottom of each spine. In many ways, Humdrum encodings can be seen as normal scores rotated clockwise by 90 degrees. Items aligned horizontally occur at the same time, and a single horizontal line of data is known as a *record*. Items marked with asterisks (*,**) are metadata events or *interpretations* that give instructions as to how data spines are to behave. For instance, spines can be added, subtracted, or exchanged, among other operations. Interpretations are also used to indicate key and time signature.

The first item in each column indicates the type of Humdrum format in use. In this case the standard `Kern` format that is more or less analogous to common-practice music notation is being employed, with notes encoded as a combination of numeric value and letter, representing the rhythmic and pitch values. Measure numbers are preceded by equal signs and are repeated in all columns. The letter c indicates middle *c* (261.63 Hz), with lower octaves being represented by upper-case letters, higher octaves by lower-case letters, and larger distances from middle *c* by accumulated lettering. A set of *c* notes going from lower to higher octaves would be encoded as "CCC, CC, C, c, cc, ccc".

While the `Kern` format is the most common type of Humdrum encoding, many other types of encodings exist, including frequency, scale degrees, melodic and harmonic intervals, solfege symbols, set-theory representations, and even psychoacoustical representations such as barks or critical band rates. Spines may be added or removed from a file as necessary, or manipulated in various other ways. Multiple encoding schemes can also be used in a single file, and the user can define their own types to accommodate specific needs.

Coupled with data storage part of Humdrum is the toolkit, a set of programs for working with *Humdrum* data files. The programs encompass a rich set of functionality, including devices for manipulating and querying, and summarizing this kind of data. It is difficulties with these tools in particular that *PerlHumdrum* is designed to address.

3 *PERLHUMDRUM* USAGE

The original Humdrum toolkit operates using the standard UNIX command line paradigm. Commands in Humdrum generally consist of Awk scripts that take command line options and an input Humdrum file, and produce another Humdrum file. Multiple commands can be strung together using pipes, and intermediate results can be stored as temporary files to undergo further processing. For instance, one of the simplest of all Humdrum programs is the `census` command, used here to obtain a list of overall statistics for a kern file called *bach.krn*:

```
census -k bach.krn
```

This produces a text file containing various statistics about the original piece, such as the number of notes, number of comments, and other basic information.

In *PerlHumdrum*, the basic ingredient is the `Humdrum` object. The equivalent program to the above using *PerlHumdrum* is shown below.

```
use Humdrum;

my $c=Humdrum->new('bach.krn');
my $dat=$c->census(k=>1);
```

Command line options from the original programs are available as parameters to the *PerlHumdrum* methods. Great care has been taken to ensure the best possible compatibility between the original toolkit and this one; most of the original programs have been converted to methods, and great

care has been taken to ensure that all options have been accounted for and behave identically.

Applying this command to multiple files, to get a sum of notes for instance, is almost as simple:

```
use Humdrum;

my $sum=0;

foreach my $file(@filelist){
    my $c=Humdrum->new($file);
    my $dat=$c->census(k=>1);
    $sum+=$dat->get_datatokens();
```

In the original toolkit, processes could be chained together by saving the results of each stage as a temporary file and using it as input to the next, or by using the pipe symbol. The example below demonstrates a short processing chain that removes the non-note elements from the `kern` file "bach.krn" and then performs a similar census operation.

```
kern -x  bach.krn | census -k
```

In *PerlHumdrum*, under normal circumstances, a `Humdrum` object is considered immutable; that is, the output of one operation produces another `Humdrum` object.

The previous set of commands can be translated into a *PerlHumdrum* program as follows:

```
use Humdrum;

my $c=Humdrum->new('bach.krn');
my $k=$c->kern(x=>1);
my $dat=$k->census(k=>1);
```

Here is an even more succinct version:

```
use Humdrum;

my $c=Humdrum->new('bach.krn');
my $dat=$c->kern(x=>1)->census(k=>1);
```

Obviously, much more complex examples are possible.

4 ADVANTAGES

The *PerlHumdrum* toolkit has several advantages over the original Humdrum toolkit.

The primary advantage is that the object oriented paradigm makes it simple to apply complex data analysis and modification operations to multiple files, through the use of loops or other control structures. While this is possible in the original command-line syntax using command-line bash shell loops and judicious use of piping, the results are often brittle and not well-suited to large-scale analyses of symbolic data as is often required in MIR research. We believe that these limitations of the current toolkit have limited the uses of existing `Kern` repositories in many MIR research areas.

A second, perhaps less-obvious advantage is that the original Humdrum toolkit makes complete sequential passes through entire Humdrum files and then passes that data on to further commands through pipes. While many operations are possible in this way, others are extremely difficult. Consider the situation of trying to analyze all of the cadences in a piece or set of pieces. Cadences are most easily-identified by working backwards through a file, finding the resolution in reverse (I-V) and then working back until the "start" of the cadence is detected. Doing such work using only forward sequential passes, perhaps with an additional operation once the cadence is located, is extremely difficult. *PerlHumdrum* provides several different methods of proceeding through a Humdrum file, including reverse iteration and ranged iteration. The array of objects is also available to the user, making it possible to define any kind of non-sequential operation the user might like (only even tokens, for instance).

Other advantages are:

- consistent use of OOP principles, instead of "imperative" Awk-style scripts simplifies the addition of new commands.
- The use of perl provides a common base for other "data-munging" operations that Perl is commonly used for. This makes it easy to connect Humdrum operations to databases, linear algebra libraries, or even CGI scripts for web pages, to give just a few examples.
- Unlike the Awk and C basis of the original toolkit, *PerlHumdrum* runs natively on many different computing platforms, including Unix, Macintosh, and Windows systems without recompiling.

5 *PERLLILYPOND*

A parallel package, *PerlLilypond*, has been developed that can take information from a Humdrum object and convert it into common practice music notation. As the name suggests, *PerlLilypond* uses *LilyPond* [14] as the underlying notation engine.

PerlLilypond has two primary virtues. First, *PerlLilypond* can be easily scripted and called within other programs. While *PerlLilypond* can be used to generate entire scores, and can function as an alternative, non-proprietary notation system. The intended use is for displaying multiple excerpts from a collection of Humdrum scores, such as all of the cadences across a corpus, without individual human intervention and editing. This is extremely difficult to do with any of the more-familiar commercial notation programs.

Secondly, and perhaps more importantly, *PerlLilypond* displays the results of symbolic searches in a form that is accessible to most musicians. Experience has shown that many traditional musicologists and theorists without a background in MIR have difficulties with non-traditional notation formats. Displaying results in a traditional notation format makes it much easier for many music researchers to use this technology, and we feel that the lack of such a system has probably been the single biggest impedement to the adoption of these technologies within that community. Also, MIR researchers themselves, with traditional music training, may find it much easier and more succinct to display results in this format.

There are two ways to use *PerlLilypond*. The easiest is simply to pass it a *PerlHumdrum* object, which is the common output of most *Humdrum* operations, and *PerlLilypond* will do all the work of converting the object into its own internal format. Thus, the entire process of printing a Humdrum score or any Humdrum data can be reduced to the procedure outlined below.

```
use Humdrum;
use LilyPond;

my $humdrumexcerpt=Humdrum->new('Bach.krn');
my $l=Lilypond->new($humdrumexcerpt);
$l->print();
```

A second way is to gradually build up an object by adding notes, staves, and other musical structures sequentially. In this mode, a *PerlLilypond* object can be considered a kind of long pipe that we keep pushing objects into until our excerpt is finished. Other "container" objects, such as chords or tuplets, can be created as needed and added to a stave. No facility is provided for editing materials inside a *PerlLilypond* object once they have been allocated, although references to the underlying raw arrays is available for end users who wish to build their own subroutines. This is because such editing tools are already in *PerlHumdrum*.

The procedure for creating a short excerpt consisting of a note, a chord, and a final note is shown below:

```
my $ll=Lilypond->new();
$ll->addstave('soprano');
$ll->addnote('soprano',{step=>'d'});

my $lh=Lilypond::Chord->new;
$lh->add(Lilypond::Note->new(
    {duration=>8}));
$lh->add(Lilypond::Note->new(
    {step=>'e',duration=>8}));
$lh->add(Lilypond::Note->new(
    {step=>'g',duration=>8}));
$ll->addchord('soprano',$lh);
$ll->addnote('soprano',
    {step=>'g',duration=>4,dots=>1});
$ll->makeexcerpt();
```

First, a new `Lilypond` object is created and a stave named *soprano* is added. Following the addition of the first note, a `Lilypond::Chord` object is created and three notes are "poured" into it in sequence. The resulting object is then added to the stave, followed by a final note. Finally, the `makexcerpt` method is called, calling the *Lilypond* binary, and generating the excerpt in all output formats.

Once a user is finished constructing a *PerlLilypond* object, various commands are available to actually get output as PNG or postscript output, transparently created by calling the actual *LilyPond* binary.

Facilities are also provided for displaying large numbers of excerpts as PNG graphics on an HTML page. HTML was chosen as the primary output format for excerpts because web pages are essentially unlimited in length, can easily align images, and can be easily viewed in practically any computing environment. An example of some output generated directly from a Humdrum file is shown in Figure 2. Here, a cadence has been automatically detected and marked with an 'X'.

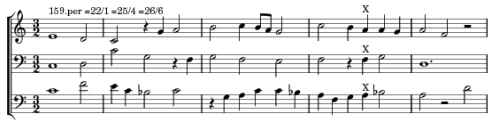

Figure 2. A simple example of web page output

One especially useful facility of *PerlLilypond* is the ability to introduce "hidden" elements that are not seen, but instead act as spacers between other notes. This is extremely useful for displaying the results of aligned sequences of notes, such as those that might be produced by musical applications of various alignment algorithms [17, 9]. At present, *PerlLilypond* captures about 60 percent of the capabilities of *LilyPond*, including almost everything connected to common notation. Development is ongoing to add the remaining alternate notation styles such as percussion notation.

6 EXTENSIONS

PerlHumdrum also provides a number of additional capabilities that are not present in the regular Humdrum toolkit.

6.1 Support for Analyses of Multiple Scores

With the exception of the `patt` and `grep` tools, Humdrum is cumbersome for the analysis of large scores. Such procedures, in the original toolkit, often involves some tricky

shell scripting, writing the results to a log file, and then providing an additional program to process these separately. This is especially difficult if the cumulative results need to be recorded, or if the results of each analysis needed to be inserted into a database (for instance). *PerlHumdrum* provides several "meta-tools" that make it simple to apply a referenced procedure or set of procedures to a collection of files. Figure 3 shows the final results of one such operation, in which the number of common rule violations in the original Bach Chorales have been accumulated as percentages of files of the original collection ([2, 10]).

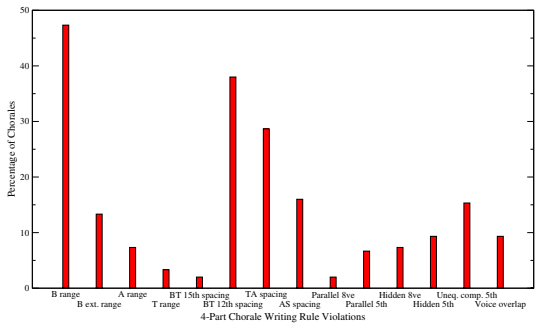

Figure 3. Common rule violations in Bach Chorales

6.2 Graphing Capabilities

In a similar vein as the above, a set of procedures has been provided to facilitate easy graphing of extracted results. The most common variety of this is the "scatterplot" diagram, such as that shown in Figure 4. This provides an easy way to visualize aspects of entire collections, and such graphs are of great interest to musicologists ([7, 8, 18]).

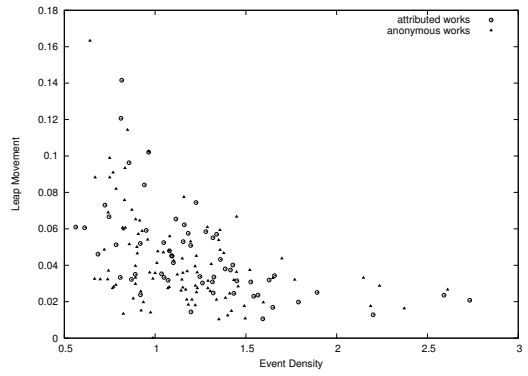

Figure 4. Comparing an entire collection

6.3 MIDI File Conversion

The original toolkit has the ability to convert Humdrum files to MIDI format. There is also a limited facility for converting direct MIDI input from a MIDI-in port into Humdrum, but this relies on a specific hardware library and only functions under Microsoft DOS. *PerlHumdrum* provides a `MakeMidi` method that will convert a MIDI file into Humdrum notation. `MakeMidi` is not currently very good at making assumptions, and relies heavily on the MIDI files being well-behaved To get good results, MIDI files must include tempo and key signature, isolate each stave to a different track, and should probably be heavily quantized. It currently does not handle more complicated tuplets such as quintuplets. However, the current capabilities work well enough for most common music notation scores, and additional improvements are being implemented.

6.4 Integration with Other MIR Systems

While not technically an aspect of *PerlHumdrum*, one very unique aspect of Perl is the *Inline* facility, which allows programmers to actually include foreign code directly inside a Perl source code file. This makes it possible to mix C, Java, Octave, and many other languages directly into Perl. The MIR field currently has many different disparate toolkits designed to handle specific problems, but very few complete systems. Nor do such systems seem likely in the future, as writing programs in multiple languages is usually extremely complex. Using Inline and Perl, it is possible to construct larger programs combined of multiple languages and source code files, including *PerlHumdrum*. We have used this facility in the past, and found it extremely effective, yet it still remains unknown in the MIR community as a whole.

7 FUTURE WORK

Most of the functionality of the original toolkit has been implemented in *PerlHumdrum*. However, a few of the more obscure options of some of the original toolkit programs are currently unsupported, have undergone changes in function in this new context, or have unfortunately been recently demonstrated as somewhat less than robust. Regression testing has helped considerably with these situations, and the entire toolkit is approaching something close to a real, distributable replacement for the original.

It is intended that *PerlHumdrum* be made freely available as is the original Humdrum toolkit, probably under the Gnu Public License [13]. The standard method for making code available to the Perl community is by distributing the modules through the CPAN network [15]. This makes it extremely easy for Perl users to download and install the entire set of modules from within Perl itself. This conversion has yet to be completed; however, current versions of *Perl-*

Humdrum are available through direct email contact with the author.

Several other extensions to *PerlHumdrum* are currently planned or in progress, including: conversion classes to other notation formats such as GUIDO, connections to databases such as PostGreSQL [16] for large-scale storage of music fragments, better integration with the Perl MIDI classes, and the ability to synthesize and play fragments directly using Timidity.

The *PerlLilypond* notational facility is still under development and has difficulty with some advanced notational situations. It has not been well-tested on many types of contemporary music, primarily because there are fewer examples of this style of music in the various *Humdrum* repositories.

The object-oriented nature of *PerlHumdrum* makes it natural to consider building a graphical interface for Humdrum. This has been attempted before [12, 11, 19], but the results have never been satisfactory. Development of such a resource would also be extremely helpful in aiding the adoption of these tools by musicologists and music theorists.

Finally, the set of operations available in Humdrum are probably the most comprehensive set of symbolic music operations in the MIR world. We would like to adapt these tools to operate on other possible file formats. The best candidate at this time is the MusicXML format. In fact a rudimentary version of this already exists, and will be completed as time and necessity permit.

8 ACKNOWLEDGMENTS

The author wishes to express his sincere thanks to Frauke Jürgensen for her ongoing and enthusiastic support of this project; her advice and patience have been invaluable. Additionally, I would like to acknowledge the encouragement of both Craig Stuart Sapp and Don Byrd in these efforts.

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COMBINING FEATURES EXTRACTED FROM AUDIO, SYMBOLIC AND CULTURAL SOURCES

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ABSTRACT

This paper experimentally investigates the classification utility of combining features extracted from separate audio, symbolic and cultural sources of musical information. This was done via a series of genre classification experiments performed using all seven possible combinations and subsets of the three corresponding types of features.

These experiments were performed using jMIR, a software suite designed for use both as a toolset for performing MIR research and as a platform for developing and sharing new algorithms.

The experimental results indicate that combining feature types can indeed substantively improve classification accuracy. Accuracies of 96.8% and 78.8% were attained respectively on 5 and 10-class genre taxonomies when all three feature types were combined, compared to average respective accuracies of 85.5% and 65.1% when features extracted from only one of the three sources of data were used. It was also found that combining feature types decreased the seriousness of those misclassifications that were made, on average, particularly when cultural features were included.

1. INTRODUCTION

Music is a multi-faceted area of inquiry, and there are many factors that influence any individual's experience of music. Given the variety of possible approaches to studying music, researchers can have a tendency to focus their attentions on limited types of music or on particular specialized approaches to studying music. One of the most exciting aspects of the contemporary music information retrieval (MIR) field is that it attempts to break down these barriers using a multi-disciplinary approach to music.

In the past, MIR research has tended to focus on studying either audio (e.g., MP3), symbolic (e.g., MIDI) or cultural (e.g., web data, user tags, surveys, etc.) sources of information about music. Traditionally, research using each of these sources has been relatively segregated based on whether a researcher had a corresponding background in signal processing, music theory or library sciences and data mining. In recent years, however, MIR researchers have increasingly begun to study these sources of information in combination, with a particular emphasis on research combining audio and cultural sources of data.

Features extracted from all three types of data can provide valuable information for use in music classification and similarity research. Audio is clearly useful because it is the essential way in which music is consumed, and cultural data external to musical content is well-known to have a large influence on our understanding of music [8].

Symbolic data has recently been receiving less attention from MIR researchers than it did in the past. The value of symbolic data should not be overlooked, however, as much of the information associated with high-level musical abstractions that can be relatively easily extracted from symbolic formats is currently poorly encapsulated by the types of features that are typically extracted from audio, which tend to focus primarily on timbral information

Symbolic formats can thus, at the very least, be a powerful representational tool in automatic music classification. This characteristic will become increasingly valuable as polyphonic audio to symbolic transcription algorithms continue to improve. Even though such technologies are still error-prone, it has been found that classification systems can be relatively robust to such errors [6].

This paper focuses on an experimental investigation of the extent to which combining features extracted from the three types of data can be advantageous in automatic music classification. If the orthogonal independence of the feature types is high, then performance boosts can potentially be attained in a variety of applications by combining features extracted from the different data types.

This investigation was performed via sets of automatic genre classification experiments. Genre classification in particular was chosen because it is a complex and difficult task that combines diverse musical variables. The experiments could just as easily have been performed using other types of classification, however, such as mood or artist classification. The essential issue being investigated remains the potential performance improvements attained by combining features extracted from the three types of data.

2. RELATED RESEARCH

There has been a significant amount of research on combining audio and cultural data. Whitman and Smaragdis

[14] performed particularly important early work on combining audio features with cultural data mined from the web, and achieved substantial performance gains when doing so. Dhanaraj and Logan [3] took a more content-based approach by combining information extracted from lyrics and audio. Others have combined audio and cultural data for the purpose of generating music browsing spaces (e.g., [5]). Aucouturier and Pachet [2] used a hybrid training approach based on acoustic information and boolean metadata tags. Research has also been done on using audio data to make correlations with cultural labels, which can in turn improve other kinds of classification (e.g., [12]).

There has been much less work on combining symbolic data with audio data. Lidy et al. [6] did, however, find that combining audio and symbolic data can result in improved performance compared to when only audio data is used.

To the best knowledge of the authors, no previous research has been performed on combining symbolic and cultural features or on combining all three feature types.

Far too many papers have been published on automatic genre classification in general to cite with any completeness here. One influential work that bears particular mention, however, is that of Tzanetakis and Cook [13].

3. THE JMIR SOFTWARE SUITE

jMIR is a suite of software tools developed for use in MIR research. It was used to perform the experiments described in this paper. jMIR includes the following components:

- **jAudio** [7]: An audio feature extractor that includes implementations of 26 core features. jAudio also includes implementations of "metafeatures" and "aggregators" that can be used to automatically generate many more features from these core features (e.g., standard deviation, derivative, etc.).
- **jSymbolic** [10]: A symbolic feature extractor for processing MIDI files. jSymbolic is packaged with 111 mostly original features [8].
- **jWebMiner** [11]: A cultural feature extractor that extracts features from the web based on search engine co-occurrence page counts. Many user options are available to improve results, including search synonyms, filter strings and site weightings.
- **ACE** [9]: A meta-learning classification system that can automatically experiment with a variety of different dimensionality reduction and machine learning algorithms in order to evaluate which are best suited to particular problems. ACE can also be used as a simple automatic classification system.

The jMIR components can be used either independently or as an integrated suite. Although the components can read and write to common file formats such as Weka ARFF, jMIR also uses its own ACE XML file formats that offer a number of significant advantages over alternative data mining formats [9].

The components of jMIR were designed with the following goals in mind:

- Provide a flexible set of tools that can easily be applied to a wide variety of MIR-oriented research tasks.
- Provide a platform that can be used to combine research on symbolic, audio and/or cultural data.
- Provide easy-to-use and accessible software with a minimal learning curve that can be used by researchers with little or no technological training.
- Provide a modular and extensible framework for iteratively developing and sharing new feature extraction and classification technologies.
- Provide software that encourages collaboration between different research centers by facilitating the sharing of research data using powerful and flexible file formats.

In order to improve accessibility, each of the jMIR components is, with the temporary exception of ACE, packaged with an easy-to-use GUI. The jMIR components also include user manuals and help systems.

The jMIR components are all implemented in Java, in order to make them as platform-independent as possible. They are open-source and distributed free of charge.[1]

4. THE SAC DATASET

The SAC (Symbolic, Audio and Cultural) dataset was assembled by the authors in order to provide matching symbolic, audio and cultural data for use in the experiments described in Section 5. SAC consists of 250 MIDI files and 250 matching MP3s, as well as accompanying metadata (e.g., title, artist, etc.). This metadata is stored in an iTunes XML file,[2] which can be parsed by jWebMiner in order to extract cultural features from the web.

It was decided to acquire the matching MIDI and audio recordings separately, rather than simply synthesizing the audio from the MIDI. Although this made acquiring the dataset significantly more difficult and time consuming, it was considered necessary in order to truly test the value of combining symbolic and audio data. This is because audio generated from MIDI by its nature does not include any additional information other then the very limited data encapsulated by the synthesis algorithms.

SAC is divided into 10 different genres, with 25 pieces of music per genre. These 10 genres consist of 5 pairs of similar genres, as shown in Figure 1. This arrangement makes it possible to perform 5-class genre classification experiments as well as 10-class experiments simply by combining each pair of related genres into one class. An additional advantage is that it becomes possible to meas-

[1] jmir.sourceforge.net

[2] Contact cory.mckay@mail.mcgill.ca for access to the file.

ure an indication of how serious misclassification errors are in 10-class experiments by examining how many misclassifications are in an instance's partner genre rather than one of the other 8 genres.

Blues: Modern Blues *and* Traditional Blues
Classical: Baroque *and* Romantic
Jazz: Bop *and* Swing
Rap: Hardcore Rap *and* Pop Rap
Rock: Alternative Rock *and* Metal

Figure 1: The ten genres found in the SAC dataset.

5. EXPERIMENTAL PROCEDURE

The first step of the experiment was to use jMIR's three feature extractors to acquire features from each matched audio recording, MIDI recording and cultural metadata. Details on the particular features extracted are available elsewhere ([7], [8], [10] and [11]).

To provide a clarifying example, features might be extracted from a Duke Ellington MP3 recording of *Perdido,* from an independently acquired MIDI encoding of the same piece, and from automated search engine queries using metadata such as artist and title. Three types of feature sets were therefore extracted for each piece, one corresponding to each of the three data types.

These three types of features were then grouped into all seven possible subset combinations. This was done once for each of the two genre taxonomies, for a total of fourteen sets of features (as shown in Table 1), in preparation for fourteen corresponding classification experiments designed to measure how well the different feature sets performed relative to one another.

Feature Type	5-Genre Code	10-Genre Code
Symbolic	S-5	S-10
Audio	A-5	A-10
Cultural	C-5	C-10
Symbolic + Audio	SA-5	SA-10
Audio + Cultural	AC-5	AC-10
Symbolic + Cultural	SC-5	SC-10
Symbolic + Audio + Cultural	SAC-5	SAC-10

Table 1: The identifying codes for the 14 experiments.

ACE was then trained on and used to classify each of the fourteen feature sets in fourteen independent 10-fold cross-validation experiments. This resulted in two classification accuracy rates for each of the seven feature type combinations, one for each of the two genre taxonomies. As a side note, ACE includes dimensionality reduction functionality, so training was actually performed with automatically chosen subsets of the available features.

It is desirable not only to determine how effective each of the feature type combinations are at achieving correct classifications, but also how serious those misclassifications that do arise are. Two classifiers with similar raw classification accuracy rates can in fact be of very different value if the misclassifications made by one classifier result in classes that are more similar to the "correct" class.

A normalized weighted classification accuracy rate was therefore calculated for each of the 10-genre experiments in order to provide insight on error types. This was calculated by weighting a misclassification within a genre pair (e.g., Alternative Rock instead of Metal) as 0.5 of an error, and by weighting a misclassification outside of a pair (e.g., Swing instead of Metal) as 1.5 of an error.

6. RESULTS AND DISCUSSION

6.1. Results

The average classification accuracy rates across cross-validation folds for each of the fourteen experiments outlined in Table 1 are shown in Table 2, including weighted and unweighted results. Figures 2 and 3 illustrate the unweighted results for the 5-genre and 10-genre experiments respectively. These results are summarized in Table 3 and Figure 4, which show the average results for all experiments using one feature type, all experiments using two feature types and all experiments using three feature types.

6.2. Effects of Combining Feature Types on Accuracy

As can be seen in Figures 2 and 3, all combinations of two or three feature types performed substantially better than all single feature types classified independently. Furthermore, combining all three feature types resulted in better performance than most pairs of feature types.

This is illustrated in Figure 4, which shows important average increases in performance when feature types are combined. Combining all three feature types resulted in increases in performance of 11.3% on the 5-genre taxonomy and 13.7% in the 10-genre taxonomy, compared to the average performances of each of the single feature types classified individually. Considered in terms of percentage reduction in error rate, this corresponds to impressive improvements of 78.0% and 39.3% for the 5 and 10-genre taxonomies, respectively.

A Wilcoxon signed-rank test indicates that, with a significance level of 0.125, the improvements in performance of two or three feature types over one type were statistically significant in all cases. However, the improvements when three feature types were used instead of two were not statistically significant. The corresponding average increases in performance were only 2.3% and 2.7% for the 5 and 10-genre taxonomies, respectively.

Overall, these results indicate that, at least to some extent, the three different types of features contain orthogonally independent information, and can therefore be profitably combined for a variety of purposes.

	S	A	C	SA	AC	SC	SAC
5-UW	86.4	82.8	87.2	92.4	95.2	94	96.8
10-UW	66.4	67.6	61.2	75.6	78.8	75.2	78.8
10-W	66.4	67.4	66.6	78.6	84.6	81.2	84.2

Table 2: The unweighted classification accuracy rates for the 5-genre (5-UW) experiments and both the unweighted (10-UW) and weighted (10-W) accuracy rates for the 10-genre experiments. Results are reported for each feature type combination, as described in Table 1. All values are average percentages calculated over cross-validation folds.

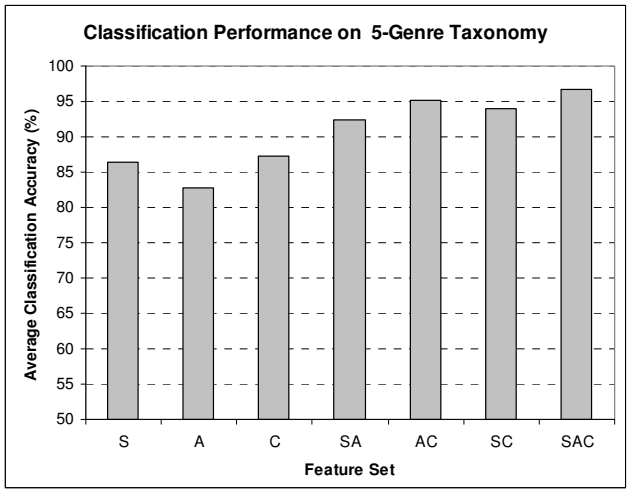

Figure 2: The classification accuracy rates for the 5-genre taxonomy, as described in Table 1.

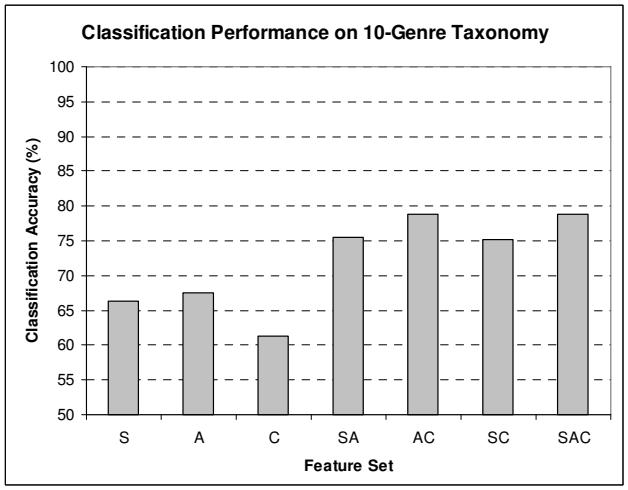

Figure 3: The unweighted classification accuracy rates for the 10-genre taxonomy, as described in Table 1.

	1 Feature Type	2 Feature Types	3 Feature Types
5-UW	85.5	93.9	96.8
10-UW	65.1	76.5	78.8
10-W	66.8	81.5	84.2

Table 3: The average classification accuracy rates for all experiments employing just one type of feature (S, A and C), two types of features (SA, AC and SC) or all three types of features (SAC). Results are specified for the 5-genre taxonomy (5-UW), the unweighted 10-genre taxonomy (10-UW) and the weighted 10-genre taxonomy (10-W). All values are percentages, and are calculated as simple mean averages from Table 2.

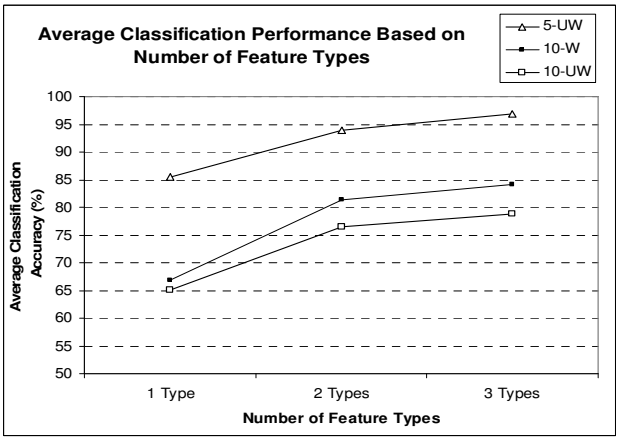

Figure 4: The average classification accuracy rates for all experiments employing just one type of feature (S, A and C), two types of features (SA, AC and SC) or all three types of features (SAC). The three trend lines refer to the 5-genre taxonomy (5-UW), the unweighted 10-genre taxonomy (10-UW) and the weighted 10-genre taxonomy (10-W). This data corresponds to the values in Table 3.

6.3. Types of Misclassifications

As described in Section 5, normalized weighted classification accuracy rates were calculated for the experiments on the 10-genre taxonomy in order to evaluate the seriousness of the particular misclassifications that were made. The results, and how they compare to the unweighted classification accuracies, are shown in Table 2 and Figure 5.

The weighted and unweighted accuracies were not significantly different when the audio and symbolic features were processed individually. However, the weighted performance was 3% higher than the unweighted performance when these two feature types were combined. Although this is not a dramatic increase, it is an indication that combining these feature types can make those misclassifications that do occur result in classes closer to the model classes in addition to increasing classification accuracy itself, as discussed in Section 6.2.

The differences between the weighted and unweighted classification accuracies were greater in all feature sets that included cultural features. These weighted rates were higher than the unweighted rates by an average of 5.7%, a difference that, based on Student's paired t-test, is statistically significant with a significance level of 0.005.

Overall, these results indicate that the types of misclassifications that occur when cultural features are used are less serious than when audio or symbolic features are used alone. Quite encouragingly, it also appears that this improvement in error quality carries through when cultural features are combined with audio and symbolic features.

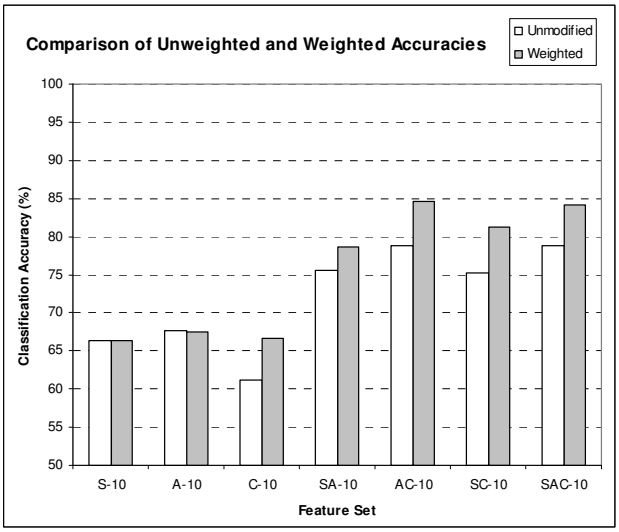

Figure 5: The differences between the unweighted and weighted classification accuracies on the 10-genre taxonomy for each of the seven feature type combinations. This data corresponds to the last two rows of Table 2.

6.4. General Genre Classification Performance

In order to put the experimental results described here in context, it is appropriate to compare them with classification accuracies achieved by high-performing existing specialized genre classification systems. It is important, however, to keep in mind the essential caveat that different classification systems can perform dramatically differently on different datasets, so direct comparisons of classification accuracies calculated on different datasets can give only a very rough indication of comparative performance.

The MIREX (Music Information Retrieval Evaluation eXchange) evaluations offer the best benchmarking reference points available. Although no evaluations of genre classification based on cultural data have been carried out yet at MIREX, both symbolic and audio genre classification evaluations have been held, most recently in 2005 and 2007, respectively. The highest accuracy for symbolic classification was 84.4%, attained on a 9-genre taxonomy by McKay and Fujinaga's Bodhidharma system [15]. The highest classification accuracy attained in audio classification was 68.3%, achieved by the University of Illinois's International Music Information Retrieval Systems Evaluation Laboratory (IMIRSEL) team, on a 10-genre taxonomy [16].

The experiments described in this paper achieved classification accuracies of 67.6% using only features extracted from audio and 66.4% using only features extracted from symbolic data. This is comparable to the best MIREX audio result of 68.3%, but significantly lower than the best MIREX symbolic result of 84.4%, which was achieved on a taxonomy only smaller by one class.

This latter result is intriguing, as jSymbolic uses the same features and feature implementations as Bodhidharma. The difference is likely due at least in part to the specialized and sophisticated hierarchical and round-robin learning classification ensemble algorithms used by Bodhidharma [8], whereas ACE only experiments with general-purpose machine learning algorithms.

When all three feature types were combined, the jMIR experiments described in this paper achieved a success rate of 78.8% which was still lower than Bodhidharma's performance, but significantly better than the best audio MIREX results to date.

Taken in the context of the particular difficulty of the SAC dataset, and when it is considered that the accuracy on the 10-genre taxonomy improves to 84.2% when weighted, the results attained here are encouraging, and may be an indication that the ultimate ceiling on performance might not be as low as some have worried [1]. It may well be that the use of more sophisticated machine learning approaches, such as those used by Bodhidharma or by DeCoro et al. [4], combined with the development of new features, could significantly improve performance further.

7. CONCLUSIONS AND FUTURE RESEARCH

The experimental results indicate that it is indeed substantively beneficial to combine features extracted from audio, symbolic and cultural data sources, at least in the case of automatic genre classification. Further research remains to be performed investigating whether these benefits generalize to other areas of music classification and similarity research.

All feature groups consisting of two feature types performed significantly better than any single feature types classified alone. Combining all three feature types resulted in small further improvement over the feature type pairs on average, but these additional improvements were not as uniform nor were they statistically significant.

The results also indicate that combining feature types tends to cause those misclassifications that do occur to be less serious, as the misclassifications are more likely to be

to a class that is more similar to the model class. Such improvements were particularly pronounced when cultural features were involved.

Encouragingly high genre classification accuracy rates were attained. The results of the experiments as a whole provide hope that any ultimate ceiling on genre classification performance might not be as low as has been worried.

The jMIR software suite was demonstrated to be an effective and convenient tool for performing feature extraction and classification research. The SAC dataset was also found to provide a good basis for performing combined audio, symbolic and cultural experiments.

An important next step is to repeat the experiments performed here with MIDI files transcribed from audio, in order to investigate more practically significant use cases where MIDI files do not have to be manually harvested. This would also enable experiments on larger datasets.

There are also plans to combine feature types in more sophisticated ways, such as by segregating them among different weighted specialist classifiers collected into blackboard ensembles. Sophisticated classification techniques that take advantage of ontological structuring, such as those utilized in Bodhidharma, also bear further investigation, and could fruitfully be incorporated into ACE. There are also plans to expand SAC as well as to apply jMIR to a wider variety of MIR research applications, such as mood and artist classification.

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THE QUEST FOR MUSICAL GENRES: DO THE EXPERTS AND THE WISDOM OF CROWDS AGREE?

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ABSTRACT

This paper presents some findings around musical genres. The main goal is to analyse whether there is any agreement between a group of experts and a community, when defining a set of genres and their relationships. For this purpose, three different experiments are conducted using two datasets: the *MP3.com* expert taxonomy, and *last.fm* tags at artist level. The experimental results show a clear agreement for some components of the taxonomy (*Blues*, *Hip-Hop*), whilst in other cases (e.g. *Rock*) there is no correlations. Interestingly enough, the same results are found in the MIREX2007 results for audio genre classification task. Therefore, a multi–faceted approach for musical genre using expert based classifications, dynamic associations derived from the wisdom of crowds, and content–based analysis can improve genre classification, as well as other relevant MIR tasks such as music similarity or music recommendation.

1 INTRODUCTION

Music genres are connected to emotional, cultural and social aspects, and all of them influence our music understanding. The combination of these factors produce a personal organization of music which is, somehow, the basis for (human) musical genre classification. Indeed, musical genres have different meanings for different people, communities, and countries [2].

The use of musical genres has been deeply discussed by the MIR community. A good starting point is the review by McKay [5]. The authors suggested that musical genres are an inconsistent way to organize music. Yet, musical genres remain a very effective way to describe and tag artists.

Broadly speaking, there are two complementary approaches when defining a set of genre labels: (*i*) the definition of a controlled vocabulary by a group of experts or musicologists, and (*ii*) the collaborative effort of a community (social tagging). The goal of the former approach is the creation of a list of terms, organised in a hierarchy. A hierarchy includes the relationships among the terms; such as hyponymy. The latter method, social tagging, is a less formal bottom–up approach, where the set of terms emerge during the (manual) annotation process. The output of this approach is called folksonomy.

The aim of this paper is, then, to study the relationships between these two approaches. Concretely, we want to study whether the controlled vocabulary defined by a group of experts concord with the tag annotations of a large community.

Section 2 introduces the pros and cons of expert–based taxonomies and music folksonomies. To compare the similarities between both approaches, we gathered data from two different websites: a musical genre taxonomy from *MP3.com*, and a large dataset of artists' tags gathered from the *last.fm* community. Section 3 presents these datasets. The experimental results, presented in section 4, are conducted in order to analyse the relationships between the genres used in the *MP3.com* taxonomy, and the genre–tags annotated in the artist dataset from *last.fm*. Finally, section 5 concludes and summarizes the main findings.

2 MUSICAL GENRES CLASSIFICATION

2.1 Expert–based taxonomies

Depending on the application, taxonomies dealing with musical genres can be divided into different groups [6]: Music industry taxonomies, Internet taxonomies, and specific taxonomies.

Music industry taxonomies are created by recording companies and CD stores (e.g. RCA, Fnac, Virgin, etc.). The goal of these taxonomies is to guide the consumer to a specific CD or track in the shop. They usually use four different hierarchical levels: (1) Global music categories, (2) Sub-categories, (3) Artists (usually in alphabetical order), and (4) Album (if available).

Internet taxonomies are also created under commercial criteria. They are slightly different from the music industry taxonomies because of the multiple relationships that can be established between authors, albums, etc. The main property is that music is not exposed in a physical space (shelves). Obviously, exploiting the relationships among the items allows the end–user a richer navigation and personalization of the catalogue.

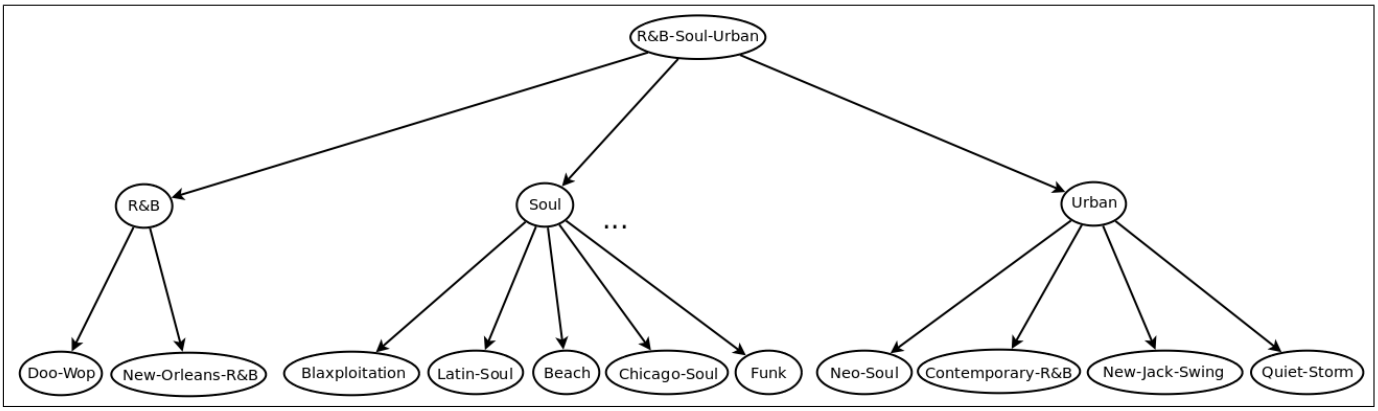

Figure 1. Partial view of the *MP3.com* taxonomy, starting with the seed genre *R&B–Soul–Urban*.

Furthermore, [6] shows that there is little consensus among the experts when defining a taxonomy. As an example, using three different musical genre taxonomies (*AllMusicGuide*, *Amazon*, and *MP3.com*) only 70 terms from more than 1500 were common in all the taxonomies.

2.2 Music Folksonomies

Since 2004, the explosion of Web 2.0 (e.g. tagging, blogging, user–generated content, etc.) questioned the usefulness of controlled vocabularies [11]. Internet sites with a strong social component, like *Last.fm*, allow users to tag music according to their own criteria. This scenario made the world of taxonomies even more complex.

Nowadays, users can organize their music collection using personal tags like *late night*, *while driving*, etc. As mentioned in the introduction, new strategies for music classification have emerged. Folksonomies exploit user–generated classification through a bottom–up approach [9].

On the one hand, this non-hierarchical approach allows users to organize their music with a better confidence. On the other hand, it creates difficulties for the design and maintenance of expert–based taxonomies, as new terms may emerge from time to time. Thus, in this scenario, up to date expert–based taxonomies become more and more difficult. Yet, it seems reasonable to analyse whether the genres derived from the tagging process share some patterns with the experts' controlled vocabulary.

3 DATASETS

3.1 Expert–based taxonomy from *MP3.com*

The *MP3.com* dataset was gathered during September 2005. Table 1 shows the relevant information about the genre taxonomy. Experts and musicologists from *MP3.com* identified 744 genres, and organized them in 13 different components, or in other words, the taxonomy has 13 seed-genres. The maximum depth is 6 levels, however, in most of the cases each component has 3 levels (plus the seed–genre at the top). Furthermore, this vocabulary still remains as it was three years ago (only a few more genres were added), showing its lack of evolution.

Total number of genres	744
Levels	7
Seed–genres	13
Num. genres at level 1	84
Num. genres at level 2	500
Num. genres at level 3	13
Num. genres at level 4	85
Num. genres at level 5	39
Num. genres at level 6	10

Table 1. Dataset gathered during September 2005 from the *MP3.com* expert–based taxonomy.

A partial view of the *MP3.com* taxonomy is depicted in Figure 1. It shows a component of the taxonomy. The seed genre is *R&B–Soul–Urban*, and the component consists of 3 different levels. A directed edge, e.g. *Urban* → *New-Jack-Swing*, represents the parent genre (*Urban*) and its subgenre(s), *New-Jack-Swing*.

3.2 Folksonomy from the *last.fm* community

A large dataset of artists' tags was gathered from the *last.fm* community during December 2007. Table 2 shows some basic information about the dataset. It is interesting to note that from the average number of tags per artist, 39% correspond to matched genres from the expert taxonomy, whilst the other 61% are distributed among other kinds of tags; including unmatched genres, decades, instruments, moods, locations, etc. For example, the artist *Jade* has the following tags (with their corresponding last.fm normalised weight):

```
Jade: urban(100), rnb(81), 90s(68),
      new jack swing(55), illinois(50),
      r and b(36), ...
```

Number of mapped genres	511
Number of artists	137,791
Number of distinct tags	90,078
Avg. tags per artist	11.95
Avg. *MP3.com* genres per artist	4.68

Table 2. Dataset gathered from the *last.fm* community during December 2007.

Nevertheless, since the experiments aim to analyse the agreement between expert genres and genre–tags, we need to match artists' tags with the expert defined genres.

3.2.1 Matching MP3.com genres and last.fm artist tags

In order to match those tags from the folksonomy that correspond to a genre in the expert taxonomy, a two-step process is followed:

- Compute a normalised form for all the folksonomy tags and expert genres, by:
  - converting them into lowercase,
  - unifying separators to a single common one,
  - treating some special characters (such as "&", which can be expanded to "and" and "n").
- Compute a string matching between the normalised folksonomy tags and expert genres.

The former step is inspired from [3]. For the latter, a string matching algorithm by Ratcliff and Metzener [8] is used to get all possible matches of a tag against a genre from the taxonomy. The similarity value goes from 0 to 1. Values close to 0 mean that the two strings are very dissimilar, and a 1 value means that the strings are identical. Deciding which is the threshold for identifying "nearly-identical" words is not trivial. Yet, [10] shows that a threshold of 0.85 gives the highest *F-measure*.

The following example shows artist *Jade*'s tags that are mapped to an *MP3.com* genre (*90s* and *illinois* tags disappear, and *rnb* and *r and b* are merged, combining their weights—with a maximum value of 100):

```
Jade: Urban(100), R&B(100),
      New-Jack-Swing(55)
```

Once the matching process is complete, the next step is to analyse whether the tagging behaviour of the community shares any resemblance with the expert taxonomy. The following section presents the experimental results.

4 EXPERIMENTAL RESULTS

In order to measure the agreement between expert genres and the genre–tags defined by the wisdom of crowds, we perform several experiments. Beforehand, we have to compute the similarities among genres. Section 4.1 explains the process of computing distances in the expert taxonomy (using the shortest path between two genres), and the tag distances in the folksonomy (by means of a classic Information Retrieval technique, called Latent Semantic Analysis).

The experiments are divided in two main groups. The first set of experiments deal with measuring the agreement at component level (a seed–genre and its subgenres). That is, to validate whether this taxonomy partition (13 components) correspond to the view of the community. Section 4.2 present these experiments. The other experiment focuses on the hierarchical structure (levels) of the expert taxonomy. In this experiment the goal is to reconstruct the taxonomy based on the genre distances from the folksonomy (section 4.3).

4.1 Computing genre distances

4.1.1 Expert taxonomy

To compute genre distances in the expert taxonomy we simply choose the shortest path between two genres, as an analogy with the number of mouse clicks to reach one genre from a given one (e.g. distance between genres *New–Jack–Swing* and *Soul* is 3, according to Figure 1). Since the taxonomy contains 13 seed genres, a virtual root node is added at the top, thus making the graph fully connected. This way we can compute the path between any genre in the graph. Whenever a path traverses the virtual root node, a penalty in the distance is added to emphasize that the two genres come from different components.

4.1.2 Folksonomy

Latent Semantic Analysis (LSA), plus cosine similarity, is used as a measure of distance among genres within the folksonomy. LSA assumes a latent semantic structure that lies underneath the randomness of word choice and spelling in "noisy" datasets [1], such as the one we are using. A significant paper that applies LSA in the music domain is [4]. The authors show the usefulness of social tags—in a low 10^2 space—to several relevant MIR problems, such as music similarity and mood analysis.

LSA makes use of algebraic techniques such as Singular Value Decomposition (SVD) to reduce the dimensionality of the Artist–Genres matrix. After this step, either artist or genre similarity can be computed using a cosine distance. Moreover, Information Retrieval literature [1, 7] states that, after raw data has been mapped into this *latent semantic space*, topic (in our case, genre) separability is improved.

	Folk	Bluegrass	Country	Electronic-Dance	New-Age	Rock-Pop	Jazz	Hip-Hop	R&B-Soul-Urban	Gospel-Spiritual	Vocal-Easy-Listening	Blues	World-Reggae
Folk	***0.184***	0.144*	0.048	0.002	0.040	-0.022	0.007	-0.005	0.003	0.095	0.001	-0.002	0.107
Bluegrass	0.144	***0.987***	0.738*	0.006	-0.018	0.008	-0.019	-0.006	-0.022	0.096	0.014	0.019	-0.033
Country	0.048	**0.738**	*0.430*	0.001	-0.034	0.048	0.007	0.007	0.009	0.054	0.031	0.008	-0.042
Electronic-Dance	0.002	0.006	0.001	*0.056*	0.019	0.012	0.025	0.004	0.019	0.036	**0.171**	0.002	-0.007
New-Age	0.040	-0.018	-0.034	0.019	***0.306***	0.125	0.001	0.022	0.018	0.068	0.102	0.037	0.303*
Rock-Pop	-0.022	0.008	0.048	0.012	0.125*	*0.043*	0.036	0.005	0.006	0.040	0.043	0.006	**0.132**
Jazz	0.007	-0.019	0.007	0.025	0.001	0.036	***0.211***	-0.008	0.030	-0.036	0.150	0.046	0.018
Hip-Hop	-0.005	-0.006	0.007	0.004	0.022	0.005	-0.008	***0.599***	-0.005	0.027	0.003	-0.005	0.021
R&B-Soul-Urban	0.003	-0.022	0.009	0.019	0.018	0.006	0.030	-0.005	***0.393***	0.355*	0.009	0.002	0.005
Gospel-Spiritual	0.095	0.096	0.054	0.036	0.068	0.040	-0.036	0.027	**0.355**	*0.134*	0.003	0.163	-0.015
Vocal-Easy-Listening	0.001	0.014	0.031	**0.171**	0.102	0.043	0.150	0.003	0.009	0.003	*0.167*	0.012	0.040
Blues	-0.002	0.019	0.008	0.002	0.037	0.006	0.046	-0.005	0.002	0.163	0.012	***0.657***	-0.004
World-Reggae	0.107	-0.033	-0.042	-0.007	**0.303**	0.132	0.018	0.021	0.005	-0.015	0.040	-0.004	*0.142*

Table 3. Confusion matrix for the inter–components coarse grained similarity. For clarification purposes, the diagonal contains the intra–component similarity. The values marked with an asterisk are as significative as the highest value (in bold).

For each artist we create a vector based on their (last.fm normalised) genre–tags' frequencies. Once the matrix is decomposed by columns, using SVD with 50 dimensions, we obtain genre similarities. For example, the closest genres to *Heavy Metal* in the semantic space are *Power Metal*, *British Metal* and *Speed Metal*, all with a similarity value above 0.6. On the other hand, similarity between *Heavy Metal* and *Pop* yields a near–zero value.

4.2 Agreement between expert and community genres

To measure the agreement between expert defined genre components and community genres we perform two experiments. The first one carries out a coarse–grained similarity (at genre component level), where the main goal is to *separate* the expert genre clusters according to the genre distances in the folksonomy. The second experiment performs a fine–grained similarity (at genre node level) in order to see the correlations between the genre distance in the taxonomy and the distance in the LSA space derived from the folksonomy.

4.2.1 Coarse-grained similarity

The first experiment aims to check how *separable* are the expert defined genre components according to the genre distances in the folksonomy (as defined in section 4.1.2). The experiment is performed in two steps: (*i*) compute the LSA cosine similarity among all the subgenres within a component (*intra–component* similarity); and (*ii*) compute the LSA cosine similarity among components, using the centroid of each component (*inter–component* similarity).

The results for intra–component similarity are presented in Figure 2. The most correlated components are *Bluegrass*, *Hip-Hop* and *Blues*. Note however that the *Bluegrass* component has only 3 subgenres mapped in our *last.fm* dataset. The components with less community–expert agreement are *Electronic-Dance* and *Rock-Pop*. For the latter genre, it is worth noting that it is an ill–defined seed-genre, and it is also the one including the highest number of subgenres. Some of these *Rock-Pop* subgenres are so eclectic that they could belong to more than one component. For instance, *Obscuro* subgenre is defined in *Allmusic* [1] as "*...a nebulous category that encompasses the weird, the puzzling, the ill-conceived, the unclassifiable, the musical territory you never dreamed*

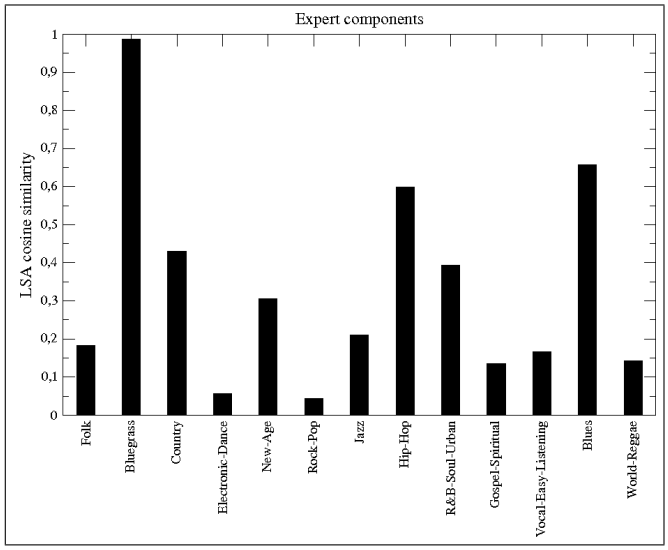

Figure 2. Intra–component coarse grained similarity.

[1] http://www.allmusic.com/

existed".

Regarding the inter–component similarity, we proceed as follows: we compute the centroid vector of each component, and then compare it with the remaining components' centroids. The results are presented in table 3. Note that the results in the diagonal represent the intra–components similarity. For each row, we mark in bold the highest value. Subgenres of *Bluegrass*, *Hip-Hop* and *Blues*, as it has been observed for the intra–component case, are highly correlated in the semantic space. Thus, they are the ones with more agreement between the community and the experts classification. However, only *Hip-Hop* and *Blues* are clearly distinguishable from the rest. Furthermore, according to the community, *Bluegrass* and *Country* genres are very similar. Indeed, other available internet taxonomies, such as *Amazon* or *Allmusic*, include *Bluegrass* as a subgenre of *Country*. Similarly, *Gospel-Spiritual* genre is merged into *R&B-Soul-Urban*.

4.2.2 Fine-grained similarity

In this experiment we focus on the genre node level (instead of components). The hypothesis is that genres closer in the semantic space of the folksonomy should also be closer in the expert taxonomy, and vice versa. To validate this formulation a one–way Anova is performed. The independent groups are considered the path distances in the expert taxonomy (ranging from $1..10$, the diameter of the taxonomy), whilst the dependent variable is the LSA cosine distance.

Figure 3 depicts the box–and–whisker plot. Indeed, a large value of the *F–statistic* as well as a *p–value* $\ll 0.05$ corroborates the hypothesis. Furthermore, to determine the distances that are statistically significant we perform the Tukey's pairwise comparisons test. The results show that path distances 1 and 2 are significant among the rest of the distances, at 95% family–wise confidence level.

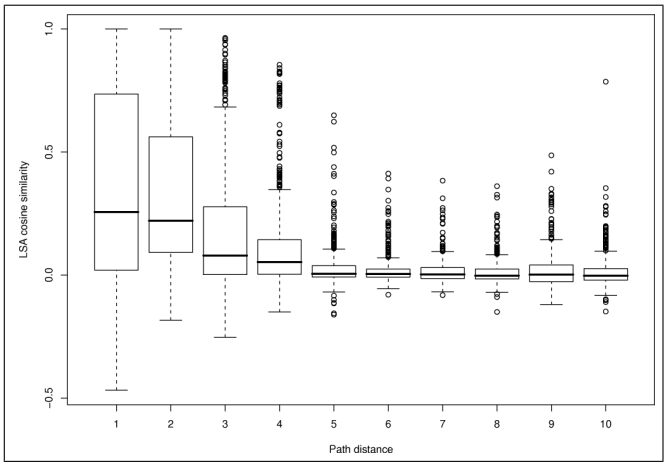

Figure 3. Box–and–whisker plot depicting the correlation between genre path distances in the taxonomy and semantic LSA cosine similarity. The Anova experiment ($p-value \ll 0.05$) shows that there is a statistical significance among path distances.

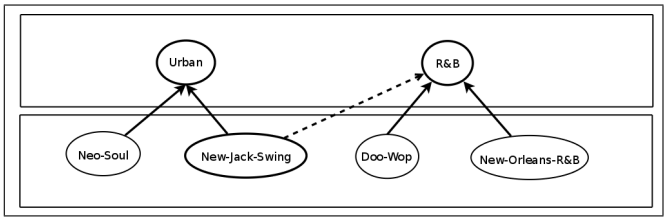

Figure 4. Reconstruction of the expert taxonomy from the folksonomy. Selection of a parent is made according to the LSA cosine similarity derived from the folksonomy.

4.3 Reconstructing the taxonomy from the folksonomy

In this experiment we try to reconstruct the taxonomy from the folksonomy's inferred semantic structure. The reconstruction of the expert taxonomy from the folksonomy is based on the correct selection of a parent genre, according to the LSA cosine similarity derived from the folksonomy. We follow a bottom–up approach, starting from the leaves of each component. At each step of the process, we record the differences between the inferred and original taxonomies in order to have a similarity metric between them.

The metrics used are: *mean reciprocal rank*, and *root hit*. The *mean reciprocal rank* (MRR) is a statistic widely used in Information Retrieval. The reciprocal rank (RR) is defined as the inverse of the correct answer's rank, $RR(tag) = 1/rank_{tag}$. For instance, given the *New–Jack–Swing* genre, see Figure 4, the closest genre parents (according to the LSA cosine distance) are: (1) *R&B*, (2) ***Urban***, (3) *Traditional–Gospel*, etc. The correct parent genre, *Urban*, is found in the second position, thus $RR(New - Jack - Swing) = \frac{1}{2} = 0.5$. Furthermore we compute whether the top–1 parent belongs to the same component as the child genre (**root hit**). In this example, it is a root hit because both the genre *New–Jack–Swing* and the selected (wrong) parent, *R&B*, belong to the same component, *R&B–Soul–Urban*.

Table 4 shows the results for each component. *Bluegrass*' perfect hit rate should be ignored, as the component has only 3 subgenres mapped. The lowest MRR is in *Rock–Pop* genre, which is also the highest (153 subgenres), and least cohesive component (lowest inter-genre similarity, see Table 3). *Hip–Hop*, on the other hand, is a highly cohesive component with a very high MRR. Finally, a lower MRR in *Folk* and *World–Reggae* could be interpreted as a consequence of the taxonomy being too geographically biased to English spoken territories. At the same time, disparate genres of all kinds, and from all over the world, are to be considered sub–genres of *Folk* and *World–Reggae*.

Component (size)	Root Hits (%)	MRR
Folk (22)	36.3	0.447
Bluegrass (3)	100	1.000
Country (35)	85.8	0.636
Electronic-Dance (36)	30.6	0.391
New-Age (5)	40.0	0.700
Rock-Pop (135)	48.2	0.295
Jazz (83)	83.2	0.638
Hip-Hop (22)	81.8	0.894
R&B-Soul-Urban (26)	75.4	0.694
Gospel-Spiritual (7)	38.6	0.558
Vocal-Easy-Listening (11)	27.3	0.446
Blues (43)	81.4	0.455
World-Reggae (83)	53.0	0.389
Weighted Avg. (511)	60.4	0.478

Table 4. Reconstruction of the expert–taxonomy, using genre similarity derived from the folksonomy.

5 CONCLUSIONS

This paper presented some interesting findings around musical genres. First of all, the consensus from a group of experts to create a universal taxonomy seems difficult. While expert taxonomies are useful for cataloguing and hierarchical browsing, the flat view of folksonomies allows better organization and access of a personal collection.

We presented three different experiments to analyse the agreement between expert–based controlled vocabulary and bottom–up folksonomies. The first two experiments focused on measuring the agreement between genres from the folksonomy and expert genres. A third experiment emphasized the hierarchical structure of a taxonomy, but using the information from a folksonomy. In all the experiments the conclusions were the same: some genres are clearly defined both from the experts and the wisdom of crowds, reaching a high agreement between these two views, while other genres are difficult to get a common consensus of its meaning.

All in all, experts, wisdom of crowds, and machines [2] agree in the classification and cohesion of some genres (e.g. *Blues*, *Hip-Hop*), and clearly disagree in others (*e.g. Rock*). A multi–faceted approach for musical genre using expert based classifications, dynamic associations derived from the community driven annotations, and content–based analysis would improve genre classification, as well as other relevant MIR tasks such as music similarity or music recommendation.

[2] See the results of MIREX 2007 in http://www.music-ir.org/mirex/2007/index.php/Audio_Genre_Classification_Results

6 ACKNOWLEDGMENTS

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Poster Session 1

TERNARY SEMANTIC ANALYSIS OF SOCIAL TAGS FOR PERSONALIZED MUSIC RECOMMENDATION

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ABSTRACT

Social tagging is the process by which many users add metadata in the form of keywords, to annotate information items. In case of music, the annotated items can be songs, artists, albums. Current music recommenders which employ social tagging to improve the music recommendation, fail to always provide appropriate item recommendations, because: (i) users may have different interests for a musical item, and (ii) musical items may have multiple facets. In this paper, we propose an approach that tackles the problem of the multimodal use of music. We develop a unified framework, represented by a 3-order tensor, to model altogether users, tags, and items. Then, we recommend musical items according to users multimodal perception of music, by performing latent semantic analysis and dimensionality reduction using the Higher Order Singular Value Decomposition technique. We experimentally evaluate the proposed method against two state-of-the-art recommendations algorithms using real Last.fm data. Our results show significant improvements in terms of effectiveness measured through recall/precision.

1 INTRODUCTION

Social tagging is the process by which many users add metadata in the form of keywords to annotate and categorize information items such as songs, pictures, products. In general, social tagging is associated to the "Web 2.0" technologies and has already become an important source of information for recommendation. In the music domain, popular web systems such as Last.fm and MyStrands provide possibility for users to tag with free text labels an item of interest - e.g., artist, song, album. Such systems can further exploit these social tags to improve the search mechanisms and the personalized music recommendation.

Recent research in the music field has also focused on exploiting the social tags in various ways. For example, a partial solution to the cold-start problem of music recommenders has been proposed in [4]: social tags are used for the automatic generation of new tags, which then can be used to label the untagged music. In [9], the social tags are investigated as a source of semantic metadata for music, which can be used to generate a psychologically-motivated search-space representing musical emotion.

However, the social tags carry useful information not only about the musical items they label, but also about the users who tagged. This aspect is not being fully exploited, neither by the music recommenders, neither in the research field. Music is an artistic concept, and the musical items (artists, songs, albums) have a rich and complex view, which is only partially perceived by particular users, depending on their emotional and cultural perspective on music. Social tags are a powerful mechanism that reveal 3-dimensional correlations between users–tags–items. This triplet information can project for each user his perception of a particular musical item. However, the current music recommender systems are commonly using collaborative filtering techniques, which traditionally exploit only pairs of 2-dimensional data. Thus, they are not capable of capturing well the multimodal use of music.

As a simple example, let us consider the social tagging system of artists in Last.fm. Assume two users. One is very fond of young female singers and therefore has tagged Christina Aguilera as "sexy" and Beyonce as "sensual". Another is fond of male singers and has tagged Lenny Kravitz as "sexy" and "male vocalists". When wanting to listen to "sexy" music, both users are recommended male and female singers, while the first user is expecting female singers and the other prefers the opposite.

Recent research has focused on developing recommendation algorithms [7, 14], which try to exploit tags given by users on specific items. However, the existing algorithms do not consider the 3 dimensions of the problem altogether, and therefore they miss a part of the semantics that is carried by the 3-dimensions.

In this paper, we address the problem of music recommendation by capturing the multimodal perception of music by particular users. We perform 3-dimensional analysis on the social tags data, attempting to discover the latent factors that govern the associations among the triplets user–tag–item. Consequently, the musical items (artists, songs or albums) can be recommended according to the captured associations. That is, given a user and a tag, the purpose is to predict whether and how much the user is likely to label with this tag a specific musical item.

Our strategy in dealing with the 3-dimensional social

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tagging data, is to develop a unified framework to model the three dimensions. Thus, user-tag-item data is represented by a 3-order tensor. Consequently, we have to deal with the data sparsity problem: the three-way data is highly sparse, especially that each user only tags a small number of items. Latent Semantic Indexing (LSI) has been proved useful to address the data sparseness in 2-dimensional data recommender systems, however, it is still an open problem for the 3-dimensional data case. Therefore, we perform 3-mode analysis, using the Higher Order Singular Value Decomposition (HOSVD) technique.

The contributions of our approach are as follow: **(1)** we provide a method to improve music recommendation by capturing users multimodal perception of music; **(2)** we develop a unified framework, represented by a 3-order tensor, to model the three types of entities that exist in social tagging data; **(3)** we apply dimensionality reduction in 3-order tensors to reveal the latent semantic associations between users, tags, and items; **(4)** we perform experimental comparison of the proposed method against two state-of-the-art recommendations algorithms, using Last.fm data; our results show significant improvements in terms of effectiveness measured through recall/precision.

The rest of this paper is organized as follows. Section 2 summarizes the related work, whereas Section 3 briefly reviews background techniques employed in our approach. A motivating example and the proposed approach are described in Section 4. Experimental results are given in Section 5. Finally, Section 6 concludes this paper.

2 RELATED WORK

Music recommendation has been addressed in various work. For example, in Logan [11] music recommendation is done based solely on using acoustic-based similarity measure. Other approaches try to bridge the semantic gap and employ hybrid music recommendation methods. Thus, Yoshii et al. [15] model collaborative filtering (CF) data and audio-content data together, and unobservable user preferences are statistically estimated. Li et al. [10] employ a probabilistic model estimation for CF, where musical items are clustered based on audio-content and user rating, and predictions are made considering the Gaussian distribution of ratings. Celma [2] mines music information from the Web (album releases, MP3 blogs, etc.) and is using it together with user profiling and audio-content descriptions.

The above work can be used to improve the music recommendation by addressing the cold-start problem and the bias of CF towards mainstream music. Along the same lines, an innovative use of social tags has been recently proposed in [4]. Eck et al. [4] predict new tags using audio features extracted from music and supervised learning. These automatically-generated tags resemble the characteristics of those generated by social taggers, and can be used to label new or poorly tagged music.

However, current music recommenders fail to always provide good recommendations, because they do not capture well the interest of particular users in musical items that have multiple facets. Recent research work [4] envisaged that musical items have multiple facets, but it did not address their multimodal perception by particular users. Since social tagging data carry simultaneous information about both the items and the users who tagged them, we propose to use such data as means to improve music recommendation by capturing the multimodal use of music.

The characteristics of social tagging systems have been already studied in the literature. Halpin et al. [6] claimed that there are three main entities in any tagging system: users, items, and tags. In contrast to the above ternary relation, recommender systems apply collaborative filtering on 2-dimensional spaces. For example, the approach of projecting the 3-dimensional space of social tagging into pair relations {user, item}, {user, tag}, {tag, item}, is applied in well-known recommendation algorithms such as Penalty-Reward and FolkRank. The Collaborative Tag Suggestions algorithm [14], also known as Penalty-Reward (PR), uses an authority score for each user, which measures how well each user has tagged in the past. This authority score can be computed via an iterative algorithm such as HITS [8]. FolkRank algorithm [7] is inspired by the seminal PageRank [12] algorithm. The key idea of FolkRank is that an item, which is tagged with important tags by important users, becomes important itself (and the same holds for tags and users).

However, the above state-of-art algorithms miss a part of the total interaction between the three dimensions of the social tagging space. In contrast, our approach develops a unified framework to concurrently model the three dimensions by employing a 3-order tensor, on which latent semantic analysis is performed using HOSVD technique [3]. The HOSVD technique has been successfully applied for computer vision problems. We also use in our approach the work proposed in Wang and Ahuja [13], which present a novel multi-linear algebra-based method to reduce the dimensionality representation of multi-dimensional data.

3 PRELIMINARIES - TENSORS AND HOSVD

In the following, we denote tensors by calligraphic uppercase letters (e.g., $\mathcal{A}$, $\mathcal{B}$), matrices by uppercase letters (e.g., A, B), scalars by lowercase letters (e.g., a, b), and vectors by bold lowercase letters (e.g., $\mathbf{a}$, $\mathbf{b}$).

SVD and Latent Semantic Indexing

The singular value decomposition (SVD) [1] of a matrix $F_{I_1 \times I_2}$ can be written as a product of three matrices, as shown in Equation 1:

$$F_{I_1 \times I_2} = U_{I_1 \times I_1} \cdot S_{I_1 \times I_2} \cdot V^T_{I_2 \times I_2,} \qquad (1)$$

where U is the matrix with the left singular vectors of F, V^T is the transpose of the matrix V with the right singular vectors of F, and S is the diagonal matrix of (ordered) singular values of F.

By preserving only the largest $c < \min\{I_1, I_2\}$ singular values of S, SVD results to matrix $\hat{F}$, which is an approximation of F. In Information Retrieval, this technique is used by Latent Semantic Indexing (LSI) [5], to deal with the latent semantic associations of terms in texts

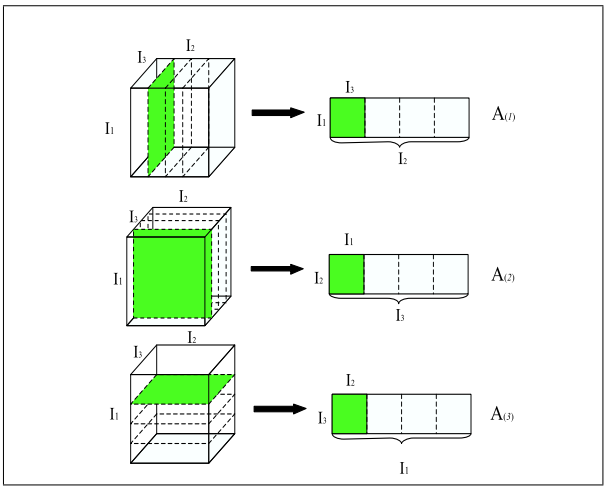

Figure 1. Visualization of the three unfoldings of a 3-order tensor.

and to reveal the major trends in F. The tuning of c is empirically determined by the information percentage that is preserved compared to the original matrix [3].

Tensors

A *tensor* is a multi-dimensional matrix. A N-order tensor $\mathcal{A}$ is denoted as $\mathcal{A} \in R^{I_1 \cdots I_N}$, with elements $a_{i_1,\ldots,i_N}$. In this paper, for the purposes of our approach, we only use 3-order tensors.

HOSVD

The high-order singular value decomposition [3] generalizes the SVD computation to multi-dimensional matrices. To apply HOSVD on a 3-order tensor $\mathcal{A}$, three *matrix unfolding* operations are defined as follows [3]:

$$A_1 \in R^{I_1 \times I_2 I_3}, \qquad A_2 \in R^{I_2 \times I_1 I_3}, \qquad A_3 \in R^{I_1 I_2 \times I_3}$$

where A_1, A_2, A_3 are called the 1-mode, 2-mode, 3-mode matrix unfoldings of $\mathcal{A}$, respectively. The unfoldings of $\mathcal{A}$ in the three modes is illustrated in Figure 1.

Example: Define a tensor $\mathcal{A} \in R^{3\times2\times3}$ by $a_{111} = a_{112} = a_{211} = -a_{212} = 1, a_{213} = a_{311} = a_{313} = a_{121} = a_{122} = a_{221} = -a_{222} = 2, a_{223} = a_{321} = a_{323} = 4, a_{113} = a_{312} = a_{123} = a_{322} = 0$. The tensor and its 1-mode matrix unfolding $A_1 \in R^{I_1 \times I_2 I_3}$ are illustrated in Figure 2.

Next, we define the n-mode product of an N-order tensor $\mathcal{A} \in R^{I_1 \times \ldots \times I_N}$ by a matrix $U \in R^{J_n \times I_n}$, which is denoted as $\mathcal{A} \times_n U$. The result of the n-mode product is an $(I_1 \times I_2 \times \ldots \times I_{n-1} \times J_n \times I_{n+1} \times \ldots \times I_N)$-tensor, the entries of which are defined as follows:

$$(\mathcal{A} \times_n U)_{i_1 i_2 \ldots i_{n-1} j_n i_{n+1} \ldots i_N} = \sum_{i_n} a_{i_1 i_2 \ldots i_{n-1} i_n i_{n+1} \ldots i_N} u_{j_n i_n} \quad (2)$$

Since we focus on 3-order tensors, $n \in \{1, 2, 3\}$, we use 1-mode, 2-mode, and 3-mode products.

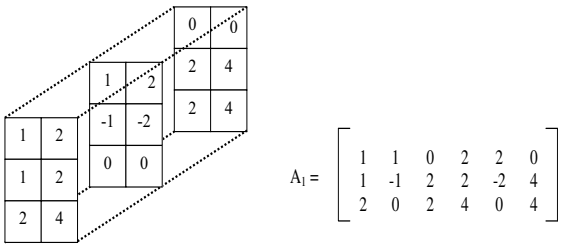

Figure 2. Visualization of tensor $\mathcal{A} \in R^{3\times2\times3}$ and its 1-mode matrix unfolding.

In terms of n-mode products, SVD on a regular two-dimensional matrix (i.e., 2-order tensor), can be rewritten as follows [3]:

$$F = S \times_1 U^{(1)} \times_2 U^{(2)} \quad (3)$$

where $U^{(1)} = (u_1^{(1)} u_2^{(1)} \ldots u_{I_1}^{(1)})$ is a *unitary* $(I_1 \times I_1)$-matrix, $U^{(2)} = (u_1^{(2)} u_2^{(2)} \ldots u_{I_1}^{(2)})$ is a *unitary* $(I_2 \times I_2)$-matrix, and S is a $(I_1 \times I_2)$-matrix with the properties of: (i) pseudo-diagonality: $S = \mathrm{diag}(\sigma_1, \sigma_2, \ldots, \sigma_{\min\{I_1,I_2\}})$ (ii) ordering: $\sigma_1 \geq \sigma_2 \geq \ldots \geq \sigma_{\min\{I_1,I_2\}} \geq 0$.

By extending this form of SVD, HOSVD of 3-order tensor $\mathcal{A}$ can be written as follows [3]:

$$\mathcal{A} = \mathcal{S} \times_1 U^{(1)} \times_2 U^{(2)} \times_3 U^{(3)} \quad (4)$$

where $U^{(1)}$, $U^{(2)}$, $U^{(3)}$ contain the orthonormal vectors (called the 1-mode, 2-mode and 3-mode singular vectors, respectively) spanning the column space of the A_1, A_2, A_3 matrix unfoldings. S is the core tensor and has the property of all orthogonality.

4 THE PROPOSED APPROACH

We first provide the outline of our approach, which we name Tensor Reduction, through a motivating example. Next, we analyze the steps of the proposed algorithm.

4.1 Outline

When using a social tagging system, to be able to retrieve information items easily, a user u labels with a tag t an item i. After some time of usage, the tagging system accumulates a collection of data – hence, *usage data*, which can be represented by a set of triplets $\{u, t, i\}$.

Our Tensor Reduction approach applies HOSVD on the 3-order tensor constructed from these usage data. In accordance with the HOSVD technique introduced in Section 3, the Tensor Reduction algorithm receives as input the usage data of $\mathcal{A}$ and outputs the reconstructed tensor $\hat{\mathcal{A}}$. $\hat{\mathcal{A}}$ measures the associations among the users, tags, and items. The elements of $\hat{\mathcal{A}}$ can be represented by a quadruplet $\{u, t, i, p\}$, where p measures the likeliness that user u will label with tag t an item i . Therefore, items can be recommended to u according to their weights associated with $\{u, t\}$ pair.

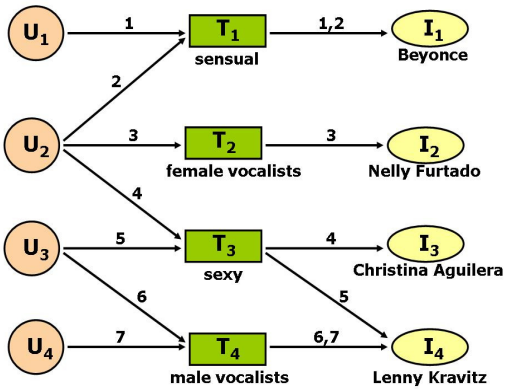

Figure 3. Usage data of the running example

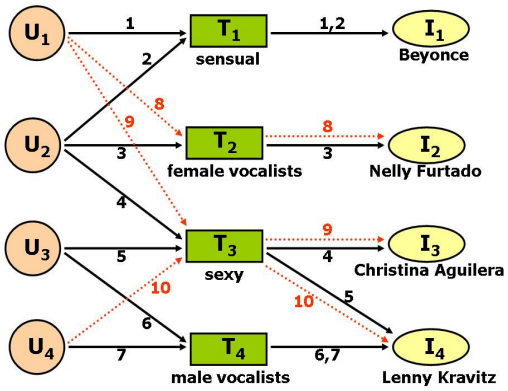

Figure 4. Illustration of the Tensor Reduction Algorithm output for the running example

In order to illustrate how our approach works, we apply the Tensor Reduction algorithm to a running example, which is illustrated in Figure 3. As it can be seen, 4 users tagged 4 different items. In the figure, the arrow lines and the numbers on them give the correspondence between the three types of entities. For example, user U_1 tagged with tag "sensual" (denoted as T_1) the item "Beyonce" (denoted as I_1). From Figure 3, we can see that users U_1 and U_2 have common interests on female singers, while users U_3 and U_4 have common interests in male singers.

A 3-order tensor $\mathcal{A} \in R^{4\times4\times4}$ can be constructed from these usage data. We use the co-occurrence frequency of user, tag and item as the elements of tensor $\mathcal{A}$, which are given in Table 1.

Arrow Line	User	Tag	Item	Weight
1	U_1	T_1	I_1	1
2	U_2	T_1	I_1	1
3	U_2	T_2	I_2	1
4	U_2	T_3	I_3	1
5	U_3	T_3	I_4	1
6	U_3	T_4	I_4	1
7	U_4	T_4	I_4	1

Table 1. Tensor Constructed from the usage Data of the running example.

After performing the Tensor Reduction analysis (details are given in the section 4.2), we get the reconstructed tensor of $\hat{\mathcal{A}}$. Table 2 gives the output of the Tensor Reduction algorithm, which is also illustrated in Figure 4.

We can notice that the algorithm outputs new associations among the involved entities (see the last rows in the Table 2 and the dotted lines in Figure 4). Even though in the original data, user U_1 did not tag items I_2 and I_3, the algorithm is capable to infer that if U_1 would tag them, then U_1 would likely (likelihood 0.35) use tags "female vocalists", and respectively "sexy". As well, the algorithm can infer that if U_4 would tag item I_4 with another tag, then U_4 would likely (likelihood 0.44) use the tag "sexy".

We judge the obtained results as reasonable since U_1 appears to be concerned with female singers rather than with male singers, and viceversa for U_4 . That is, the Tensor Reduction approach is able to capture the latent associations among the multi-type data entities: users, tags and musical items. These associations can further be used to improve the recommendation procedure of musical items.

Arrow Line	User	Tag	Item	Weight
1	U_1	T_1	I_1	0.50
2	U_2	T_1	I_1	1.20
3	U_2	T_2	I_2	0.85
4	U_2	T_3	I_3	0.85
5	U_3	T_3	I_4	0.72
6	U_3	T_4	I_4	1.17
7	U_4	T_4	I_4	0.72
8	$\mathbf{U_1}$	$\mathbf{T_2}$	$\mathbf{I_2}$	**0.35**
9	$\mathbf{U_1}$	$\mathbf{T_3}$	$\mathbf{I_3}$	**0.35**
10	$\mathbf{U_4}$	$\mathbf{T_3}$	$\mathbf{I_4}$	**0.44**

Table 2. Tensor Constructed from the usage data of the running example.

4.2 The Tensor Reduction Algorithm

In this section, we elaborate on how HOSVD is applied on tensors and on how the recommendation of musical items is performed according to the detected latent associations.

Our Tensor Reduction algorithm initially constructs a tensor, based on usage data triplets $\{u, t, i\}$ of users, tags and items. The motivation is to use all three entities that interact inside a social tagging system. Consequently, we proceed to the unfolding of $\mathcal{A}$, where we build three new matrices. Then, we apply SVD in each new matrix. Finally, we build the core tensor $\mathcal{S}$ and the resulting tensor $\hat{\mathcal{A}}$. All these can be summarized in 6 steps, as follows.

Step 1. The initial construction of tensor $\mathcal{A}$

From the usage data triplets (user, tag, item), we construct an initial 3-order tensor $\mathcal{A} \in R^{u\times t\times i}$, where u, t, i are the numbers of users, tags and items, respectively. Each tensor element measures the preference of a (user u, tag t) pair on an item i.

Step 2. Matrix unfolding of tensor $\mathcal{A}$

As described in Section 3, a tensor $\mathcal{A}$ can be matricized i.e., to build matrix representations in which all the column (row) vectors are stacked one after the other. In our approach, the initial tensor $\mathcal{A}$ is matricized in all three modes. Thus, after the unfolding of tensor $\mathcal{A}$ for all three modes, we create 3 new matrices A_1, A_2, A_3, as follows:

$$A_1 \in R^{I_u \times I_t I_i}, \qquad A_2 \in R^{I_t \times I_u I_i}, \qquad A_3 \in R^{I_u I_t \times I_i}$$

Step 3. Application of SVD in each matrix

We apply SVD on the three matrix unfoldings A_1, A_2, A_3. We result to total 9 new matrices.

$$A_1 = U^{(1)} \cdot S_1 \cdot V_1^T \tag{5}$$

$$A_2 = U^{(2)} \cdot S_2 \cdot V_2^T \tag{6}$$

$$A_3 = U^{(3)} \cdot S_3 \cdot V_3^T \tag{7}$$

For tensor dimensionality reduction, there are three parameters to be determined. The numbers c_1, c_2, and c_3 of left singular vectors of matrices $U^{(1)}$, $U^{(2)}$, $U^{(3)}$ which are retained, are determinative for the final dimension of the core tensor S. Since each of the three diagonal singular matrices S_1, S_2, and S_3 are calculated by applying SVD on matrices A_1, A_2 and A_3 respectively, we use different c_1, c_2, and c_3 values for each matrix $U^{(1)}, U^{(2)}, U^{(3)}$.

The numbers c_1, c_2, and c_3 are empirically chosen by preserving a percentage of information of the original S_1, S_2, S_3 matrices after appropriate tuning (usually the percentage is set to 50% of the original matrix).

Step 4. The core tensor $\mathcal{S}$ construction

The core tensor S governs the interactions among user, item and tag entities. Since we have selected the dimensions of $U^{(1)}$, $U^{(2)}$, and $U^{(3)}$ matrices, we proceed to the construction of the core tensor S, as follows:

$$\mathcal{S} = \mathcal{A} \times_1 {U_{c_1}}^{(1)^T} \times_2 {U_{c_2}}^{(2)^T} \times_3 {U_{c_3}}^{(3)^T}, \tag{8}$$

where $\mathcal{A}$ is the initial tensor, ${U_{c_1}}^{(1)^T}$ is the transpose of the c_1-dimensionally reduced $U^{(1)}$ matrix, ${U_{c_2}}^{(2)^T}$ is the transpose of the c_2-dimensionally reduced $U^{(2)}$, and ${U_{c_3}}^{(3)^T}$ is the transpose of the c_3-dimensionally reduced $U^{(3)}$.

Step 5. The tensor $\hat{\mathcal{A}}$ construction

Finally, tensor $\hat{\mathcal{A}}$ is build by the product of the core tensor $\mathcal{S}$ and the mode products of the three matrices $U^{(1)}$, $U^{(2)}$ and $U^{(3)}$ as follows:

$$\hat{\mathcal{A}} = \mathcal{S} \times_1 {U_{c_1}}^{(1)} \times_2 {U_{c_2}}^{(2)} \times_3 {U_{c_3}}^{(3)}, \tag{9}$$

where $\mathcal{S}$ is the c_1, c_2, c_3 reduced core tensor, ${U_{c_1}}^{(1)}$ is the c_1-dimensionally reduced $U^{(1)}$ matrix, ${U_{c_2}}^{(2)}$ is the c_2-dimensionally reduced $U^{(2)}$ matrix, ${U_{c_3}}^{(3)}$ is the c_3-dimensionally reduced $U^{(3)}$ matrix.

Step 6. The generation of the item recommendations

The reconstructed tensor $\hat{\mathcal{A}}$ measures the associations among the users, tags and items, so that the elements of $\hat{\mathcal{A}}$ represent a quadruplet $\{u, t, i, p\}$ where p is the likeliness that user u will tag musical item i with tag t. Therefore, musical items can be recommended to u according to their weights associated with $\{u, t\}$ pair.

5 EXPERIMENTAL CONFIGURATION

In this section, we study the performance of our approach against two well-known recommendation algorithms: Collaborative Tag Suggestions [14] (known as Penalty-Reward) and FolkRank [7], denoted as PR and FolkRank, respectively. We denote our algorithm as Tensor Reduction.

5.1 Data Set

To evaluate the aforementioned algorithms we have chosen a real data set mined from Last.fm. The data was gathered during October 2007, by using Last.fm web services. The musical items correspond to artist names, which are already normalized by the system. There are 12,773 triplets in the form user–tag–artist. To these triplets correspond 4,442 users, 2,939 tags and 1,620 artists. (We only kept tags expressing positive preference.)

To get more dense data, we follow the approach of [7] and we adapt the notion of a p-core to tri-partite hypergraphs. The p-core of level k has the property, that each user, tag and item occurs in at least k posts. We use k = 5, and we finally retain 112 users, 567 tags, and 234 artists.

5.2 Evaluation Metrics

We perform 4-fold cross validation and the default size of the training set is 75% – we pick, for each user, 75% of his posts randomly. The task of each item recommendation algorithm is to predict the items of the user's 25% remaining posts. As performance measures we use precision and recall, which are standard in such scenarios.

For a test user that receives a list of N recommended items (top-N list), the following are defined:
– *Precision*: is the ratio of the number of relevant items in the top-N list (i.e., those items in the top-N list that are used/tagged by the test user) relative to N.
– *Recall*: is the ratio of the number of relevant items in the top-N list relative to the total number of relevant items (all items used/tagged by the test user).

5.3 Settings of the Algorithms

Tensor Reduction algorithm: The numbers c_1, c_2, and c_3 of left singular vectors of matrices $U^{(1)}$, $U^{(2)}$, $U^{(3)}$ after appropriate tuning are set to 40, 80 and 190. Due to lack of space we do not present experiments for the tuning of c_1, c_2, and c_3 parameters. The core tensor dimensions are fixed, based on the aforementioned c_1, c_2, and c_3 values.
FolkRank algorithm: We set the damping factor d = 0.7 and stop computation after 10 iterations or when the distance between two consecutive weight vectors is less than 10^{-6}. For the preference vector $\mathbf{p}$, we give higher weights to the user and the tag from the post which is chosen. While each user, tag and item gets a preference weight

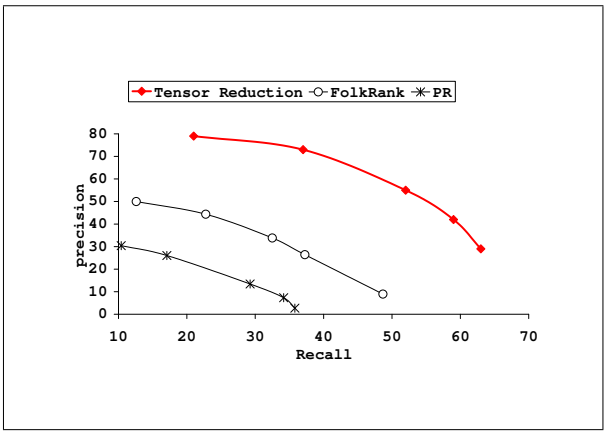

Figure 5. Comparison of the Tensor Reduction, FolkRank and the PR algorithms for the Last.fm data set

of 1, the user and tag from that particular post gets a preference weight of $1 + |U|$ and $1 + |T|$, respectively.
PR algorithm: Initially, we set the uniform authority score for each user equal to 1.0. Then, the authority score of users is computed via an iterative algorithm similar to HITS.

5.4 Comparison Results

We compare the Tensor Reduction algorithm with FolkRank and PR, in terms of precision and recall. This reveals the robustness of each algorithm in attaining high recall with minimal losses in terms of precision. We examine the top-N ranked list, which is recommended to a test user, starting from the top item. In this situation, the recall and precision vary as we proceed with the examination of the top-N list. For our data set, N is between [1..5].

In Figure 5, we plot a precision versus recall curve for all three algorithms. As it can be seen, Tensor Reduction algorithm attains the best performance. The reason is that Tensor Reduction exploits altogether the information that concerns the three entities (users, tags, items) and thus, it is able to provide more accurate recommendations.

6 CONCLUSIONS

We developed a unified framework to model the three types of entities that exist in social tagging systems: users, tags and items. This data is represented by a 3-order tensor, on which latent semantic analysis and dimensionality reduction is applied using the higher-order singular value decomposition technique. Our approach can be used to generate personalized recommendations of musical items, which can capture users multimodal perception of music. We performed thorough experimental comparison of our approach against two state-of-the-art recommendations algorithms on real Last.fm data. Our method achieves significant improvements measured through recall/precision.

As future work, we intend to apply different weighting schemes for the initial construction of a tensor. Also, we will adjust our Tensor Reduction framework so that the newly emerged users, tags or items of a social tagging system, can be handled online.

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THE LATIN MUSIC DATABASE

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ABSTRACT

In this paper we present the Latin Music Database, a novel database of Latin musical recordings which has been developed for automatic music genre classification, but can also be used in other music information retrieval tasks. The method for assigning genres to the musical recordings is based on human expert perception and therefore capture their tacit knowledge in the genre labeling process. We also present the ethnomusicology of the genres available in the database as it might provide important information for the analysis of the results of any experiment that employs the database.

1 INTRODUCTION

The maturation of the music information retrieval (MIR) field has required that researchers move beyond work involving simplistic musical databases to more expansive studies that require much larger, more varied and carefully annotated collections [19]. McKay et al. [19] have suggested a list of desired features for the development of new music databases in order to have music databases that can be employed in different types of research instead of only a few experiments. The suggested list of desired features in new databases can be divided into three major groups: related to the musical recordings, related to the underlying framework to efficiently use the musical recordings and considering the accessibility and distribution of the database.

The desired features related to the musical recordings are: (1) The database should contain many different types of music; (2) The database should include many thousands of recordings; (3) The database should include a significant amount of commercial music; (4) Each recording should be annotated with as diverse a range of metadata fields as possible, in order to make the database usable as ground truth for as wide a range of research as possible; (5) Entire recordings should be accessible, at least indirectly, not just short excerpts; (6) Given that different compression methods can influence extracted feature values, the audio format(s) most commonly used by the public should be adopted in order to reflect realistic conditions.

The desired features related to the underlaying framework are: (7) It should ideally be possible to label segments of recordings as well as recordings as a whole; (8) Annotations of subjective fields such as genre or mood should include a wide range of candidate categories; allowing ten or so coarse categories is unrealistic; (9) It should be possible to assign multiple independent values to a single field so that, for example, a recording could be classified as both swing and blues; (10) The ability to construct ontological structures between fields could be useful; (11) Metadata should be made available to users in formats that are easy to both manually browse and automatically parse; (12) Automatic tools should be available to validate metadata and generate profiles of the database; (13) It should be easy to add new recordings and their metadata.

The last desired feature is concerned with the accessibility and distribution of the database: (14) Data should be freely and legally distributable to researchers.

Another question that often arises is how to assign a music genre in order to create a musical genre database. In the work of McKay and Fujinaga [18] this issue has been extensively covered. We consider the following definition of a musical genre: "a kind of music, as it is acknowledged by a community for any reason or purpose or criteria" [9].

In this paper we present the Latin Music Database (LMD), a 3.227 musical recordings database from ten different genres. This database was firstly developed to the task of musical genre classification, but it can be used by any MIR task since it provides not only the full recordings in the MP3 format but also information about the music title and performing artist. The main motivation behind the development of this database is due to the lack of ground-truth in the area. Most of databases normally contains few musical recordings, sometimes only excerpts of the full music recording, and also a small number of instances per class. One of the reasons that hinders the development of novel databases is the difficultly in distributing them due to copyright restrictions of musical recordings. However with the creation of the OmeN (On-demand Metadata Extraction Network) tool [17] researchers now have a tool to overcomes this limitation and make databases widely available to the MIR community. In this work an approach to assign musical genres that corroborates with the definition of [9] is used. We also present information concerning how the LMD attends this list of desired features along with a brief ethnomusicology

description of the musical genres used in the database.

This paper is organized as follows: Section 2 presents the related work; Section 3 presents the development process of the LMD; The concluding remarks and a brief discussion about future work are presented in the last section; Appendix A presents the ethnomusicology of the genres used in the LMD.

2 RELATED WORK

The work of Tzanetakis and Cook [22] have strongly motivated the research in automatic music genre classification. The contributions from this work were three fold: First, it showed the task of an automatic music genre classification as a pattern recognition problem; Second, it not only proposed a feature set for the task but also developed an open source software for feature extraction know as Marsyas; and Third, the database used in the experiments is available. This database contains 1.000 music pieces from 10 musical genres (*Blues*, *Classic*, *Country*, *Disco*, *Hiphop*, *Jazz*, *Metal*, *Pop*, *Reggae* and *Rock*). In the remainder of this work this database will be as referred the GTZAN database. From this work onwards, different methods were developed based on extracting features from the audio signal [12].

Current work in the field is trying to break through the possible glass ceiling that content-based audio classification using timbral features has reached [2]. In [3] the authors proposed a method for considering context attributes in order to improve classification accuracy. The experiments were performed with database containing 4.936 songs with 801 boolean attributes grouped in 18 categories, some of which being correlated with some acoustic aspect of the sound ("Main Instrument", "Dynamics"), while others seem to result from a more cultural aspect of the sound ("Genre", "Mood"). This method achieved an improvement of 5% accuracy in average when compared to using only features extracted from the music signal. In [16] a framework for augmenting genre classification accuracy using an ensemble that considers two different views of the same audio signal. The first view is based on the audio signal content, while the second is based on a symbolic form that is obtained using a transcription system. The experiments were performed on three databases: GTZAN, ISMIR Rhythm and ISMIR Genre. The achieved results shown improvements over the previous results on the same databases.

Another important aspect that concerns current research is how the evaluation of music genre classification have been performed. In [10] is verified that performing classification without artist filters (i.e. using music pieces from the same artist in both training and test sets) not only lower the classification accuracy but may also erode the differences in accuracies between different techniques. In [7] is presented a discussion on how to define ground-truth for the task of music genre classification. In this work they define ground-truth as an artefact of an individual response to music, not an artefact of the audio itself; to establish any ground-truth is therefore a cultural study. An approach that comply with this definition is presented by Gouyon et al. [12]. In this work they used the instantly ability that dancers have to recognize different ballroom musical genres (*Jive*, *Quickstep*, *Tango*, *Waltz*, *Viennese Waltz*, *Cha Cha Cha*, *Samba* and *Rumba*) to label their database. These approaches are similar to the one used to define ground-truth for the LMD, as we will see in section 3.1.

Despite the efforts on the development on new techniques for the task of automatic music genre classification, relatively less work have been done concerning the development of music databases that can be used as ground-truth. Besides the GTZAN database, other known databases are: the RWC database [11], the CODAICH database [19], the magnatune database which was used in the MIREX 2004 [8] genre contest, and the database developed by Homburg et al. [14].

3 THE DEVELOPMENT OF THE DATABASE

We start the development of the LMD [1] aiming to collect at least 3.000 musical recordings from ten Latin musical genres. However, there are many issues involved in the development of a new audio database, as we will see in the following.

3.1 The Process of Assigning Musical Genres

In the development of the LMD we used an approach motivated by the human perception of how to assign musical genres to the musical recordings. The genre is defined according to how human listeners dance the musical recordings. The musical recordings were labeled by professional teachers with over ten years of experience in teaching ballroom and Brazilian cultural dances. In the first stage of the process these professionals made a selection of the musical recordings that they judged representative of a specific genre, according to how that musical recording is danced. In the second stage of the process the first author of this work verified each one of the selected songs in order to avoid mistakes that were expected to happen due to the stress produced by manually listening and labeling each one of the songs. About 300 musical recordings were classified by month, and the total duration of the development of the LMD took a year.

As a result of this attempt, we developed the Latin Music Database which currently has 3.227 musical recordings from ten Latin music genres in MP3 format. Table 1 presents information concerning the informetrics of the LMD. The original source of the recordings are CD's (which were encoded using 256 kbps and 44 kHz) and from different websites.

[1] Information available on: http://www.ppgia.pucpr.br/∼silla/lmd/

			Track Duration		
Genre	Artists #	Tracks #	Min.	Max.	Length
Axé	37	313	01:22	08:37	19:09:07
Bachata	64	313	02:54	06:32	23:23:22
Bolero	99	315	01:50	05:45	19:19:44
Forró	27	313	01:17	07:50	18:38:16
Gaúcha	92	311	01:13	07:32	17:32:29
Merengue	96	315	01:39	07:30	20:21:39
Pagode	16	307	00:52	11:12	19:41:27
Salsa	54	311	01:55	09:12	28:53:48
Sertaneja	9	321	02:04	06:34	23:03:10
Tango	7	408	01:19	12:06	19:02:26

Table 1. Latin Music Database Informetrics

The employed protocol of human inspection based on the perception about how that musical recordings are danced contrasts with the suggestion given by Aucouturier and Pachet [1]. They suggest to use complete CDs from collections of a given genre or theme. We justify the approach because the latter proposal, in the case of Latin music, is highly inefficient for labeling. For example in the case of the four CD collection *Los 100 Mayores Exitos De La Musica Salsa* only half (50 out of 100) of the songs can be classified as *Salsa*, the remaining of the collection belongs to other music genres like *Merengue*, *Lambada* and even *Samba*.

Another approach that could be used for automatically labeling the music is the classification of albums based on the artist profile, a common practice in the area. However this approach does not seems to be adequate again. For example, if we want to add songs by Carlos Gardel (the famous Tango Composer), all his songs would be labeled as *Tango*. Although he has over 500 compositions, only about 400 of them are actually *Tangos*, so the straightforward labeling introduces unnecessary noise from a machine learning perspective. Even with other artists from a specific genre hardly ever all the tracks from an album are from the same genre. One interesting fact that was perceived during the database creation is that about one of three songs in each album are not from the main style of the Artist profile.

3.2 Storage and Retrieval of the Musical Recordings

Besides the acquisition and genre labeling of the music pieces, much effort was employed in order to develop a database with the following characteristics: (1) it must be easily used in other MIR tasks; (2) it must allow the reproducibility of the performed experiments; (3) it must avoids duplicity in the stored musical recordings; and (4) it must be easily extended, with the storage of new music pieces, new music genres or any other kind of desired metadata (i.e. mood). Taking this issues in consideration a prototype system for storing, labeling and handling the musical recordings was developed. The proposed system is in accordance with the list of desirable features proposed by McKay et al. [19].

The process of adding a new music recording to the framework is as follows. First, a musical genre is assigned to the recording following the procedure explained at Section 3.1. Next, the ID3v2 tag (which contains meta-information) of the musical recording is manually inspected to verify if the fields have been correctly fulfilled. Sometimes we need to correct / adapt them according to a standardization procedure, such as in the case of names that contain the special character &, indicating more than one artist in the same song. This procedure could be replaced by an automated process like the one suggested by McKay et al. [19] as it mainly involves converting the music titles and artists to a pre-defined standard. To add a new music to the database the mandatory fields in the ID3v2 tag are the artist and the song title. The reason is simple: even if only one person is working on adding new songs to the database, eventually albums from the same artist can contain musical recordings that also appear in other albums, e.g. in the case of albums with greatest hits of a particular artist. The contribution of this simple procedure is two-fold: first, it avoids having duplicate songs in the database; second, it enforces that all music added to the database have the valuable metadata of song artist and title.

Finally, the musical recording is stored in the database. In this step the framework obtains the metadata from the ID3v2 tag of the musical recording and verify if the song is not already present in the database, to avoid duplicity. If there is no duplicity the framework assigns the song an identifier code, associate this song to the defined genre and creates an internal copy of the song. The information concerning the music genre is stored into a relational database, because the Genre field of ID3v2 tag is not reliable. Another reason for this approach is flexibility, as it makes the assignment of multiple genres per musical recording as simple as adding another entry into the relational database.

The process of retrieval in the framework is straightforward: it is started by the user, who specifies how many recordings from each musical genre should be retrieved. The framework also implements other modules that execute specific tasks, such as the integration of the database with feature extraction frameworks – like Marsyas [22] – or to allow the user to retrieve the feature set in a specific data format – such as in the arff format for using the WEKA machine learning tool [24].

Another important aspect of the framework is related to the reproducibility of performed experiments. Since every musical recording in the database has information concerning the song artist and title, it is possible to create along the output of the feature sets in arff format a list of the songs in the same order in which they are used by classification modules. The file used to store this list is called SAL (Song Artist List). There are three reasons for representing this information in the SAL format: (1) Sometimes different artists interpret the same songs using the same or a different style;

Database	No. Songs	No. Genres	Commercial Music	Meta Data	Full Recordings
CODAICH	20.849	53	Yes	Yes	Yes
Homburg et al.	1.886	9	No	Yes	No
GTZAN	1.000	10	Yes	No	No
LMD	3.227	10	Yes	Yes	Yes
Magnatune	2.181	6	No	Yes	Yes
RWC	100	10	No	Yes	Yes

Table 2. Comparison of the existing databases for musical genre classification

(2) Using only the system assigned ID we cannot obtain a reliable information, because if for some reason (i.e. backup) the musical recordings need to be restored in the system, they will probably not be added in the same order they were originally; (3) It might be easier to understand and interpret the achieved results if one knows which songs / artists were used in the experiment.

3.3 Analysis of the Database

In this section we present an explanation on how the LMD deals with the list of desired features in new databases indicated in [19]. As discussed in Section 1, the list of desired features can be divided into three major groups: related to the musical recordings, related to the underlying framework to efficiently use the musical recordings and considering the accessibility and distribution of the database.

Concerning the features related to the musical recordings, the LMD satisfy all the desired criteria because: (1) it has 10 Latin music genres; (2) 3.227 music pieces; (3) every musical recording in the LMD is commercial; (4) the fields from artist, tile, and musical genre have all been carefully annotated; (5) only full recordings are available; (6) every recoding is in the MP3 format.

Concerning the features related to the underlying framework, some of them are not fully implemented yet. However, they are less critical than the features related to the musical recordings, because doing implementation improvements is easier than doing carefully and correct annotation. In the current framework the issues that can be handled are as follows: features (7) and (9) can be handled by creating a table in the relational database of the framework to consider the segments of the music recordings, or to allow more than one genre per musical recording; (8) Although only 10 Latin music genres are presented in the database, it is important to stress that the classification was done by human experts and not using website listings in order to achieve ground-truth. Therefore, we would like to encourage more approaches as the one presented in this paper, even if few genres are presented as the classification by human experts is a really time-consuming task; (10) is beyond the scope of the designed framework; (11) and (12) the framework outputs information in arff and SAL formats, and could be easily modified to deal with other formats as needed; (13) as seen in section 3.2 it is a straightforward process.

The last desired feature is concerned with the accessibility and distribution of the database. Due to copyright reasons, the LMD cannot be made freely available to the research community; however there are on-going efforts in two directions to disseminate the use of the database. The first direction is making the LMD available through the OmeN tool [17]. The second direction is contributing the LMD to the MIREX (Music Information Retrieval Evaluation eXchange) community [8].

A summary of the different databases according to the list of desired features related to the musical recordings as seen in Section 1 is presented in Table 2. An analysis of this table shows that besides CODAICH, only the LMD has all the desired features related to the musical recordings, due to its large number of entire commercial songs associated to metadata.

4 CONCLUDING REMARKS

In this work we present a novel and carefully developed database for the task of automatic music genre classification, namely the Latin Music Database (LMD). An analysis of the database shows that the LMD has all the desired features listed in the work of McKay et al. [19], with one exception: the legal questions related to music recording distribution. However two efforts are being made in order to make the LMD available to the research community: (a) contributing the database to the MIREX community; and (b) using it in the OMEN framework. The method for assigning musical genres although similar in concept to the one used in [12] differs from it for the following reasons: (1) the LMD has a greater number of musical recordings; (2) the genre labeling process is based on two human experts instead of assuming that the classification given by a website is the ground-truth; and (3) most of the genres used are different.

Another contribution of this work is that it presented briefly the ethnomusicology aspects of the genres used in LMD. This is an important aspect of the database, considering the current state of the art in the field of automatic music genre classification: understanding the ethnomusicology of the genres being worked is crucial to understand the social context to which the genres belong. This information along with the full metadata for the music pieces in the database allow that

context information may be explored in novel ways. One interesting aspect of future research is training a classifier based on the lyrics of the musical recordings, in order to provide further information about the context of specific genres or sub-genres. Another goal for future research is to expand the LMD and further classify the existing songs into their sub-genres categories.

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A ETHNOMUSICOLOGY OF THE DATABASE

One important aspect of the research on music genre classification that is often forgot is how it can contribute to music-related fields like musicology. This might be due to the fact that most approaches still use only content based information extracted from the audio signal [12, 22]. However, as any other application in data mining, in order to better understand the results achieved by the state of art algorithms, prior knowledge about the data itself is needed. For the task of music genre classification, this can be achieved from the field of Ethnomusicology, which considers not only the origin of the music genres but also the social and cultural context in which it exists. In this section, we present a brief analysis of the music genres used in the database in the hope of: (a) a better understanding of any results using the database; (b) a briefly description of what some of the unknown genres are; (c) act as starting reference points to books that discuss the social and cultural aspects of the musical genres used in the LMD.

Axé: It is a Brazilian music genre, originated in Salvador, Bahia, where rhythm has been the key ingredient,

which typically feature simple melodies and harmonies. It is energetic, good-time party music, designed for dancing and Carnival. The lyrics usually focus on romance, sex, and Carnival [20].

Bachata: Closely associated with poor rural migrants residing in urban shantytowns in the Dominican Republic, was considered too crude, too vulgar, and too musically rustic to be allowed entrance into the mainstream musical landscape, since its emergence in the early 1960's. In 1991, Juan Luis Guerra and his group were responsible for transforming the genre into a success with the song *Bachata Rosa*. In his album only 4 of the 10 musics were considered as bachata because of the standard instrumentation (guitar, maracas and bongos) although the arrangements were enhanced by synthesizers and Guerra sang the romantic lyrics with the heightened emotion typical of bachata singers. Since 1990 the most significant change is bachata is the instrumentation, while still guitar led, bachata is increasingly being played with hollow-body electric guitars rather than acoustic guitars, which has dramatically changed its timbre[13].

Bolero: It has the same Afro-Cuban roots as the Rumba and is thought to have originated from Cuban or Spanish folk dances such as the Danzon and Beguine. However the Bolero is strongly associated with Mexico, because of its appropriation by the Mexican *Trios Romanticos*. Bolero poems often deal with themes of bittersweet, unrequited, betrayed or eternal love. The music is frequently arranged with Spanish vocals and a subtle percussion effect, usually implemented with guitar, conga or bongos [5].

Forró: Originated in 1946 when Luiz Gonzaga recorded the evolutionary song named *baião*, whose title become the name of a new genre. The *baião* derived from an old northeastern circle dance of African origin. It has a steady beat from beginning to end, making it easier to dance to. The basic instruments are the accordion, a triangle and a zabumba bass drum which are used to create a syncopated 2/4 rhythm. Around 1950, Luiz Gonzaga popularized others musical genres such as *Xaxado* and *Xote*. Nowadays much of his music is referred to as *Forró*, a word that originally meant a party or place to play music, however the etymology of the word is subject to much speculation [20].

Gaúcha: Is related to the southern states in Brazil and is related to its cultural identity. As with the Forró, this label is used to refer to different music genres such as the *Vaneira*, *Vaneirão*, *Buggio*, *Xote Gaúcho*, among others. The lyrics usually refer to a specific set of values like respect for the women, the love for the countryside, the culture and the animals.

Merengue: Is a century-old music has been used to tell news and gossip an to spread political propaganda since the birth and is also the national dance of the Dominican Republic. There is no mistaking its crisp.,zippy beat, hissed and scratched out on a metal grater quira in jaunty 2/4 time, or the bubbling triple beat roll of tambora rhythms and sharp twitter of interlocked saxophone vamps[21]. The mainstream Merengue is also called merengue cibaeño. There are other types of Merengue like the merengue palo eacho (or pri-prì) of the south and east. While few regional merengues are played today, a vital pri-prí culture persists. This music class for one singer, the single-headed basie drum, the guira, and the one-row accordion and its 12/8 rhythm is markedly different from the 4/4 of meregue cibaeño[4].

Pagode: It is a style of Brazilian samba, that surged in the mid-1970's, when a group of musicians associated with the Carnival Block *Cacique de Ramos* started getting together for a *pagode*, a party where people played samba. In these informal get-togethers they introduced the tan-tan, a type of atabaque, which replaced the larger and heavier surdo. This was more practical for informal samba get-togethers, as the tan-tan could be more easily carried on buses, the mode of transportation for Rio de Janeiros' Working class. Almir Guineto added a banjo, which was louder than a cavaquinho and better for open-air gatherings. Ubirany started playing a hand-held repique, called *repique de mão* and dispensed with the usual use of drum sticks. And Bira played the pandeiro in unusual ways. The sound was intimate and earth with new percussive textures. Their lyrics were unpretentious, focusing on situations from their daily life [20].

Salsa: The term salsa did not come into common usage until the early 1970s, when it appeared in Latin New York magazine and was adopted as a category for the 1975 Latin New York Music Awards [15]. There are different theories around the origins of Salsa, one of them is that it surged in Cuba and later went to Puerto Rico. However, studies from the lyrics show that Puerto Rican musical identity and salsa are inseparable, because Puerto Ricans in New York made salsa a political movement in the 1970's. The most important aspect of the instrumentation in Salsa is the Clave as every musician in the band must understand it [23].

Sertaneja: A type of Brazilian country music, surged in national popularity and was the single biggest category in term of record sales. It is a pop-music version of *música caipira*, the rural folk music from Brazil's South, Southeast and central regions. Most *Sertaneja* artists are *duplas* (duos) who strum guitars and ten-strings violas, harmonizing plaintively as they croon about romance and rural life [20].

Tango: The etymology of the word tango cannot be traced completely, but it seems high probable that it reached the Western hemisphere with the slave-ships and on the lips of slaves. Regardless of the origin, in many parts of the Spanish-American empire, the word *Tango* acquired the standard meaning of a place where African Slaves assembled for the purpose of dancing. The dance as we know today come from a parodistic way of dancing the African-Argentine *tango* where the movements of *Quebradas* (an improvised dramatic contortion) and *Cortes* (a sudden break in the standard figures of the dance) were incorporated into dances in which the partners danced together [6].

N-GRAM CHORD PROFILES FOR COMPOSER STYLE REPRESENTATION

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ABSTRACT

This paper studies the problem of using weighted N-grams of chord sequences to construct the profile of a composer. The N-gram profile of a chord sequence is the collection of all N-grams appearing in a sequence where each N-gram is given a weight proportional to its beat count. The N-gram profile of a collection of chord sequences is the simple average of the N-gram profile of all the chord sequences in the collection.

Similarity of two composers is measured by the cosine of their respective profiles, which has a value in the range $[0, 1]$. Using the cosine-based similarity, a group of composers is clustered into a hierarchy, which appears to be explicable. Also, the composition style can be identified using N-gram signatures.

1 INTRODUCTION

The chord progression is an important component in music. Musicians and listeners speak of novel and influential chord progressions, typified in the First Act Prelude of "Tristan und Isolde" by Richard Wagner, in "Because" by The Beatles, and in "Giant Steps" by John Coltrane. Among many genres of music the role of chord progressions is the most significant in Jazz, where performances are improvisational and thus performers often choose to play tunes with interesting chord progressions. While many jazz compositions are constructed based on well-known chord progressions, such as 12-bar blues progressions and "I Got Rhythm" by George Gershwin, there are composers, such as Thelonius Monk and Wayne Shorter, whose chord progressions are thought of as unique. The importance of chord progressions in Jazz raises the questions of whether they can be effectively used for music retrieval and whether they can be used to characterize composers, which we will address in this paper.

An approach to the problem of comparing two chord progressions is sequence alignment, as often used in melody-based music retrieval (see, e.g., [2, 6, 11, 12]). The basis for the sequence-alignment approach is a theory that explains transformation of a chord progression to another (see, e.g., [8, 10]). Such a generative theory offers musical understanding of how two progressions are similar, but has a limitation that not all pairs of progressions are necessarily comparable. Also, for comparing multiple progressions with each other, generative-theory-based comparisons may be too computation-intensive. This consideration suggests the use of N-grams—the patterns consisting of N-consecutive chords that appear in a chord sequence. The use of N-grams is very popular in natural language understanding (see, e.g., [5]). In music information retrieval, N-grams have been shown effective for melodic contour analysis [3,4,9]. Also, the recent work of Mauch et al. [7] use 4-grams to examine the chord sequences of The Beatles.

While the chord sequences are an important subject in musicology, we believe that they can be incorporated into a tune retrieval system where the tunes are indexed with metadata and chord sequence. In such a system, a user provides a chord sequence (either typed or copy-pasted from a sequence on screen) as input and the system retrieves tunes that contain a part with either exactly the same as (with the possibility of allowing transposition of the key) or similar to the input sequence, where the input chord progression is specified using an unambiguous notation system (such as the one in [1]).

Highly prominent in the Jazz harmony are the 6th, 7th and major 7th notes and the tensions (the 9th, the 11th, and the 13th notes). The former signify the functions that chords possess while the latter add color to triads. Chord progression analysis in terms of triads is likely to enable fundamental understanding of the chord structure, but perhaps deeper understanding can be obtained by examining these non-triad notes. Our work extends [7] by considering functional and additional notes such as the 6th, 7th, and tensions, by creating a profile out of N-gram data, and then by assigning the similarity between two profiles.

1.1 Organization of the paper

The rest of the paper is organized as follows: Section 2 describes the method for obtaining an N-gram profile of chord sequences; Section 3 describes the experiments; and Section 4 describes conclusions and discusses future work.

2 THE METHOD

2.1 N-Grams

2.1.1 N-Grams

Let U denote the universe of chords. For integers $N \geq 1$, an N-gram is an ordered N-tuple $(u_1, \ldots, u_N)$ such that $u_1, \ldots, u_N \in U$. We say that an N-gram $(u_1, \ldots, u_N)$ such that $u_1, \ldots, u_N \in U$ is *proper* if for all i, $1 \leq i \leq N-1$, $u_i \neq u_{i+1}$.

2.1.2 Chord simplification

The universe, U, of chord names is vast. There are twelve possible choices for the root (without distinguishing between sharps and flats); four for the 3rd (Minor, Major, Sus4, Omitted 3rd); four for the 5th ($^{\flat}5$, $^{\natural}5$, $^{\sharp}5$, Omitted 5th); four for the 6th/7th (the 6th, Minor 7th, Major 7th, no 6th or 7th); four for the 9th ($^{\flat}9$, $^{\natural}9$, $^{\sharp}9$, no 9th); three for the 11th ($^{\natural}11$, $^{\sharp}11$, no 11th); and three for the 13th ($^{\flat}13$, $^{\natural}13$, $^{\sharp}13$, no 13th). These make the total number of choices more than 27,000, and so, the total number of possible N-grams becomes 761 millions for $N = 2$ and 21 trillions for $N = 3$. In addition, chords can be augmented with a use of a bass note different than the root of the chord, which further increases the number of N-grams. While the space of N-grams can be enormous, the N-grams are sparse. In fact, the N-grams appearing in a progression with M chord changes is only $M - N + 1$. Even though the distributions of chords are often very skewed (towards certain keys and towards chords without tension notes), the vastness may make it unlikely for the N-gram profile of a chord progression with highly enriched chords to intersect with the N-gram profile of another chord progression. This problem can be overcome by simplifying chords.

The concept of simplification corresponds well with that of stemming in document processing, which is the process of removing modifiers of words thereby making words generated from the same root with difference modifiers treated as identical words. We divide of the process of simplifying a chord into two parts: (1) turning a fractional chord (a chord with an attached bass note, such as AMI7 on B) into a non-fractional chord and (2) simplifying the tensions and the 6th/7th. We consider three options for the first part:

- $(B0)$ simply removing the bass note (for example, AMI7 on B is turned into AMI7),
- $(B1)$ reorganizing the notes so that the bass note becomes the root (for example, AMI7 on B is turned into B7sus4 ($^{\sharp}5$, $^{\flat}9$)), and
- $(B2)$ incorporate the bass note as a tension (for example, AMI7 on B is turned into AMI9).

We consider three options for the second part:

- $(T0)$ removing entirely the tensions and the 6th/7th note,
- $(T1)$ removing entirely the tensions but keeping the 6th/7th note, and
- $(T2)$ replacing the whole tension notes with a single bit of information as to whether the chord has any tension and keeping the 6th/7th note.

We also consider the possibility of keeping all the tensions intact and keeping the 6th/7th note. We will denote this option (not a form of simplification though) by T_3. Then our simplification method is expressed as a pair (Bi, Tj) such that $0 \leq i \leq 2$ and $0 \leq j \leq 3$. The most aggressive simplifications are $(Bi, T0), 0 \leq i \leq 3$. Each of these simplifications reduces a chord to a triad and thus reduces the number of possibilities for a chord name to 192. For a progression Π and a simplification method τ, we will use $\tau(\Pi)$ to denote the progression Π after applying τ.

We will be interested only in proper N-grams. So, given an input chord progression, after simplification we collapse all the consecutive entries whose chord names are identical to each other into one entry whose duration is the total of the durations of the subsequence. After this modification every block of N-consecutive entries corresponds to an N-gram, so we call such a chord progression a *proper* chord progression. Note that simplification applied to a proper N-gram may produce a non-proper N-gram.

2.1.3 N-gram transposition

Also, since popular songs are transposed to different keys, in our analysis of N-grams we are only interested in how chords are changed after the first chord. We will thus transpose each N-gram locally, in such a way that each N-gram starts with a code having A as the root. For example, from a five-chord sequence [FMI7, B$^{\flat}$7, E$^{\flat}$MA7, CMI7, B7], we obtain three 3-grams, FMI7 – B$^{\flat}$7 – E$^{\flat}$MA7, B$^{\flat}$7 – E$^{\flat}$MA7 – CMI7, and E$^{\flat}$MA7 – CMI7 – B7, which are then transposed respectively to AMI7 – D7 – GMA7 A7 – DMA7 – BMI, and AMA7 – F$^{\sharp}$MI7 – F7. We call this process *A-transpose*.

2.1.4 Chord sequences and Weight of N-grams

We assume that each chord progression is a series of chord names each accompanied by a positive rational, which represents the number of beats during which its chord is to be played. In other words, a progression Π is a sequence $[(u_1, d_1), \ldots, (u_m, d_m)]$ such that for all i, $1 \leq i \leq m$, u_i is a chord name (belonging to the universe U of chord names) and d_i is a nonnegative rational.

We next assign a weight to each N-gram produced from a proper chord progression, for the purpose of considering the contribution that the chord progression makes to the tune.

For example, we consider that the contribution of a 4-chord pattern DMI7 - G7 - EMI7 - A7 when one beat is allocated to each chord is different from that when four beats are allocated to each. We approximate the contribution of an N-chord pattern by the total number of beats assigned to the chords. For an N-gram $(a_i, a_{i+1}, \ldots, a_{i+N-1})$ of a proper chord progression $\Pi = [(a_1, \ell_1), \ldots, (a_N, \ell_N)]$, its contribution is thus $\ell_i + \cdots + \ell_{i+N-1}$.

2.1.5 N-gram profile of chord sequences

For a chord progression Π, a simplification method τ, and a positive integer N, the N-gram profile of the progression Π with respect to τ, denoted $\Theta[\tau, N](\Pi)$, is the set of all pairs (w, c), where w is a proper N-gram appearing in Π and c is the total contribution of w (since w may appear at more than one place in Π) divided by the total contribution of all N-grams appearing in Π. By assuming that the weight is 0 for each N-gram not appearing in Π, $\Theta[\tau, N](\Pi)$ can be viewed as the set

$$\{(w, c) \mid w \text{ is an A-transposed proper } N\text{-gram appearing in } \tau(\Pi) \text{ after applying simplification } \tau \text{ and } c \text{ is the weight of } w \text{ in } \tau(\Pi)\}.$$

Given a collection C of chord sequences $\Pi_1, \ldots, \Pi_s$, the N-gram profile of C with respect to τ, $\Theta[\tau, N](C)$, is the average of $\Theta[\tau, N](\Pi_1), \ldots, \Theta[\tau, N](\Pi_s)$, that is, $\frac{1}{s}\sum_{j=1}^{s} \Theta[\tau, N](\Pi_j)$.

Figure 1 shows the melody and the chord progression of "Yesterdays" composed by Jerome Kern. Due to the space constraint on the drawing software, the minus sign is used for minor chords. With the $B0$-simplification, the two fractional chords at the end of the first line, DMI/C$^\sharp$ and DMI7/C, are respectively turned into DMI and DMI7. The chord progression has the following 3-grams after A-transpose: AMI–BMI7($^\flat$5)–E7 (16 beats), AMI7($^\flat$5)–D7–GMI (14 beats), A7–DMI–EMI7($^\flat$5) (8 beats), A7–DMI–DMI7 (8 beats), AMI–AMI7–F$^\sharp$7($^\flat$5) (8 beats), AMI7–F$^\sharp$7($^\flat$5)–B7 (12 beats), AMI7$^\flat$7–D7–G7($^\sharp$5) (12 beats), A7–D7($^\sharp$5)–G7 (12 beats), A7($^\sharp$5)–D7–G7 (12 beats), A7–D7–G7 (24 beats), A7–D7–GMA7 (12 beats), A7–DMA7–G$^\sharp$7($^\flat$5) (12 beats), AMA7–D$^\sharp$7($^\flat$5)–G$^\sharp$7 (12 beats).

2.2 Profile Comparison Using Cosine-based Similarity Measure

Given two profiles Π_1 and Π_2, we compute the similarity between them by the cosine of the two. More precisely, if $\Pi_1 = (u_1, \ldots, u_k)$ and $\Pi_2 = (v_1, \ldots, v_k)$, then the similarity between the two is:

$$\frac{u_1 v_1 + \cdots + u_k v_k}{\sqrt{u_1^2 + \cdots + u_k^2}\sqrt{v_1^2 + \cdots + v_k^2}}.$$

The cosine can be thought of as representing the similarity between the two because the value of cosine is 1 for two identical N-gram profiles and 0 for two N-gram profiles with no common N-gram patterns.

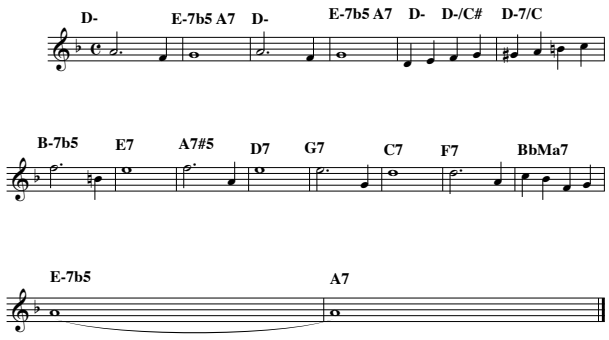

Figure 1. The melody and the chord progression of a Jerome Kern composition "Yesterdays".

3 EXPERIMENTS

3.1 Data

We collect from jazz fake books (Real Book 1, 2, and 3; New Real Book 1, 2, and 3; Jazz Limited) 218 chord progressions of the following modern jazz composers: John Coltrane (28 tunes), Chick Corea (25 tunes), Duke Ellington (25 tunes), Herbie Hancock (16 tunes), Freddie Hubbard (17 tunes), Thelonius Monk (27 tunes), Wayne Shorter (47 tunes), and Horace Silver (33 tunes). We exclude Fusion compositions of them, in particular, the Wayne Shorter compositions for the Weather Report and the later periods, the Herbie Hancock compositions for The Headhunters, and the Chick Corea compositions for The Elektric Band, fearful that addition of such tunes would make these composers too different from the rest of the composers. We also collect 63 "standard" tunes from Real Book 1. These are restricted to contain neither compositions by modern jazz musicians, nor Bossa Nova. Finally, we collect 20 compositions of The Beatles from the Hal Leonard Publishing "Anthology Volume 3". We consider these 20 compositions to be something not similar to the Jazz Fakebook tunes. An archive of the data files can be obtained at: www.cs.miami.edu/∼ogihara/chord-sequence-files.zip.

3.2 Comparison of the simplification methods

3.2.1 The choice of N and bass note simplification

To determine the value for N and and to choose the bass note simplification, we examine the cosine-based similarity between the standards and D. Ellington with respect to each of the twelve simplification methods. The similarity values are shown in Table 1. Since D. Ellington played the

most prominent role in founding the modern jazz theory and the chord progressions of the Fakebook standard tunes in some sense summarize the chord sequences resulting from Jazz reharmonization, it is anticipated that the two groups are very similar, in particular, when the tension notes are excluded (namely, $T0$ simplification). So this comparison suggests that $N = 3$ or $N = 4$ will be a good choice.

The choice of the bass note simplification (the B-part) does not seem to affect much the similarity measure, while the choice for the tension note simplification (the T-part) makes a substantial difference, in particular, for the 3- and 4-gram similarity. The phenomenon that the selection on

Method		N			
T	B	1	2	3	4
$T0$	$B0$	0.990	0.950	0.818	0.579
	$B1$	0.990	0.950	0.818	0.579
	$B2$	0.990	0.950	0.818	0.576
$T1$	$B0$	0.954	0.835	0.628	0.319
	$B1$	0.953	0.836	0.630	0.320
	$B2$	0.952	0.834	0.626	0.310
$T2$	$B0$	0.950	0.798	0.504	0.197
	$B1$	0.949	0.797	0.500	0.190
	$B2$	0.947	0.796	0.497	0.187
$T3$	$B0$	0.952	0.805	0.502	0.194
	$B1$	0.951	0.804	0.500	0.189
	$B2$	0.950	0.804	0.500	0.185

Table 1. The cosine-based similarity between the standards and D. Ellington with respect to various simplification methods.

the bass note simplification doesn't affect similarity much can be explained by the fact that only a small fraction (less than 5%) of the chords appearing the data had a bass note. This observation leads us to choose $B0$ (bass note omission) for the bass note simplification, because it is the simplest operation.

3.2.2 Tension simplification

We next examine how different the similarity value is depending on the choice for the T-part in the method. It is anticipated that the more aggressive the simplification is, the higher similarity values become, and this is clearly exhibited in Table 1 that shows comparisons between the standards and the D. Ellington tunes. According to the table, there is much difference between the $T2$ and $T3$ simplifications. Since $T2$ is more aggressive than $T3$, and thus, the resulting chord notation is generally simpler with $T2$ than with $T3$, we should choose $T2$ over $T3$.

We then compare $T0$ and $T1$ using the songs by The Beatles and those by the others. The numbers are shown in Table 2. There is a substantial difference in the similarity

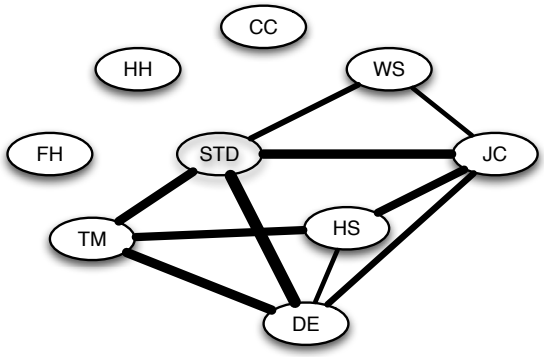

Figure 2. The similarity graph of the Jazz composers.

value between $T0$ and $T1$. Given that The Beatles is in the Pop/Rock genre and the rest are in Jazz, we feel that $T1$ is more appropriate than $T0$. Since the similarity of The Beat-

Composer	1-gram		2-gram	
	$T0$	$T1$	$T0$	$T1$
CC	0.933	0.594	0.527	0.250
DE	0.993	0.521	0.715	0.239
FH	0.921	0.570	0.456	0.114
HH	0.827	0.354	0.346	0.078
HS	0.962	0.483	0.621	0.178
JC	0.983	0.562	0.790	0.241
TM	0.998	0.551	0.691	0.243
WS	0.950	0.373	0.500	0.164

Table 2. Comparison between $T0$ and $T1$.

les to these composers seems very high for $T0$, we consider using $T1$ instead of $T0$.

These observations narrow our choices down to $(B0, T1)$ and $(B0, T2)$.

Table 3 shows the comparison of The Beatles, T. Monk, and H. Hancock with respect to the $(B0, T1)$-simplification and the $(B0, T3)$-simplification. We note that as N increases the similarity of the standards more quickly decays with The Beatles and Herbie Hancock than with Thelonius Monk and the decay is more dramatic with the 6th/7th kept in the chord names and further more dramatic with the tensions kept.

Figure 2 shows the cosine-based similarity of the profiles among the Jazz composers with respect to 3-grams and $(B0, T2)$-simplification. Two composers are connected if the similarity is 0.2500 or higher. The thicker the line is, the higher the similarity value is. The actual similarity values of these lines are summarized in Table 4. Since the similarity is symmetric, the upper right portion of the table is left blank and the two $<$'s appearing in the last line indicate that

N	Standards Versus		
	The Beatles	T. Monk	H. Hancock
1	0.430	0.922	0.875
2	0.163	0.716	0.390
3	0.040	0.437	0.114
4	0.017	0.199	0.038

N	Standards Versus		
	The Beatles	T. Monk	H. Hancock
1	0.414	0.886	0.829
2	0.162	0.676	0.185
3	0.040	0.378	0.051
4	0.018	0.1580	0.010

Table 3. Cosine-based comparison of N-gram profiles between the standards and each of The Beatles, T. Monk, and H. Hancock. for $N = 1, 2, 3, 4$. Top: with respect to the $(B0, T1)$-simplification. Bottom: : with respect to the $(B0, T3)$-simplification.

	STD	DE	HS
DE	0.504		
HS	0.349	0.376	
TM	0.379	0.422	0.363
JC	0.402	0.278	0.349
WS	0.267	$<$	$<$

Table 4. Composer Similarity

the similarity value is not more than 0.2500.

This graph coincides seems to reflect well the relations among the composers from the historical perspective. According to the year of the first recording session as a leader, these composers are ordered as follows: Ellington (1924), Monk (1947), Silver (1955), Coltrane (1957), Shorter (1959), Hubbard (1960), Hancock (1962), and Corea (1966). The graph connects among the first five along with the standards and disconnects the remaining three from every one else.

3.3 Artist clustering using profiles

This historical is more strongly represented in hierarchical clustering of the composers. Figures 3 and 4 show the hierarchical clusters of the composers generated using 3-grams, the former with the $(B0, T1)$-simplification and the latter with the $(B0, T2)$-simplification.

3.4 Unique N-gram signatures

We also consider the question of whether N-gram profiles can be used to identify a composition style. To study this

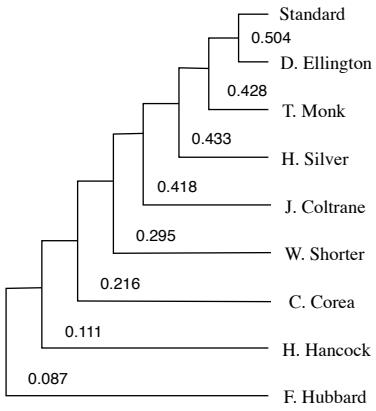

Figure 3. Hierarchical clustering of the composers with respect to the $(B0, T1)$-simplification. The Beatles is not included.

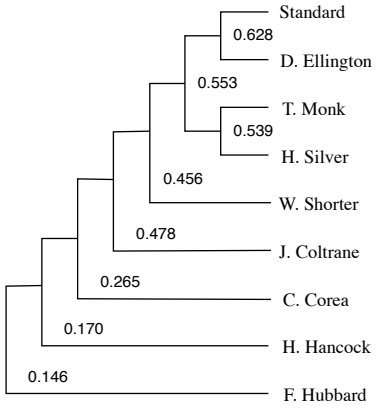

Figure 4. Hierarchical clustering of the composers with respect to the $(B0, T3)$-simplification. Again, The Beatles is not included.

question, for each composer we look for an N-gram w such that the frequency value of w with respect to that composer has a large positive gap from the frequency value of w with respect to any other composer. For each simplification method and for each $N = 1, 2, 3, 4$, we compute top 20 N-grams in terms of the narrowest positive gap from the value with respect to any other composer.

Table 5 shows the most distinctive 4-grams of the composers when compared against the rest. We note that the most prominent patterns in the standards in contrast with the rest are perfect-4th movements. This agrees with the observation of Mauch et al. [7] that the standard jazz tunes have frequent occurrences of perfect-4th movements. Also, the distinctive patterns of other composers contain chromatic, major 2nd, and minor 3rd movements.

Name	4-gram	Freq. Value	Max. Others
STD	AMI7–DMI7–G7–CMA7	0.891	0.304
	A7–DMI7–G7–CMA7	0.789	0.225
	AMI7–D7–GMA7–EMI7	0.815	0.300
	AMI7–D7–GMA7–CMA7	0.776	0.291
BTLS	A–E–A–E	4.396	0.345
	A–D–A–D	3.964	0.357
FH	AMI7–C7–B7–A♯MA7	6.250	0.000
	A7–G7–A7–G7	3.274	0.823
CC	AMI–DMI–AMI–F7	1.000	0.000
	AMI7–EMI–C–D	0.680	0.000
DE	Adim.–A♯MI7–D♯7–D♯MI7	0.799	0.000
	A7–DMI6–A7–DMI6	0.785	0.000
HH	AMI7–F7–F♯SUS7–AMI7	3.125	0.000
	A7–A♯SUS7–C♯MI7–A7	1.563	0.000
WS	A7–G7–A7–AMI7	1.084	0.000
	A7–F♯7–B–A♯MI7	1.042	0.000
HS	AMI–D7–AMI–G♯7	1.299	0.000
	AMI6–F7–AMI6–E7	1.299	0.000
JC	A7–G7–A7–CMI7	1.587	0.000
	A7–DMA7–F7–A♯MA7	1.449	0.210

Table 5. The most distinctive 4-grams of the composers with respect to the $(B0, T1)$-simplification. For each 4-gram, the "Freq. Value" column shows its frequency in the composer and the "Max. Others" column shows its maximum frequency in any other composer.

4 CONCLUSIONS AND FUTURE WORK

In this paper we explored the use of N-gram profiles generated from chord progressions to find a composition style. Of the twelve possible methods of chord simplification, we identified two to be the most promising. We used the chord profile to cluster artists in a hierarchy, which seems to be consistent with the composition styles from the historical perspective. It will be interesting to conduct more extensive studies to examine usefulness of chord progression profiles with more genres and more composers. The fact that the frequencies are very small for most of the N-grams raises the question of whether there is a more appropriate alternative for the distance measure that accounts for the sparseness. Another interesting direction will be to study the relation between the melody line and the chord progression and question whether the relation shows unique characteristics of composers.

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A WEB OF MUSICAL INFORMATION

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ABSTRACT

We describe our recent achievements in interlinking several music-related data sources on the Semantic Web. In particular, we describe interlinked datasets dealing with Creative Commons content, editorial, encyclopedic, geographic and statistical data, along with queries they can answer and tools using their data. We describe our web services, providing an on-demand access to content-based features linked with such data sources and information pertaining to their creation (including processing steps, applied algorithms, inputs, parameters or associated developers). We also provide a tool allowing such music analysis services to be set up and scripted in a simple way.

1 INTRODUCTION

Information management has become a primary concern for multimedia-related technologies, from personal collections management to the construction of large content delivery services. However, the solutions that have emerged still exist in isolation. For example, large online databases (such as Musicbrainz, MyStrands, etc.) and personal music collection management tools such as iTunes or Songbird do not interact with each other, although they actually deal with the same kind of data. The information one of these solutions manages does not benefit from the information another may hold.

The problem becomes acute when narrowing our view to the exchange of results between music technology researchers. If providing access to content-based feature extraction through web services is a first step [10, 6], the results they produce, in order for them to be meaningful, must be interlinked with other data sources. Just giving back a set of results from a particular digital audio item is useless unless we know *what* has been processed, and *how*.

In this paper, we show the benefits of using a set of web standards, often referred to as Semantic Web technologies, to achieve these goals. First, we give an overview of these standards and how they can be used to create a 'web of data'—a distributed, domain-independent, web-scale database. We give a brief summary of the Semantic Web ontology we described in [12], able to deal with music-related data. We then focus on the music-related interlinked datasets we published since, and give examples of the type of queries they can answer and of tools consuming their data. We describe our lightweight extension of this music ontology to make content-based features part of this 'web of data'. We also describe our web services, providing on-demand access to such resources, as well as the framework underlying them.

2 TOWARDS A MUSIC-RELATED WEB OF DATA

2.1 Web identifiers and structured representations

The web is built around the concept of URI (Uniform Resource Identifier [1]). A URI can identify anything, not just a document: a person, a particular performance, a signal, a particular content-based feature, etc. Web resources can have multiple associated representations. For example, to a URI identifying a particular signal, we may want to associate an HTML representation of it (providing some textual information about its characteristics) or an image depicting the actual waveform. The web aspect comes into place when other web identifiers are mentioned within such a representation. For example, the HTML representation of our signal URI might link to an URI identifying the corresponding recording device.

Now, these representations can be *structured*: they can provide explicit machine-processable information. The Resource Description Framework (RDF [2]) allows such representations to be made, by expressing *statements* about web resources in the form of *triples*: subject, predicate and object. When such representations quote other resource identifiers, enabling access to corresponding structured representation, we create a 'web of data'.

For example, starting from the URI of a band available in our Jamendo dataset [3]: `http://dbtune.org/jamendo/artist/5`, two possible representations are available. One is in HTML and can be rendered for human consumption through a traditional web browser; the other is structured and machine readable, holding explicit statements about this band. The RDF representation, requested via the HTTP `Accept` header, is

[1] http://www.ietf.org/rfc/rfc2396.txt
[2] `http://www.w3.org/RDF/`
[3] See `http://dbtune.org/` for all our DBTune datasets.

as follows: [4]

```
<http://dbtune.org/jamendo/artist/5>
    a mo:MusicGroup;
    foaf:made <http://dbtune.org/jamendo/record/174>;
    foaf:made <http://dbtune.org/jamendo/record/33>;
    owl:sameAs <http://zitgist.com/music/artist/0781a3f3-
            645c-45d1-a84f-76b4e4decf6d>;
    foaf:based_near <http://sws.geonames.org/2991627/>;
    foaf:homepage <http://www.both-world.com>;
    foaf:img <http://img.jamendo.com/artists/b/both.jpg>;
    foaf:name "Both"^^xsd:string.
```

Using this representation, we can for example follow the `foaf:based_near` link to a resource in the Geonames dataset [5] in order to get detailed information about the place where this band is based.

2.2 Accessing and querying Semantic Web data

The new SPARQL W3C recommendation [6] gives a way to access and query such data, through a simple SQL-like syntax allowing requests ranging from simple DESCRIBEs ("return all information about resource x") to complex structured queries (eg. "return the latest album from artists located in Mexico, produced by a French person").

SPARQL allows us to specify explicitly the location of the RDF data, access a particular aggregation of data, or drive a Semantic Web "user agent", such as the Semantic Web Client Library [7], which crawls and aggregates data in order to satisfy a particular query. A place on the web accepting SPARQL queries is called an *end-point*.

2.3 The Music Ontology

The Music Ontology we described in [12] aims at providing a set of web identifiers and corresponding structured representations for an ontology (defining the main concepts and relationships) of the music domain. It is divided into three different parts, respectively dealing with editorial information (track names, people, labels, releases, etc.), production workflow information (compositions, arrangements, performances, recording, etc.) and event decomposition (eg. "the piano player was playing in that particular key at that time") The following Music Ontology example describes a set of resources involved in the description of a performance of Antonio Vivaldi's Four Seasons:

```
:vivaldi a mo:MusicArtist;
    foaf:name "Antonio Vivaldi";
```

[4] All our RDF examples are written in RDF/Turtle: `http://www.w3.org/TeamSubmission/turtle/`. Each block corresponds to a set of statements (subject, predicate, object) about one subject. Web identifiers are either between angle brackets or in a prefix:name notation (with the namespaces defined at the end of the paper). Literals are enclosed in double quotes and can hold an explicit typing information.

[5] `http://geonames.org/`.

[6] `http://www.w3.org/TR/rdf-sparql-query/`

[7] `http://sites.wiwiss.fu-berlin.de/suhl/bizer/ng4j/semwebclient/`

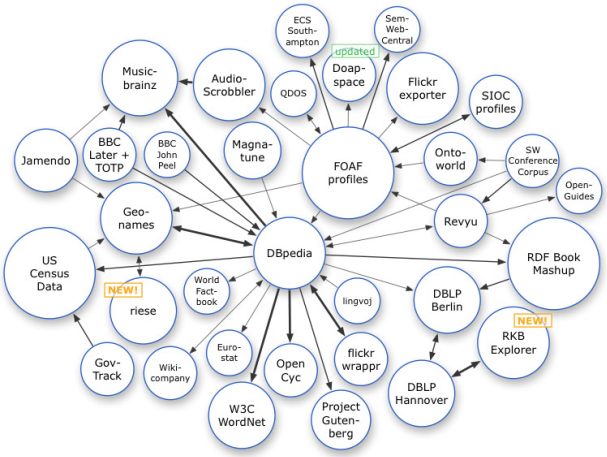

Figure 1. Available interlinked dataset, as of March 2008.

```
    owl:sameAs <http://dbpedia.org/resource/Antonio_Vivaldi>.
:compositionevent a mo:Composition;
    mo:composer :vivaldi;
    mo:produced_work :thefourseasons;
    event:time [time:year "1723"].
:thefourseasons a mo:MusicalWork;
    dc:title "The Four Seasons";
    owl:sameAs
<http://dbpedia.org/resource/The_Four_Seasons_(Vivaldi)>;
    mo:part :spring.
:spring a mo:MusicalWork;
    dc:title "Concerto No. 1 in E major, Op. 8, RV 269, La primavera";
    mo:movement :largo.
:largo a mo:Movement;
    dc:title "Largo".
:performance a mo:Performance;
    owl:sameAs <http://dbtune.org/magnatune/performance/369>;
    mo:perfomer :american_baroque;
    mo:performance_of :largo.
:americanbaroque a mo:MusicalGroup;
    foaf:name "American Baroque";
    owl:sameAs
<http://dbtune.org/magnatune/artist/american_baroque>.
```

Following the principles outlined above, this example interlinks two datasets: DBpedia and Magnatune, which we study in the next section.

3 CURRENT DATASETS

The "Linking Open Data on the Semantic Web" project [3] hosted by the W3C Semantic Web Education and Outreach group aims at publishing a wide range of datasets on the Semantic Web, and creating links between them. The currently available interlinked datasets (regardless of their domain) can be depicted as in fig. 1. In this section, we discuss a few of these datasets and their links to other datasets. We especially focus on the music-related datasets we published within our DBTune service, using the Music Ontology mentioned above. We illustrate their use by examples of queries they can answer, or by tools using their data.

3.1 DBpedia

The DBpedia project described by Auer et al. in [1] extracts structured information from Wikipedia, and republishes it on the Semantic Web, together with links to other datasets. The whole DBpedia dataset holds around 100 million RDF statements, describing 2.4 million entities across a wide range of domains, and 2 million links (statements that quote an identifier outside DBpedia). For example, the following SPARQL query issued on the DBpedia dataset returns names, descriptions and birth dates of guitarists born in Paris between 1939 and 1945: [8]

```
SELECT ?name ?birth ?description ?person WHERE {
?person
  p:birthPlace <http://dbpedia.org/resource/Paris> ;
  a yago:Guitarist110151760 ;
  p:birth ?birth ;
  foaf:name ?name ;
  rdfs:comment ?description .
  FILTER (LANG(?description) = 'en' &&
   ?birth > "1939-01-01"^^xsd:date &&
   ?birth < "1945-01-01"^^xsd:date ) . }
```

The DBpedia relationship finder [9] allows this dataset to be queried for arbitrary RDF 'paths' between two different resources. For example, it would answer the query 'what do Mozart and Metallica have in common?' as depicted in fig. 2.

3.2 DBTune

We created the DBTune service to experiment with heterogeneous music-related datasets published as Music Ontology linked data. We published the Magnatune record label catalogue, and we interlinked it with corresponding DBpedia identifiers (as illustrated in § 2.3). We also published the Jamendo music platform database, and we interlinked it with Geonames and Musicbrainz identifiers. DBTune republishes on the Semantic Web data coming from AudioScrobbler and MySpace, therefore allowing different social networks as well as listening habits to be interlinked. DBTune also hosts a Semantic Web version of the BBC John Peel sessions, interlinked with DBpedia.

Musicbrainz is a community-maintained music database holding detailed editorial information about 300,000 artists, 500,000 releases and 6 million tracks. We published a linked data version of this dataset. Artists, records and countries are interlinked with corresponding DBpedia web identifiers. For example, the identifier corresponding to Madonna in DBTune is stated as being the same as the corresponding identifier in DBpedia. We also interlinked artist resources with corresponding web identifiers on our linked data publication of the MySpace social network.

We provide a number of SPARQL end-points, for each dataset (except AudioScrobbler and MySpace, as the RDF documents are generated dynamically). For example, the following query can be issued to our Jamendo end-point:

```
SELECT DISTINCT ?an ?lat ?long WHERE {
?a a mo:MusicArtist;
  foaf:based_near ?place;
  foaf:name ?an;
  foaf:made ?alb.
?alb tags:taggedWithTag
     <http://dbtune.org/jamendo/tag/jazz>.
?place
  geo:name ?name;
  geo:population ?population;
  wgs:lat ?lat;
  wgs:long ?long } ORDER BY ?population
```

This returns artists who made at least one album tagged as 'jazz' by a Jamendo user, sorted by the number of inhabitants of the places they are based near. Overall, DBTune gives access to approximately 13 billion RDF triples, thus making it one of the largest dataset on the Semantic Web.

3.3 Use-case: personal collection management

Such interlinked music-related data sources can be used for a wide range of use-cases, including personal collection management. We developed two tools to aggregate Semantic Web data describing arbitrary personal music collections. GNAT [9] finds, for all tracks available in a collection, the corresponding web identifiers in the Semantic Web publication of the Musicbrainz dataset mentioned earlier. GNAT uses primarily a metadata-based interlinking algorithm described in [13], which was also used to interlink the above described music-related datasets. GNARQL crawls the web of data from these identifiers and aggregate structured information about them, coming from heterogeneous data sources. GNAT and GNARQL then automatically create a tailored database, describing different aspects of a personal music collection. GNARQL provides a SPARQL endpoint, allowing this aggregation of Semantic Web data to be queried. For example, queries such as "Create a playlist of performances of works by French composers, written between 1800 and 1850" or "Sort European hip-hop artists in my collection by murder rates in their city" can be answered using this end-point. GNARQL also gives a faceted browsing interface based on /facet [8], as illustrated in fig. 3.

4 ON-DEMAND FEATURE EXTRACTION AND INTERLINKING

4.1 Why web services are not enough

Although providing web service access through technologies such as SOAP or WSDL is a first step towards a distributed framework for performing music analysis tasks (as highlighted in [10] and [6]), this is not enough. In order for results to be easily re-usable, it is desirable to prevent

[8] This query returns, among others, Babik Reinhardt, one of Django Reinhardt's sons.

[9] GNAT and GNARQL are available at `http://sourceforge.net/projects/motools`.

Figure 2. What do Mozart and Metallica have in common?

Figure 3. Management of personal music collections using GNAT and GNARQL. Here, we plot our collection on a map and display a particular artist.

the client software from having to explicitly adapt to the peculiarities of each web service. This is particularly important when trying to combine several web services (known as the 'choreography' problem). The results of various services must be interlinked with additional resources supplying information pertaining to their creation. The information could include processing steps, applied algorithms, inputs and the parameters, associated developers, etc.

The technologies described in § 2 provide a way to address this issue. The results of a music analysis task can be described in an RDF document which provides such links. The results are then part of the data web, and can be queried through their relationships to further data. For example, one could issue a SPARQL query such as "Give me a structural segmentation obtained using an implementation of algorithm x developed by y, on the signal identified by z".

In the rest of this section, we describe our framework to express music analysis results in RDF, and a tool providing on-demand access to such interlinked results. We also describe a way to easily script and publish combinations of music analysis results made available on the Semantic Web.

4.2 Representing features in RDF

We extended the Music Ontology framework to describe a wide range of musical audio features in RDF [10]. Features describing the whole audio signal (whose concept is identified by the URI `mo:Signal`) can be directly attached to the signal resource. Moreover, an audio signal is associated with a temporal extent (`tl:Interval`) defined on a time-line (`tl:TimeLine`). Then, events (`event:Event`) can be defined in order to classify particular points or intervals on such time-lines. Defining new types of audio features is then just a matter of subsuming the event concept. For example, the following defines a new zero-crossing rate feature type:

```
:ZeroCrossingRate
      a owl:Class;
      rdfs:subClassOf event:Event;
      rdfs:label "Zero Crossing Rate event".
:zcr
      a owl:DatatypeProperty;
      rdfs:subPropertyOf event:literal_factor;
      rdfs:domain :ZeroCrossingRate;
      rdfs:label "Associated zero-crossing rate".
```

Then, an instance of such a feature on the time-line of a particular signal (here, a track from the Jamendo dataset) would be represented as:

```
<http://dbtune.org/audio/Den-Nostalia.ogg> mo:encodes :sig.
:sig a mo:Signal;
      mo:time       [ # Temporal extent of the signal
        tl:timeline :tl; # Time-line on which this extent is defined
        tl:duration "PT4M50S" ]. # 4 minutes and 50 seconds
:e1 a :ZeroCrossingRate;
      :zcr "27"; # Zero-crossing rate
      event:time [
        tl:start "PT0S"; # Starts at 0
        tl:end "PT0.046439912S"; # Ends at 1024/22050 seconds
        tl:timeline :tl ]. # Defined on the time-line backing our signal
```

4.3 Evaluating music analysis predicates

We model music analysis processes as RDF properties, associating their results with the inputs and parameters used. Predicates involving these properties can be considered as *built-in* predicates, within particular SPARQL end-points. For example, the decoding of an audio file and the evaluation of a musical key detection Vamp plugin [11] implementing the template-based algorithm described by Noland and Sandler in [11] could be triggered while evaluating the following query:

```
SELECT ?sig ?result
WHERE {
<http://dbtune.org/audio/Den-Nostalia.ogg>
   mo:encodes ?sig.
?sig vamp:qm-keydetector ?result }
```

The audio file is discovered at querying time. The corresponding signal resource is associated to the `?sig` variable,

[10] `http://purl.org/ontology/af/`

[11] `http://www.vamp-plugins.org/`.

and the results given by the Vamp plugin are associated to the `?result` variable. The predicates `mo:encodes` and `vamp:qm-keydetector` both correspond to a particular process, which is triggered when the predicate is evaluated.

Moreover, in the representations of `mo:encodes` and `vamp:qm-keydetector`, we can access two statements specifying that these are *functional* properties [12]. For one subject of these properties, there is only one corresponding object. Results corresponding to the evaluation of such predicates are therefore cached, to prevent repeating similar evaluations.

In these representations, we might also have links to actual implementations. We can therefore discover at querying time how to evaluate a new predicate by retrieving such implementations.

4.4 Combining music analysis and structured web data

N3 [2] provides a way to publish rules over RDF data on the Semantic Web, by extending the RDF data model in two ways: *quoting* facilities (the ability to consider an RDF graph as a literal resource) and universal quantification. A particular RDF predicate then captures the notion of implication: `log:implies` (denoted by the => symbol in our example). N3, in our context, allows to easily script how combination of predicates (which resolution might trigger some analysis tools, or some querying of RDF data) can lead to further predicates. A N3 rule can express "statements linking two similar audio signals can be derived by computing a MFCC model for both and thresholding their Kullback-Leibler divergence", or "statements linking two similar audio signals can be derived by thresholding the cosine distance of the corresponding beat spectra" [13]. A N3 rule can also merge contextual queries and content analysis, as suggested in [7]: "A *danceable* track has an average tempo of around 120 beats per minute, a high loudness, and is in the playlist of a disc jockey.".

Our previous key detection example can be mashed up with an identification process (captured by the `gnat:match` predicate which, from an audio file, finds out the corresponding identifier in the Musicbrainz dataset) and be translated into Music Ontology terms using the following rule:

```
{ ?af mo:encodes ?sig; gnat:match ?mbzsig. ?sig vamp:qm-keydetector ?results.
?mbzsig mo:time [tl:timeline _:tl ].
(?start ?duration ?key) list:in ?results. } => {
    _:keyevent a to:Tonality;
      rdfs:label "key event"; to:key ?key; event:time [
        tl:start ?start; tl:duration ?duration; tl:timeline _:tl ]. }
```

[12] individuals of `owl:FunctionalProperty`

[13] Using the approach described in [5], we can trace back what has been achieved to derive such similarity statements.

4.5 On-demand extraction and interlinking: Henry

We developed an interpreter of such rules combining music analysis processes and contextual data: Henry [14]. Henry handles two persistent stores: an RDF and N3 store m (handling, among other things, computation metadata and RDF statements gleaned from the web) and a binary store b (handling binary representations of resources described in the RDF store: signals, features, etc.). The underlying logic of this software builds on top of Transaction Logic [4] over b. Henry provides the ability to register new built-in predicates written in a variety of languages.The main interface to interact with Henry is its SPARQL end-point. When processing a SPARQL query q, Henry executes the following steps:

1. For every newly appearing web identifier i in q, dereference it, and then:
   (a) If the representation is RDF, store it in m. If i is a property and its representation links to a built-in implementation matching the current platform, retrieve it ;
   (b) If the representation is N3, store it in m and register the corresponding rules ;
   (c) If the representation is binary, cache it in b ;
2. For every triple pattern p in q, the possible solutions are:
   (a) Instantiations of p in m ;
   (b) If p is in the conclusion part of a rule, the solutions correspond to the possible evaluations of the premises ;
   (c) if p=(s_p, p_p, o_p) where p_p is a built-in predicate, solutions are derived using this built-in, updating b accordingly.

4.6 Available services

An available Henry instance [15] provides access to a number of such results. It wraps a Vamp plug-in host, so new built-in predicates and the corresponding Vamp implementation can be discovered at querying time. This instance also discovers new audio files at querying time. It provides a SPARQL endpoint, although a future version will also include a simplified HTTP interface.

The above defined rule for key events and a number of others are pre-loaded. For example, the following query selects several attributes of key events on the time-line of an audio signal (start time, end time and corresponding key):

[14] Henry source code is available at `http://code.google.com/p/km-rdf/`

[15] Available at `http://dbtune.org/henry/`

```
SELECT ?start ?end ?key WHERE {
<http://dbtune.org/audio/Den-Nostalia.ogg>
     mo:encodes ?sig.
?sig mo:time ?time.
?time tl:timeline ?tl.
_:evt a to:Tonality;
     event:time [
        tl:start ?start; tl:end ?end; tl:timeline ?tl
        ] ;
     to:key ?key }
```

This query triggers a call to a key detection Vamp plugin if the results have not yet previously been computed for the mentioned audio file. As the predicate corresponding to the Vamp call is functional, the Vamp results are cached. The results bound with the variable $?key$ are resources within the Tonality Ontology [16], which allows us to access further information about the keys themselves.

5 CONCLUSION AND FUTURE WORK

Publishing web resources and corresponding RDF representations providing links to further resources is a big step towards a shared information space for music enthusiasts. The web environment allows us to link together heterogeneous types of information: from music analysis results to editorial or social data, but also data that we often don't include in a music information system, such as geographical, statistical or encyclopedic data. This web of data is now a reality, which is concretely used in a number of applications such as GNAT and GNARQL. We developed a tool called Henry, which dynamically creates and publishes such linked data, by working on top of on-the-fly discovery of content, content analysis processes, logical rules, and external data. Publishing a new range of music analysis results on the data web is therefore as simple as publishing a small amount of RDF specifying new processing tools (if we need any), and some N3 rules combining them together. We are now in the position of answering hybrid queries such as 'find me all works composed in New York in 1977, performed at an average tempo of 130 bpm and with an overall C major key'.

Future work includes publishing and interlinking more music-related datasets. It also includes delegating the resolution of predicates within Henry: it might be inefficient to allow every instance to be dynamically patched with the same set of plugins, handling a similar range of processing. Moreover, some plugin providers may not want to distribute their implementation, but just provide web access to it.

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[16] http://purl.org/ontology/tonality/

Namespaces

```
@prefix rdfs: <http://www.w3.org/2000/01/rdf-schema#>.
@prefix owl: <http://www.w3.org/2002/07/owl#>.
@prefix geo: <http://www.geonames.org/ontology#>.
@prefix wgs: <http://www.w3.org/2003/01/geo/wgs84_pos#>.
@prefix yago: <http://dbpedia.org/class/yago/>.
@prefix foaf: <http://xmlns.com/foaf/0.1/>.
@prefix p: <http://dbpedia.org/property/>.
@prefix xsd: <http://www.w3.org/2001/XMLSchema#>.
@prefix dc: <http://purl.org/dc/elements/1.1/>.
@prefix mo: <http://purl.org/ontology/mo/>.
@prefix event: <http://purl.org/NET/c4dm/event.owl#>.
@prefix tl: <http://purl.org/NET/c4dm/timeline.owl#>.
@prefix to: <http://purl.org/ontology/tonality/>.
@prefix vamp: <http://purl.org/ontology/vamp/>.
@prefix list: <http://www.w3.org/2000/10/swap/list#>.
@prefix gnat: <http://motools.sourceforge.net/doap.rdf#>.
@prefix tags: <http://www.holygoat.co.uk/owl/redwood/0.1/tags/>.
```

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HYPERLINKING LYRICS: A METHOD FOR CREATING HYPERLINKS BETWEEN PHRASES IN SONG LYRICS

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ABSTRACT

We describe a novel method for creating a hyperlink from a phrase in the lyrics of a song to the same phrase in the lyrics of another song. This method can be applied to various applications, such as song clustering based on the meaning of the lyrics and a music playback interface that will enable a user to browse and discover songs on the basis of lyrics. Given a song database consisting of songs with their text lyrics and songs without their text lyrics, our method first extracts appropriate keywords (phrases) from the text lyrics without using audio signals. It then finds these keywords in audio signals by estimating the keywords' start and end times. Although the performance obtained in our experiments has room for improvement, the potential of this new approach is shown.

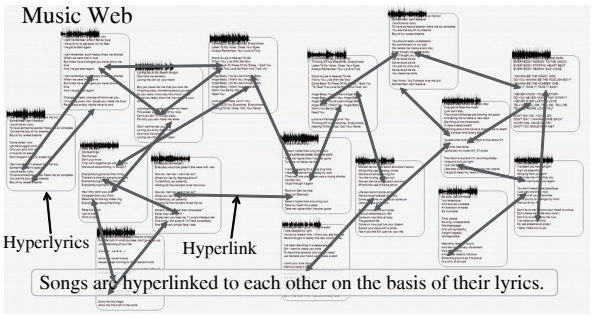

Figure 1. Hyperlinking lyrics. on Music Web

1 INTRODUCTION

The goal of this study is to enable a *Music Web* where songs are hyperlinked to each other on the basis of their lyrics (figure 1). Just as some hypertext phrases on the web are hyperlinked, so some phrases of lyrics (which we call *hyperlyrics*) on the Music Web can be hyperlinked. Such a hyperlinked structure of the Music Web can be used as a basis for various applications. For example, we can cluster songs based on the meanings of their lyrics by analyzing the hyperlinked structure or can show relevant information during music playback by analyzing the hyperlinked songs. We can also provide a new active music listening interface [1] where a user can browse and discover songs by clicking a hyperlinked phrase in the lyrics of a song to jump to the same phrase in the lyrics of another song. Although we can think of many possible ways to hyperlink musical pieces on the Music Web, focusing on song lyrics is natural because the lyrics are one of the most important elements of songs and often convey their essential messages.

Most approaches for analyzing inter-song relationship have been based on musical similarity between songs, and various music interfaces based on such song-level similarity have been proposed [2, 3, 4, 5, 6]. the methods of music browsing for intra-song navigation have also been studied, such as music browsing based on the structure [7, 8] and the lyrics [9, 10, 11]. However, hyperlinking lyrics — i.e., a combination of inter-song and intra-song navigations based on lyrics phrases — has not been proposed.

In this paper, we propose a method for hyperlinking identical keywords (phrases) that appear in the lyrics of different songs. Hyperlinking lyrics enables us to benefit from various studies dealing with sung lyrics in musical audio signals. For example, by using methods for automatic synchronization of lyrics with musical audio signals[9, 10], we can first find a keyword pair in the text lyrics for two different songs and then locate (i.e., estimate the start and end times of) the keyword in the audio signals of each song. However, it is still difficult to achieve accurate automatic lyrics recognition that enables us to find a keyword pair in sung lyrics that are recognized in polyphonic musical audio signals, despite the achievements made by studies on automatic lyrics recognition for musical audio signals [12, 13, 14]. While studies have also been done on analyzing text lyrics without using audio signals [15, 16], we cannot use their results to hyperlink lyrics.

Figure 2 shows the strategy our method uses to hyperlink lyrics. We assume that a user has a song database that is a set of audio files (e.g., MP3 files) and that we can prepare text files of the lyrics for some of the songs. Note that we do not require text lyrics for all the songs — i.e., a song database consists of songs with their text lyrics and songs without them. We therefore apply two different strategies for hyperlinking: one for hyperlinking from a phrase in other text lyrics to the same phrase in the text lyrics, and one for hyperlinking from a phrase in the text lyrics to the same phrase in the sung lyrics in a song (polyphonic music signals) without its text lyrics. Here, "text lyrics" means a text document containing the lyrics of a song, and "sung lyrics" means audio signals containing the lyrics sung by a singer in a polyphonic sound mixture. Although hyperlinking from text lyrics to text lyrics is relatively easy with the support of LyricSynchronizer [10, 1], hyperlinking from text lyrics to sung lyrics is more difficult.

2 OUR METHOD FOR HYPERLINKING LYRICS

Our method hyperlinks phrases that appear in the lyrics of different songs. In other words, if different songs share

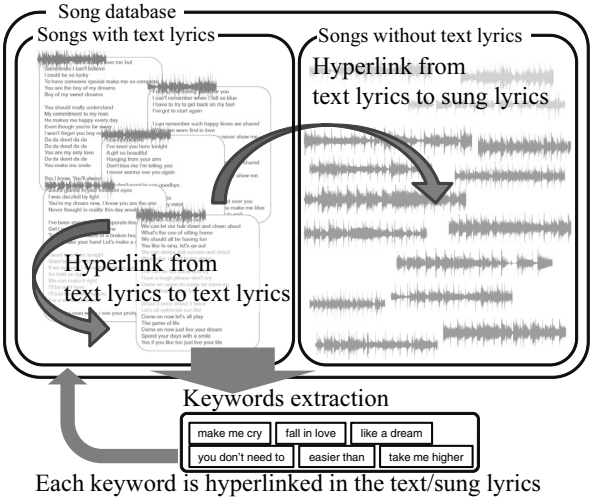

Figure 2. Two different strategies for hyperlinking: hyperlink from text lyrics to text lyrics and hyperlink from text lyircs to sung lyrics.

the same phrase (what we call a *keyword*[1]) in their lyrics, a section (temporal region) corresponding to the keyword in one song is hyperlinked with a section corresponding to the same keyword in another song. This method should deal with polyphonic music mixtures containing sounds of various instruments as well as singing voices. If a song database consists of songs with text lyrics and songs without text lyrics, this approach creates two different types of bidirectional hyperlinks: a hyperlink between songs with text lyrics and a hyperlink from a song with text lyrics to a song without them. The former hyperlink can be created by extracting potential keywords from all the text lyrics and finding sections corresponding to them (i.e., temporally locating them) in audio signals with the help of their lyrics. This estimation can be done using an automatic lyrics synchronization method described in [10]. For the latter hyperlink, our method looks for sections including voices that sing the keywords by using a keyword spotting technique for polyphonic music mixtures.

2.1 Hyperlinking from text lyrics to text lyrics

This section describes our method for hyperlinking from text lyrics to text lyrics. By using text lyrics of all the songs, the method first extracts as many keywords that can result in meaningful hyperlinks as possible. It then estimates sections (the start and end times) of each keyword in audio signals. Finally, it creates bidirectional hyperlinks between the estimated keyword sections.

2.1.1 Keyword extraction from the text lyrics

We would like to create as many hyperlinks as possible while ensuring that they are meaningful and useful. We therefore have to accordingly extract keywords. From the viewpoint of inter-song hyperlinks, each keyword must appear in multiple songs (the more songs, the better). In addition, since long keywords tend to convey important meanings, longer keyword lengths are preferred. Longer keywords are also advantageous for improving the accuracy of keyword detection, in Sec. 2.2. In contrast, short words/phrases, such as an article and a preposition, are not appropriate as keywords.

The requirements for appropriate keywords can be summarized as follows:

(a) A larger number of songs sharing a keyword is better.
(b) A larger number of phonemes in a keyword is better.

Note the trade-off between these requirements: a longer phrase is less likely to appear many times. We therefore try to maximize the number of phonemes provided each keyword appears in two songs or more.

According to these requirements, we developed the keyword extraction algorithm shown below.

1. Initialize a keyword dictionary as empty. Each keyword registered to this dictionary in the following will have a flag called a "finalized" flag only when it cannot be connected with an adjacent word to make a longer keyword (phrase).
2. Register every word of all text lyrics to the keyword dictionary.
3. Count the number of different songs whose lyrics include each keyword of the keyword dictionary. The keyword with the largest number of corresponding songs is considered the most frequent keyword.
4. Remove all keywords that appear in less than M songs from the keyword dictionary.
5. Select the most frequently occurring keyword without the "finalized" flag. If all the keywords have this flag, this algorithm is finished.
6. Try to combine adjacent words with the selected keyword. The former word and the latter word of each lyrics are respectively combined to make a longer phrase.
7. Only the best combination that results in the most frequent phrase is registered as a new keyword to the keyword dictionary, provided it appears in M songs or more. Note that the original keyword is not removed even if a combined one is thus registered.
8. If any combination for the selected keyword does not appear in M songs or more, the "finalized" flag is attached to the keyword when the number of phoneme of it is more than "N" and the keyword is removed when the number of phoneme of it is under "N".
9. Go back to 5.

First, the algorithm uses a dictionary to prepare the pronunciation of each word of all text lyrics so that each word can be represented as a sequence of phonemes and the number of phonemes can be calculated. [2]

Two parameters, M and N, correspond to the above requirements. Since their appropriate values will depend on the total volume of all lyrics, in Sec. 2.2 we discuss how they are set.

[1] In this paper, the term "keyword" means a lyrics phrase consisting of one or more consecutive words.

[2] For songs sung in the Japanese language, we need to apply a preprocessing, called morphological analysis, so that text lyrics can be divided into a sequence of words. Unlike English, a word boundary is not explicitly specified by a space character in Japanese.

2.1.2 Hyperlinking keywords

Each keyword in the keyword dictionary is used to create bidirectional hyperlinks. Given the text lyrics, our method can easily find all the positions of each keyword through a string search. At each found position, the method then finds the keyword's section (temporal region) by estimating its start and end times in the corresponding musical audio signals. This can be done by using an automatic lyrics synchronization method [10] which estimates the start and end times of all words in the lyrics, including the keywords.

2.2 Hyperlinking from text lyrics to sung lyrics

This section describes our method for hyperlinking from songs with text lyrics to songs with only the sung lyrics (audio signals). Each keyword in the keyword dictionary is again used to create bidirectional hyperlinks, but our method should find each keyword in polyphonic musical audio signals without text lyrics. The method first judges whether the sung lyrics of each song (without text lyrics) includes a keyword, and then finds the keyword section by estimating its start and end times. We enable this through a method based on a keyword spotting technique [17]. This technique uses two different statistical acoustic models: a model for the keyword pronunciation (called the *keyword model*) and a model for other sounds (called the *garbage model*). By finding the best matching of these models to singing voices segregated from polyphonic audio signals, we can find keyword sections described by the keyword model. More specifically, we first detect many candidate sections of keywords and then narrow the candidates down by rescoring them.

2.2.1 Feature extraction for singing voices in polyphonic mixtures

To use the acoustic models, we extract a sequence of feature vectors from each song without the text lyrics. Since the audio signals are polyphonic, we have to reduce various influences of the accompaniment sounds to extract feature vectors that represents only the singing voice. We therefore use the feature extraction method described in [10]. This method estimates and resynthesizes the singing voice (the sung melody) in polyphonic mixtures through three steps:

(1) estimate the most predominant F0 as the melody line (singing voice candidate) by using the *PreFEst* method [18]

(2) extract the harmonic structure corresponding to the estimated F0

(3) use sinusoidal synthesis of the harmonics to resynthesize the audio signal (waveform) corresponding to the melody line.

After resynthesizing the singing voice, we extract the MFCCs, ΔMFCCs, and Δ Power as feature vectors.

2.2.2 Preparing acoustic models by training phone models

The keyword and garbage models are represented as the hidden Markov models (HMMs). In the keyword HMMs, the pronunciation of each keyword is represented as a sequence of phone models as shown in Figure 3. The garbage HMM is defined as a phoneme typewriter in which any phonemes can appear in any order as shown in Figure 4. The keyword HMMs and garbage HMM are integrated in parallel as shown in Figure 5.

The phone models that represent acoustic characteristics of the phonemes of singing voices significantly affect the performance of keyword spotting. Because we cannot directly use phone models for typical automatic speech recognition, we built from scratch our own phone models for singing voices. We prepared precise phoneme-level annotation for 79 Japanese songs of the RWC Music Database: Popular Music (RWC-MDB-P-2001) [19], and then trained monophone models with 3-state left-to-right HMMs on those songs.

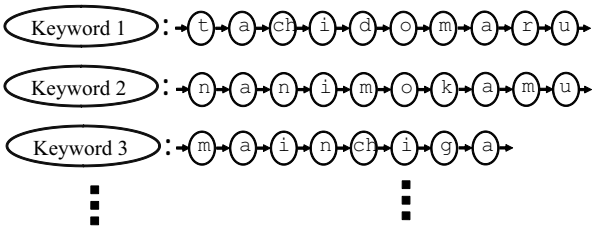

Figure 3. Keyword models (HMMs).

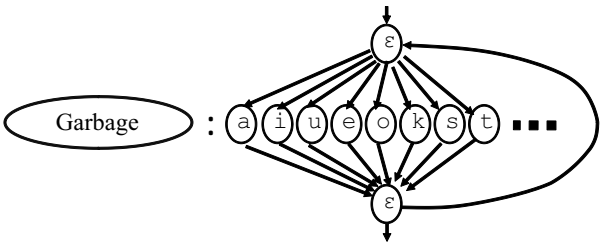

Figure 4. Garbage model (HMM).

2.2.3 Keyword candidate detection and score calculation

For each song without the text lyrics, the method detects candidate sections of each keyword by applying a Viterbi decoder to the feature vectors. As a result, the keyword HMMs can be expected to match with candidate sections where the singer sings the keyword, while the garbage HMM can be expected to match with all other sections. In this decoding, the likelihood of each candidate section against the corresponding keyword HMM can also be obtained, which represents acoustic matching between singing voices and the HMM. This decoding needs to use a word insertion penalty; each keyword in the keyword HMMs and each phoneme in the garbage HMM is regarded as a word. The word insertion penalty prevents long keywords from being substituted by short keywords of other keyword HMMs or phonemes of the garbage HMM. This decoding framework is based on typical automatic speech recognition techniques, except that the language model (Figure 5) was designed specifically to detect the keywords.

After the candidate sections of keywords are detected, we calculate the score of each candidate to narrow down the number of candidates. First, the Viterbi decoder using only the garbage HMM is applied to each candidate section to obtain its likelihood against the garbage HMM. Since the likelihood of each candidate section against the corresponding keyword HMM has already been obtained as explained above the score can be defined as the difference in the average log likelihood between the keyword HMM and the

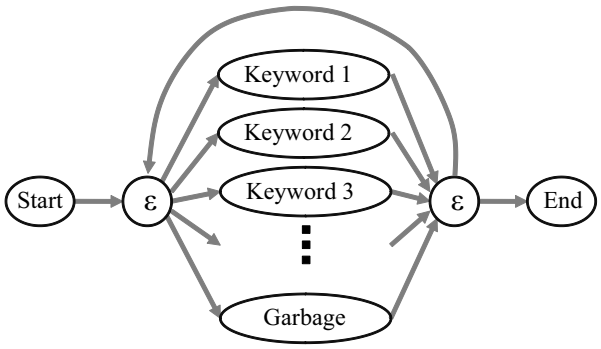

Figure 5. Integration of the keyword models and the garbage model.

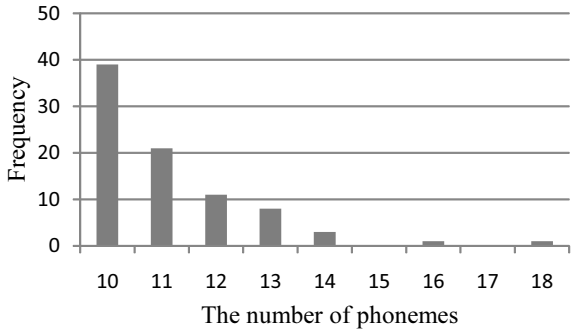

Figure 6. Distribution of the number of phonemes of extracted keywords.

garbage HMM.

2.2.4 Hyperlinking keywords

For each keyword, we select several songs that include detected keyword candidate sections with high scores. We then create bidirectional hyperlinks from keyword sections in songs with the text lyrics to the selected candidate sections in songs without the text lyrics.

3 EXPERIMENTS

We conducted preliminary experiments to evaluate our hyperlinking method by using 79 Japanese songs taken from the RWC Music Database: Popular Music (RWC-MDB-P-2001) [19].

3.1 Evaluation of hyperlinking lyrics

First, we evaluated the performance of the keyword extraction described in Sec. 2.1.1. We set parameters M and N to 2 (songs) and 10 (phonemes), respectively. As a result, 84 keywords were automatically extracted and the average number of phonemes of those keywords was 11.1. [3] Figure 6 shows a distribution of the number of phonemes and Table 1 shows examples of the extracted keywords. We found that appropriate phrases that could result in meaningful hyperlinks were chosen as the keywords.

Next, we evaluated the performance of hyperlinking from text lyrics to sung lyrics, as described in Sec. 2.2 by introducing a new evaluation measure *link success rate* [4]. The link success rate indicates the percentage of hyperlinks that correctly connected a phrase in a song to the same phrase appearing in another song. Our evaluation procedures were as follow:

1. Assuming that the lyrics of the 79 songs were unknown, we executed the keyword candidate detection and score calculation described in Sec. 2.2.3.
2. For each song in the 79 songs, we again assumed that we obtained the text lyrics of this song only and created the hyperlinks of the above 84 keywords from this song to the other 78 songs. In this experiment, the hyperlinks were created only for the keyword candidate section with the highest score.
3. We evaluated these hyperlinks by using the link success rate and averaged the rate over all 79 songs.

The link success rate, r, of a song is expressed as

$$r = \frac{\sum_{k=1}^{K} \sum_{i=1}^{I_k} s(w(k,i))}{\sum_{k=1}^{K} I_k}, \quad (1)$$

$$s(w(k,i)) = \begin{cases} 1 & \text{if } w(k,i) \text{ is appropriate} \\ 0 & \text{if } w(k,i) \text{ is not appropriate} \end{cases}, \quad (2)$$

where k denotes a different keyword, K denotes the number of keywords appearing in the song, I_k denotes the number of occurences of the k-th keyword within this song (note that the same keyword is sometimes repeated in the same song), and $w(k,i)$ expresses the i-th hyperlink of the k-th keyword. A hyperlink $w(k,i)$ is judged to be appropriate when more than half of the detected region (in the other 78 songs) hyperlinked from the k-th keyword overlaps with the ground truth region (given by the annotation we used in Sec. 2.2.2. The experimental result showed that the link success rate was 30.1%.

3.2 Number of phonemes and occurrences of keywords

We then evaluated the appropriateness of parameters M (= 2 songs) and N (= 10 phonemes). As described in Sec. 2.1.1, M indicates the minimum number of songs in which each keyword should appear and N indicates the minimum number of phonemes of each keyword (i.e., the minimum length of each keyword). Because we would like to create as many hyperlinks as possible, we first fixed M as 2 songs for this evaluation with a small number of songs. We then measured the link success rate by increasing N. Table 2 shows the dependence of the link success rate and the number of extracted keywords on the number of phonemes in the keywords. Taking the trade-off between the two parameters into consideration, we set N as 10 phonemes.

3.3 Validation of our experimental conditions

In the experiments described in Secs. 3.1 and 3.2, the 79 songs used for the evaluation of hyperlinks were the

[3] Since the lyrics used here were written in Japanese, we used a preprocessing morphological analyzer MeCab[20] to divide the input text lyrics into word sequences.

[4] In this paper, we have omitted any evaluation of the hyperlinking from text lyrics to text lyrics, described in Sec. 2.1, because its accuracy depends on only the performance of LyricSynchronizer [10, 1]

Table 1. Examples of extracted keywords. Note that English translations were done specifically are prepared just for this paper because most keywords are not complete sentences and are hard to translate.

Keyword	English translation	Phoneme sequence	number of occurrences
が教えてくれたこと	what 〜 taught me	g a o sh i e t e k u r e t a k o t o	2 songs
どこまでも続く	continue forever	d o k o m a d e m o ts u z u k u	2 songs
心の中	in one's heart	k o k o r o n o n a k a	3 songs
素敵な笑顔	nice smile	s u t e k i n a e g a o	2 songs
世界中に	all over the world	s e k a i jy u: n i	2 songs

Table 2. Dependence of the link success rate on the number of phonemes.

# of phonemes	8	9	10	11	12	13
Link success rate (%)	23.2	27.5	30.1	24.3	35.9	40.0
# of keywords	271	144	84	45	24	13

Table 3. Results of phoneme-level recognition: Comparison between closed and open conditions.

Condition	i. closed	ii. open
Accuracy	50.9%	49.8%

same as those used for the phone model training. This was done because we had to apply the laborious time-consuming phoneme-level annotation on the 79 songs for both training the phone model and preparing the ground truth for the evaluation. Moreover, since the likelihood comparison between different songs requires use of the same phone model, we could not prepare different phone models by omitting the target song in the training and conduct a typical cross-validation. Since we used 79 songs for the training, we expected little contribution from the target song (used in the evaluation). Still, we wanted to confirm that this would not be a major issue affecting the reliability of our evaluation.

We therefore evaluated the performance of phoneme-level recognition using the Viterbi decoder and the phoneme typewriter under the condition that the text lyrics were not known. As the ground truth data for measuring the average frame level accuracy, we used the phoneme-level annotation used to train the phone model. Two two conditions were compared:

i. The same 79 songs were used for both the phone model training and the evaluation (closed condition).

ii. 10-fold cross validation was done on the 79 songs for the training and the evaluation (open condition).

Note that these conditions, the purpose, and the evaluation measure differ from those in Secs. 3.1 and 3.2.

Table 3 shows the results. These conditions did not greatly affect accuracy, confirming the appropriateness of our experimental conditions.

3.4 Evaluation of phone models

Finally, we compared the phone models trained from scratch with the phone models adapted from those for speech recognition. In [10], the phone models were prepared by adapting phone models for speech recognition with a small number of training data (10 songs with the phoneme-level annotation). They performed well for the lyrics synchronization problem, but the keyword detection problem, which is more difficult, requires more accurate phone models. We therefore prepared precise phoneme-level annotation and trained the phone models from scratch. To confirm the effectiveness of training of the phone models from a large amuont of training data, we conducted experiments using three phone models:

Table 4. Results of phoneme-level recognition: Comparison of three phone models.

	i. small adapt.	ii. large adapt.	iii. train.
Accuracy	27.1%	32.7%	50.9%

i. (small adaptation) Phone models created by adapting the phone model for speech recognition using 10 songs.

ii. (large adaptation) Phone models created by adapting the phone model for speech recognition using 79 songs.

iii. (training) Phone models trained from scrach using 79 songs.

The results are shown in Table 4. We found that the averaged frame level accuracy was drastically improved under the condition iii. This indicates that, when we have enough training data, phone models trained from scratch can perform better than the adapted ones based on speech recognition.

4 DISCUSSION

In this paper, we looked at the way of hyperlinking lyrics, i.e., creating bidirectional hyperlinks connecting lyrics keywords shared by different songs. Thus, we can generate a hyperlinked (networked) structure of songs. Our technology can be used when we only know the lyrics of a part of the songs. This step towards enabling the Music Web should help lead to a lyrics-based-MIR from songs that have unknown lyrics.

We can incorporate this method into *LyricSynchronizer* [10, 1], which displays scrolling lyrics with the phrase currently being sung highlighted during playback. A user interested in a different song containing the same hyperlinked keyword of the song currently being played can simply click on a highlighted keyword to jump to and listen from that keyword in a different song.

This is just the beginning for this new research topic, and because dealing with lyrics of polyphonic music signals is challenging, the performance represented by the experimental results given in Sec. 3.1 still need to be im-

proved. We expect to improve the overall performance by refining the phone model. As shown in Sec. 3.4, we can improve it by preparing the annotation for more songs. Moreover, as speech recognition technologies have shown, once we obtain good initial phone models, we can improve them through transcribing audio signals (i.e., lyrics) without requiring precise phoneme-level annotation.

Our experience in developing the keyword extraction method described in Sec. 2.1.1 has shown that the well known tf-idf (term frequency-inverse document frequency) is not useful for extracting keywords that have as many phonemes as possible but appear in two songs or more. The tf-idf is designed based of the assumptions that the importance of a keyword for a document is proportional to the number of occurrences of the keyword in the document is and inversely proportional to the number of occurrences of the keyword in all documents. However, the number of keyword occurrences in a song in our problem is unimportant. Even if a keyword appears only once, a user can jump from that keyword to other songs of interest. Therefore, we developed our own keyword extraction method.

5 CONCLUSION

This paper described a method for creating hyperlinks between different songs at the level of lyrics phrases. We presented a method for dealing with an important problem that has received little attention. We created two kinds of hyperlinks: hyperlinks between songs where the text lyrics are known and those from songs with text lyrics to songs without text lyrics. We created these hyperlinks by extracting keywords from the text lyrics of all songs and by detecting the keywords in polyphonic audio signals by using a HMM-based keyword spotting technique. Our experimental results show that the approach is promising, but better performance is needed for practical application. We will apply our method to a much larger database in the future. We also plan to develop useful applications for this method, including song clustering based on the songs' lyrics, and a Music Web browser that enables users to browse and discover interesting songs by clicking the *hyperlyrics* of songs.

6 ACKNOWLEDGEMENTS

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JOINT STRUCTURE ANALYSIS WITH APPLICATIONS TO MUSIC ANNOTATION AND SYNCHRONIZATION

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ABSTRACT

The general goal of music synchronization is to automatically align different versions and interpretations related to a given musical work. In computing such alignments, recent approaches assume that the versions to be aligned correspond to each other with respect to their overall global structure. However, in real-world scenarios, this assumption is often violated. For example, for a popular song there often exist various structurally different album, radio, or extended versions. Or, in classical music, different recordings of the same piece may exhibit omissions of repetitions or significant differences in parts such as solo cadenzas. In this paper, we introduce a novel approach for automatically detecting structural similarities and differences between two given versions of the same piece. The key idea is to perform a single structural analysis for both versions simultaneously instead of performing two separate analyses for each of the two versions. Such a joint structure analysis reveals the repetitions within and across the two versions. As a further contribution, we show how this information can be used for deriving musically meaningful partial alignments and annotations in the presence of structural variations.

1 INTRODUCTION

Modern digital music collections contain an increasing number of relevant digital documents for a single musical work comprising various audio recordings, MIDI files, or symbolic score representations. In order to coordinate the multiple information sources, various synchronization procedures have been proposed to automatically align musically corresponding events in different versions of a given musical work, see [1, 7, 8, 9, 14, 15] and the references therein. Most of these procedures rely on some variant of dynamic time warping (DTW) and assume a global correspondence of the two versions to be aligned. In real-world scenarios, however, different versions of the same piece may exhibit significant structural variations. For example, in the case of Western classical music, different recordings often exhibit omissions of repetitions (e. g., in sonatas and symphonies) or significant differences in parts such as solo cadenzas of concertos. Similarly, for a given popular, folk, or art song, there may be various recordings with a different number of stanzas. In particular for popular songs, there may exist structurally different album, radio, or extended versions as well as cover versions.

A basic idea to deal with structural differences in the synchronization context is to combine methods from music structure analysis and music alignment. In a first step, one may partition the two versions to be aligned into musically meaningful segments. Here, one can use methods from automated structure analysis [3, 5, 10, 12, 13] to derive similarity clusters that represent the repetitive structure of the two versions. In a second step, the two versions can then be compared on the segment level with the objective for matching musically corresponding passages. Finally, each pair of matched segments can be synchronized using global alignment strategies. In theory, this seems to be a straightforward approach. In practise, however, one has to deal with several problems due to the variability of the underlying data. In particular, the automated extraction of the repetitive structure constitutes a delicate task in case the repetitions reveal significant differences in tempo, dynamics, or instrumentation. Flaws in the structural analysis, however, may be aggravated in the subsequent segment-based matching step leading to strongly corrupted synchronization results.

The key idea of this paper is to perform a single, joint structure analysis for both versions to be aligned, which provides richer and more consistent structural data than in the case of two separate analyses. The resulting similarity clusters not only reveal the repetitions within and across the two versions, but also induce musically meaningful partial alignments between the two versions. In Sect. 2, we describe our procedure for a joint structure analysis. As a further contribution of this paper, we show how the joint structure can be used for deriving a musically meaningful partial alignment between two audio recordings with structural differences, see Sect. 3. Furthermore, as described in Sect. 4, our procedure can be applied for automatic annotation of a given audio recording by partially available MIDI data. In Sect. 5, we conclude with a discussion of open problems and

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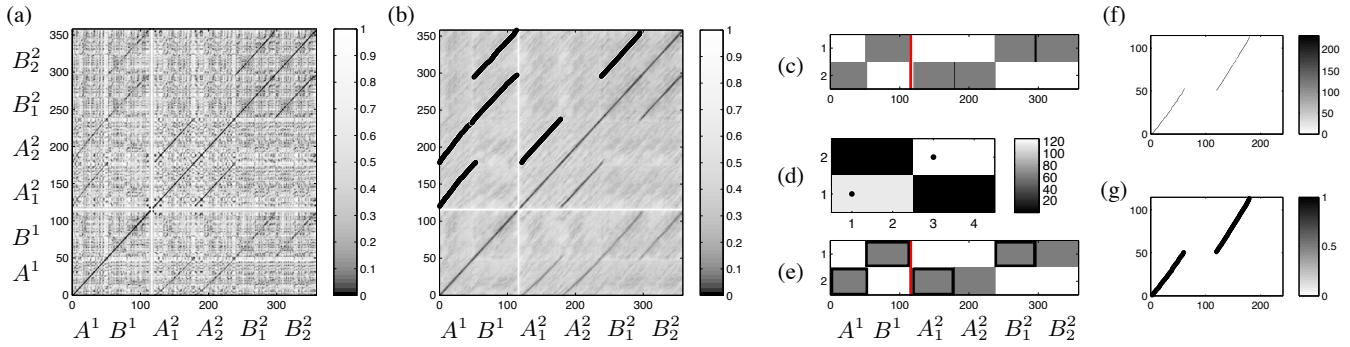

Figure 1. Joint structure analysis and partial synchronization for two structurally different versions of the Aria of the Goldberg Variations BWV 988 by J.S. Bach. The first version is played by G. Gould (musical form A^1B^1) and the second by M. Perahia (musical form $A_1^2A_2^2B_1^2B_2^2$). **(a)** Joint similarity matrix $\mathcal{S}$. **(b)** Enhanced matrix and extracted paths. **(c)** Similarity clusters. **(d)** Segment-based score matrix $\mathcal{M}$ and match (black dots). **(e)** Matched segments. **(f)** Matrix representation of matched segments. **(g)** Partial synchronization result.

prospects on future work.

The problem of automated partial music synchronization has been introduced in [11], where the idea is to use the concept of path-constrained similarity matrices to enforce musically meaningful partial alignments. Our approach carries this idea even further by using cluster-constraint similarity matrices, thus enforcing structurally meaning partial alignments. A discussion of further references is given in the subsequent sections.

2 JOINT STRUCTURE ANALYSIS

The objective of a joint structure analysis is to extract the repetitive structure within and across two different music representations referring to the same piece of music. Each of the two versions can be an audio recording, a MIDI version, or a MusicXML file. The basic idea of how to couple the structure analysis of two versions is very simple. First, one converts both versions into common feature representations and concatenates the resulting feature sequences to form a single long feature sequence. Then, one performs a common structure analysis based on the long concatenated feature sequence. To make this strategy work, however, one has to deal with various problems. First, note that basically all available procedures for automated structure analysis have a computational complexity that is at least quadratic in the input length. Therefore, efficiency issues become crucial when considering a single concatenated feature sequence. Second, note that two different versions of the same piece often reveal significant local and global tempo differences. Recent approaches to structure analysis such as [5, 12, 13], however, are built upon the constant tempo assumption and cannot be used for a joint structure analysis. Allowing also tempo variations between repeating segments makes the structure analysis problem a much harder problem [3, 10]. We now summarize the approach used in this paper closely following [10].

Given two music representations, we transform them into suitable feature sequences $U := (u^1, u^2, \ldots, u^L)$ and $V := (v^1, v^2, \ldots, v^M)$, respectively. To reduce different types of music data (audio, MIDI, MusicXML) to the same type of representation and to cope with musical variations in instrumentation and articulation, chroma-based features have turned out to be a powerful mid-level music representation [2, 3, 8]. In the subsequent discussion, we employ a smoothed normalized variant of chroma-based features (CENS features) with a temporal resolution of 1 Hz, see [8] for details. In this case, each 12-dimensional feature vector u^ℓ, $\ell \in [1:L]$, and v^m, $m \in [1:M]$, expresses the local energy of the audio (or MIDI) distribution in the 12 chroma classes. The feature sequences strongly correlate to the short-time harmonic content of the underlying music representations. We now define the sequence W of length $N := L + M$ by concatenating the sequences U and V:

$$W := (w^1, w^2, \ldots, w^N) := (u^1, \ldots, u^L, v^1, \ldots, v^M).$$

Fixing a suitable local similarity measure— here, we use the inner vector product—the $(N \times N)$-*joint similarity matrix* $\mathcal{S}$ is defined by $\mathcal{S}(i,j) := \langle w^i, w^j \rangle$, $i, j \in [1:N]$. Each tuple (i,j) is called a *cell* of the matrix. A *path* is a sequence $p = (p_1, \ldots, p_K)$ with $p_k = (i_k, j_k) \in [1:N]^2$, $k \in [1:K]$, satisfying $1 \leq i_1 \leq i_2 \leq \ldots \leq i_K \leq N$ and $1 \leq j_1 \leq j_2 \leq \ldots \leq j_K \leq N$ (monotonicity condition) as well as $p_{k+1} - p_k \in \Sigma$, where Σ denotes a set of admissible step sizes. In the following, we use $\Sigma = \{(1,1), (1,2), (2,1)\}$.

As an illustrative example, we consider two different audio recordings of the Aria of the Goldberg Variations BWV 988 by J.S. Bach, in the following referred to as *Bach example*. The first version with a duration of 115 seconds is played by Glen Gould without repetitions (corresponding to the musical form A^1B^1) and the second version with a duration of 241 seconds is played by Murray Perahia with repetitions (corresponding to the musical form $A_1^2A_2^2B_1^2B_2^2$). For the feature sequences hold $L = 115$, $M = 241$, and $N = 356$. The resulting joint similarity matrix is shown in

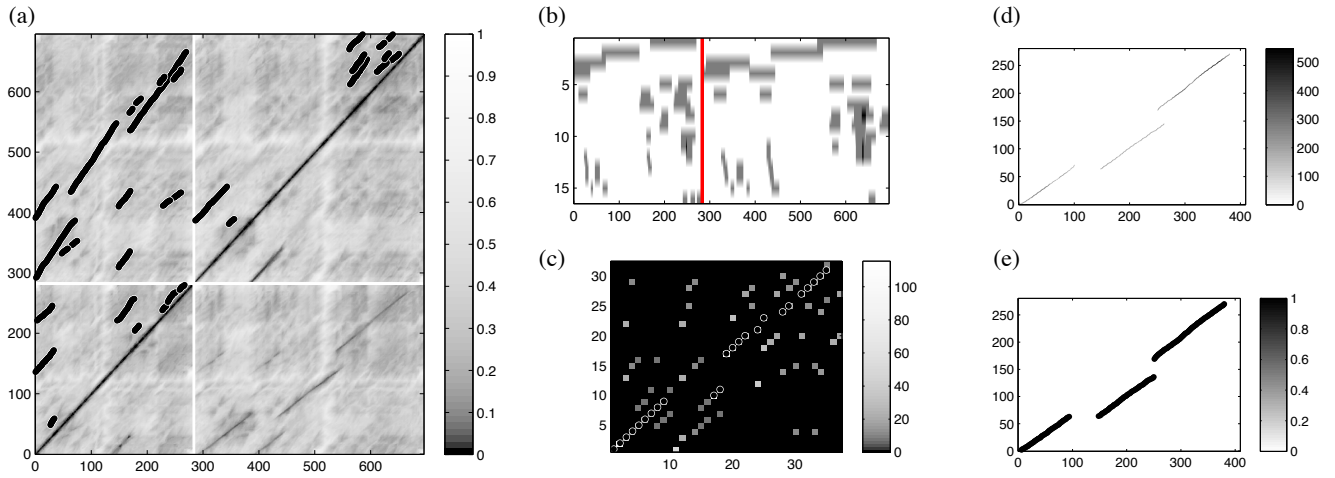

Figure 2. Joint structure analysis and partial synchronization for two structurally modified versions of Beethoven's Fifth Symphony Op. 67. The first version is a MIDI version and the second one an audio recording by Bernstein. **(a)** Enhanced joint similarity matrix and extracted paths. **(b)** Similarity clusters. **(c)** Segment-based score matrix $\mathcal{M}$ and match (indicated by black dots). **(d)** Matrix representation of matched segments. **(e)** Partial synchronization result.

Fig. 1a, where the boundaries between the two versions are indicated by white horizontal and vertical lines.

In the next step, the path structure is extracted from the joint similarity matrix. Here, the general principle is that each path of low cost running in a direction along the main diagonal (gradient $(1, 1)$) corresponds to a pair of similar feature subsequences. Note that relative tempo differences in similar segments are encoded by the gradient of the path (which is then in a neighborhood of $(1, 1)$). To ease the path extraction step, we enhance the path structure of $\mathcal{S}$ by a suitable smoothing technique that respects relative tempo differences. The paths can then be extracted by a robust and efficient greedy strategy, see Fig. 1b. Here, because of the symmetry of $\mathcal{S}$, one only has to consider the upper left part of $\mathcal{S}$. Furthermore, we prohibit paths crossing the boundaries between the two versions. As a result, each extracted path encodes a pair of musically similar segments, where each segment entirely belongs either to the first or to the second version. To determine the global repetitive structure, we use a one-step transitivity clustering procedure, which balances out the inconsistencies introduced by inaccurate and incorrect path extractions. For details, we refer to [8, 10].

Altogether, we obtain a set of similarity clusters. Each similarity cluster in turn consists of a set of pairwise similar segments encoding the repetitions of a segment within and across the two versions. Fig. 1c shows the resulting set of similarity clusters for our Bach example. Both of the clusters consist of three segments, where the first cluster corresponds to the three B-parts B^1, B_1^2, and B_2^2 and the second cluster to the three A-parts A^1, A_1^2, and A_2^2. The joint analysis has several advantages compared to two separate analyses. First note that, since there are no repetitions in the first version, a separate structure analysis for the first version would not have yielded any structural information. Second, the similarity clusters of the joint structure analysis naturally induce musically meaningful partial alignments between the two versions. For example, the first cluster shows that B^1 may be aligned to B_1^2 or to B_2^2. Finally, note that the delicate path extraction step often results in inaccurate and fragmented paths. Because of the transitivity step, the joint clustering procedure balances out these flaws and compensates for missing parts to some extent by using joint information *across* the two versions.

On the downside, a joint structural analysis is computationally more expensive than two separate analyses. Therefore, in the structure analysis step, our strategy is to use a relatively low feature resolution of 1 Hz. This resolution may then be increased in the subsequent synchronization step (Sect. 3) and annotation application (Sect. 4). Our current MATLAB implementation can easily deal with an overall length up to $N = 3000$ corresponding to more then forty minutes of music material. (In this case, the overall computation time adds up to 10-400 seconds with the path extraction step being the bottleneck, see [10]). Thus, our implementation allows for a joint analysis even for long symphonic movements of a duration of more than 20 minutes.

Another drawback of the joint analysis is that local inconsistencies across the two versions may cause an overfragmentation of the music material. This may result in a large number of incomplete similarity clusters containing many short segments. As an example, we consider a MIDI version as well as a Bernstein audio recording of the first movement of Beethoven's Fifth Symphony Op. 67. We structurally modified both versions by removing some sections. Fig. 2a shows the enhanced joint similarity matrix and Fig. 2b the set of joint similarity clusters. Note that

some of the resulting 16 clusters contain semantically meaningless segments stemming from spuriously extracted path fragments. At this point, one could try to improve the overall structure result by a suitable postprocessing procedure. This itself constitutes a difficult research problem and is not in the scope of this paper. Instead, we introduce a procedure for partial music alignment, which has some degree of robustness to inaccuracies and flaws in the previously extracted structural data.

3 PARTIAL SYNCHRONIZATION

Given two different representations of the same underlying piece of music, the objective of *music synchronization* is to automatically identify and link semantically corresponding events within the two versions. Most of the recent synchronization approaches use some variant of dynamic time warping (DTW) to align the feature sequences extracted from the two versions, see [8]. In classical DTW, all elements of one sequence are matched to elements in the other sequence (while respecting the temporal order). This is problematic when elements in one sequence do not have suitable counterparts in the other sequence. In the presence of structural differences between the two sequences, this typically leads to corrupted and musically meaningless alignments [11]. Also more flexible alignment strategies such as subsequence DTW or partial matching strategies as used in biological sequence analysis [4] do not properly account for such structural differences.

A first approach for partial music synchronization has been described in [11]. Here, the idea is to first construct a path-constrained similarity matrix, which a priori constricts possible alignment paths to a semantically meaningful choice of admissible cells. Then, in a second step, a path-constrained alignment can be computed using standard matching procedures based on dynamic programming.

We now carry this idea even further by using the segments of the joint similarity clusters as constraining elements in the alignment step. To this end, we consider pairs of segments, where the two segments lie within the same similarity cluster and belong to different versions. More precisely, let $\mathcal{C} = \{C_1, \ldots, C_M\}$ be the set of clusters obtained from the joint structure analysis. Each similarity cluster C_m, $m \in [1 : M]$, consists of a set of segments (i.e., subsequences of the concatenated feature sequence W). Let $\alpha \in C_m$ be such a segment. Then let $\ell(\alpha)$ denote the length of α and $c(\alpha) := m$ the cluster affiliation. Recall that α either belongs to the first version (i.e., α is a subsequence of U) or to the second version (i.e., α is a subsequence of V). We now form two lists of segments. The first list $(\alpha_1, \ldots, \alpha_I)$ consists of all those segments that are contained in some cluster of $\mathcal{C}$ and belong to the first version. The second list $(\beta_1, \ldots, \beta_J)$ is defined similarly, where the segments now belong to the second version. Both lists are sorted according to the start positions of the segments. (In case two segments have the same start position, we break the tie by also considering the cluster affiliation.) We define a segment-based $I \times J$-score matrix $\mathcal{M}$ by

$$\mathcal{M}(i,j) := \begin{cases} \ell(\alpha_i) + \ell(\beta_j) & \text{for } c(\alpha_i) = c(\beta_j), \\ 0 & \text{otherwise,} \end{cases}$$

$i \in [1 : I], j \in [1 : J]$. In other words, $\mathcal{M}(i,j)$ is positive if and only if α_i and β_j belong to the same similarity cluster. Furthermore, $\mathcal{M}(i,j)$ depends on the lengths of the two segments. Here, the idea is to favor long segments in the synchronization step. For an illustration, we consider the Bach example of Fig. 1, where $(\alpha_1, \ldots, \alpha_I) = (A^1, B^1)$ and $(\beta_1, \ldots, \beta_J) = (A_1^2, A_2^2, B_1^2, B_2^2)$. The resulting matrix $\mathcal{M}$ is shown in Fig. 1d. For another more complex example, we refer to Fig. 2c.

Now, a segment-based *match* is a sequence $\mu = (\mu_1, \ldots, \mu_K)$ with $\mu_k = (i_k, j_k) \in [1 : I] \times [1 : J]$ for $k \in [1 : K]$ satisfying $1 \leq i_1 < i_2 < \ldots < i_K \leq I$ and $1 \leq j_1 < j_2 < \ldots < j_K \leq J$. Note that a match induces a partial assignment of segment pairs, where each segment is assigned to at most one other segment. The *score* of a match μ with respect to $\mathcal{M}$ is then defined as $\sum_{k=1}^{K} \mathcal{M}(i_k, j_k)$. One can now use standard techniques to compute a score-maximizing match based on dynamic programming, see [4, 8]. For details, we refer to the literature. In the Bach example, the score-maximizing match μ is given by $\mu = ((1,1),(2,3))$. In other words, the segment $\alpha_1 = A^1$ of the first version is assigned to segment $\beta_1 = A_1^2$ of the second version and $\alpha_2 = B^1$ is assigned to $\beta_3 = B_1^2$.

In principle, the score-maximizing match μ constitutes our partial music synchronization result. To make the procedure more robust to inaccuracies and to remove cluster redundancies, we further clean the synchronization result in a postprocessing step. To this end, we convert the score-maximizing match μ into a sparsepath-constrained similarity matrix $\mathcal{S}^{\text{path}}$ of size $L \times M$, where L and M are the lengths of the two feature sequences U and V, respectively. For each pair of matched segments, we compute an alignment path using a global synchronization algorithm [9].Each cell of such a path defines a non-zero entry of $\mathcal{S}^{\text{path}}$, where the entry is set to the length of the path (thus favoring long segments in the subsequent matching step). All other entries of the matrix $\mathcal{S}^{\text{path}}$ are set to zero. Fig. 1f and Fig. 2d show the resulting path-constrained similarity matrices for the Bach and Beethoven example, respectively. Finally, we apply the procedure as described in [11] using $\mathcal{S}^{\text{path}}$ (which is generally much sparser than the path-constrained similarity matrices as used in [11])to obtain a purified synchronization result, see Fig. 1g and Fig. 2e.

To evaluate our synchronization procedure, we performed similar experiments as described in [11]. In one experiment, we formed synchronization pairs each consisting of two different versions of the same piece. Each pair

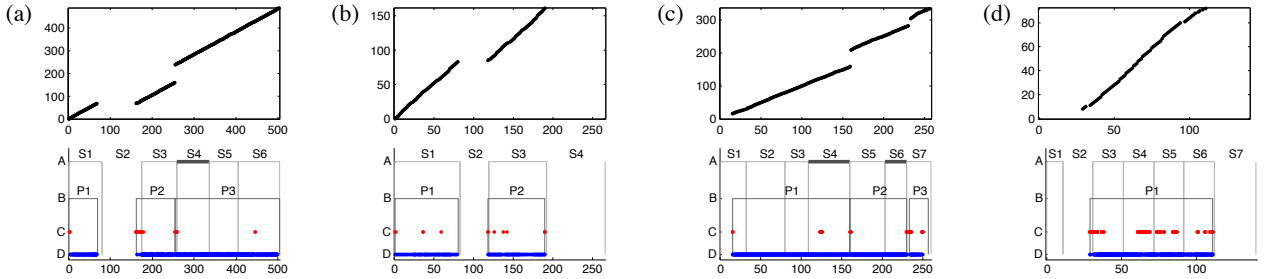

Figure 3. Partial synchronization results for various MIDI-audio synchronization pairs. The top figures show the final path components of the partial alignments and the bottom figures indicate the ground truth (Row A), the final annotations (Row B), and a classification into correct (Row D) and incorrect annotations (Row C), see text for additional explanations. The pieces are specified in Table 1. **(a)** Haydn (RWC C001), **(b)** Schubert (RWC C048, distorted), **(c)** Burke (P093), **(d)** Beatles ("Help!", distorted).

consists either of an audio recording and a MIDI version or of two different audio recordings (interpreted by different musicians possibly in different instrumentations). We manually labeled musically meaningful sections of all versions and then modified the pairs by randomly removing or duplicating some of the labeled sections, see Fig. 3. The partial synchronization result computed by our algorithm was analyzed by means of its path components. A path component is said to be *correct* if it aligns corresponding musical sections. Similarly, a match is said to be *correct* if it covers (up to a certain tolerance) all semantically meaningful correspondences between the two versions (this information is given by the ground truth) and if all its path components are correct. We tested our algorithm on more than 387 different synchronization pairs resulting in a total number of 1080 path components. As a result, 89% of all path components and 71% of all matches were correct (using a tolerance of 3 seconds).

The results obtained by our implementation of the segment-based synchronization approach are qualitatively similar to those reported in [11]. However, there is one crucial difference in the two approaches. In [11], the authors use a combination of various ad-hoc criteria to construct a path-constrained similarity matrix as basis for their partial synchronization. In contrast, our approach uses only the structural information in form of the joint similarity clusters to derive the partial alignment. Furthermore, the availability of structural information within and across the two versions allows for recovering missing relations based on suitable transitivity considerations. Thus, each improvement of the structure analysis will have a direct positive effect on the quality of the synchronization result.

4 AUDIO ANNOTATION

The synchronization of an audio recording and a corresponding MIDI version can be regarded as an automated annotation of the audio recording by means of the explicit note events given by the MIDI file. Often, MIDI versions are used as a kind of score-like symbolic representation of the underlying musical work, where redundant information such as repetitions are not encoded explicitly. This is a further setting with practical relevance where two versions to be aligned have a different repetitive structure (an audio version with repetitions and a score-like MIDI version without repetitions). In this setting, one can use our segment-based partial synchronization to still obtain musically adequate audio annotations.

We now summarize one of our experiments, which has been conducted on the basis of synchronization pairs consisting of structurally equivalent audio and MIDI versions.[1] We first globally aligned the corresponding audio and MIDI versions using a temporally refined version of the synchronization procedure described in [9]. These alignments were taken as ground truth for the audio annotation. Similar to the experiment of Sect. 3, we manually labeled musically meaningful sections of the MIDI versions and randomly removed or duplicated some of these sections. Fig. 3a illustrates this process by means of the first movement of Haydn's Symphony No. 94 (RWC C001). **Row A** of the bottom part shows the original six labeled sections S1 to S6 (warped according to the audio version). In the modification, S2 was removed (no line) and S4 was duplicated (thick line). Next, we partially aligned the modified MIDI with the original audio recording as described in Sect. 3. The resulting three path components of our Haydn example are shown in the top part of Fig. 3a. Here, the vertical axis corresponds to the MIDI version and the horizontal axis to the audio version. Furthermore, **Row B** of the bottom part shows the projections of the three path components onto the audio axis resulting in the three segments P1, P2, and P3. These segments are aligned to segments in the MIDI thus being annotated by the corresponding MIDI events. Next, we compared these partial annotations with the ground truth annotations on the MIDI note event level. We say that an alignment of a note event to a physical time position of the audio version is correct in a *weak* (*strong*) sense, if there is

[1] Most of the audio and MIDI files were taken from the RWC music database [6]. Note that for the classical pieces, the original RWC MIDI and RWC audio versions are not aligned.

Composer	Piece	RWC	Original		Distorted	
			weak	strong	weak	strong
Haydn	Symph. No. 94, 1st Mov.	C001	98	97	97	95
Beethoven	Symph. Op. 67, 1st Mov.	C003	99	98	95	91
Beethoven	Sonata Op. 57, 1st Mov.	C028	99	99	98	96
Chopin	Etude Op. 10, No. 3	C031	93	93	93	92
Schubert	Op. 89, No. 5	C048	97	96	95	95
Burke	Sweet Dreams	P093	88	79	74	63
Beatles	Help!	—	97	96	77	74
Average			96	94	91	87

Table 1. Examples for automated MIDI-audio annotation (most of files are from the RWC music database [6]). The columns show the composer, the piece of music, the RWC identifier, as well as the annotation rate (in %) with respect to the weak and strong criterion for the original MIDI and some distorted MIDI.

a ground truth alignment of a note event of the same pitch (and, in the strong case, additionally lies in the same musical context by checking an entire neighborhood of MIDI notes) within a temporal tolerance of 100 ms. In our Haydn example, the weakly correct partial annotations are indicated in **Row D** and the incorrect annotations in **Row C**.

The other examples shown in Fig. 3 give a representative impression of the overall annotation quality. Generally, the annotations are accurate—only at the segment boundaries there are some larger deviations. This is due to our path extraction procedure, which often results in "frayed" path endings. Here, one may improve the results by correcting the musical segment boundaries in a postprocessing step based on cues such as changes in timbre or dynamics. A more critical example (Beatles example) is shown Fig. 3d, where we removed two sections (S2 and S7) from the MIDI file and temporally distorted the remaining parts. In this example, the MIDI and audio version also exhibit significant differences on the feature level. As a result, an entire section (S1) has been left unannotated leading to a relatively poor rate of 77% (74%) of correctly annotated note events with respect to the weak (strong) criterion.

Finally, Table 1 shows further rates of correctly annotated note events for some representative examples. Additionally, we have repeated our experiments with significantly temporally distorted MIDI files (locally up to ±20%). Note that most rates only slightly decrease (e. g., for the Schubert piece, from 97% to 95% with respect to the weak criterion), which indicates the robustness of our overall annotation procedure to local tempo differences. Further results as well as audio files of sonifications can be found at `http://www-mmdb.iai.uni-bonn.de/projects/partialSync/`

5 CONCLUSIONS

In this paper, we have introduced the strategy of performing a joint structural analysis to detect the repetitive structure within and across different versions of the same musical work. As a core component for realizing this concept, we have discussed a structure analysis procedure that can cope with relative tempo differences between repeating segments. As further contributions, we have shown how joint structural information can be used to deal with structural variations in synchronization and annotation applications. The tasks of *partial* music synchronization and annotation is a much harder then the *global* variants of these tasks. The reason for this is that in the partial case one needs *absolute* similarity criteria, whereas in the global case one only requires *relative* criteria. One main message of this paper is that automated music structure analysis is closely related to partial music alignment and annotation applications. Hence, improvements and extensions of current structure analysis procedures to deal with various kinds of variations is of fundamental importance for future research.

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Music, Movies and Meaning: Communication in Film-makers' Search for Pre-existing Music, and the Implications for Music Information Retrieval.

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ABSTRACT

While the use of music to accompany moving images is widespread, the information behaviour, communicative practice and decision making by creative professionals within this area of the music industry is an under-researched area. This investigation discusses the use of music in films and advertising focusing on communication and meaning of the music and introduces a reflexive communication model. The model is discussed in relation to interviews with a sample of music professionals who search for and use music for their work. Key factors in this process include stakeholders, briefs, product knowledge and relevance. Searching by both content and context is important, although the final decision when matching music to picture is partly intuitive and determined by a range of stakeholders.

1. INTRODUCTION

Although evidence from the period is sparse, it is likely that music has been used to accompany moving images since the Lumiere brothers' presentations in 1896 [17]. Music was either written especially for the purpose, or consisted of a mixture of well-known favourites drawn from classical and popular repertoires. Gradually these ad hoc combinations of music and film have led to a multi-million dollar worldwide creative industry and there is an accompanying wealth of theory on how music works with film. Directors such as Quentin Tarrantino and the Coen Brothers spearhead a wave of film makers using existing music in their productions, and this widespread trend has spun off into television, advertising, and computer games. This widespread use raises the questions, 'who chooses what, why and how?', which this investigation attempts to answer. The research focuses on identifying the communicative practice and decision making undertaken by creative professionals who choose music for multimedia texts (films, adverts, television programmes). This set of professionals is an under-researched sub-group of the music industry, and this paper contributes to the literature by focussing on their work processes, which may inform the development of systems that attempt to meet those users' information needs.

2. BACKGROUND

2.1. Use of music in films

Although music was originally used as a mood enhancer and narrative aid for the cinema audience it gradually became an essential part of the film itself, 'to colour a scene, to suggest a general mood, to intensify a narrative or emotional tension' [17:145]. The interpretation of film music depends on the listener [21] and there are two main areas of analysis and interpretation, the music itself, and its interaction with the film [9]. The meaning of the music is successfully transmitted to the audience through means of agreed cultural codes – whether these are major/minor for happy/sad or consonance/dissonance as light/shade, as well as style topics, tonal design, leitmotiv, timbre and musical and filmic form [21]. Using pre-existing music means considering how the viewers' familiarity with the music might determine meaning, and being aware that these meanings can change [23].

2.2. Communication And Meaning

As the research subjects of this investigation describe their needs with words, a semiotic approach may be of value in successfully analysing the music in their collections. In semiotics, signs, (including words). are interpreted by the community according to agreed codes. Musical signs would include elements of notation (slurs, clefs), the music itself and its surrounding texts (cd sleeves, photos, videos), and genre names and theory terms [26]. Owing to the importance of connotational signification within music, where the use of signs/sounds such as a car horn sound effect within a piece is to act as music rather than to tell you a fast moving vehicle is approaching, in order to understand the meaning of music it is important to learn the cultural meanings encoded by the music. It is only through this that the intertextuality or dialogism of modern popular music, which frequently carries important codes by referencing other music, films and literature, can be fully appreciated.

If music retrieval systems are to effectively retrieve music for the end user then they need to be able to incorporate the listener's meaning of music. Different listeners have varying ways of giving meaning to the music they hear [19], and historically this has affected the

way it is organised and retrieved. The context and the content of the music both come into play. Although music semiotics initially focussed on the content of the music there have been moves towards analysis of the contextual elements [4, 20, 25, 26]. This reflects calls both within the text information retrieval (IR) [29] and music information retrieval (MIR) communities [11] for user centred research. Musical code links the sound and its context, and understanding these codes should help establish musical meaning [27]. In the case of music, codes become increasingly specific: langue, norms, sub-norms, dialects, styles, genres, sub-codes, idiolects, works/performances [4, 20]. Competences are required to successfully interpret these codes and these competences will vary according to experience, background and interests [25]. These can be incorporated into established communications models [24] in order to explain the communication process that is taking place when listening to music. Hall [12] discussed how encoding and decoding cultural messages would lead to the audience influencing the message as well as determining its meaning. A key model was developed by Tagg [26] which incorporated codes and competences as symbols and socio-cultural norms. Here the problems of mis-communication are caused by incompetence and interference. The flow of information is also described as a one-way process (from Transmitter or producer to Receiver or listener). It has been proposed that a user centred model would more accurately reflect the process of meaning-making when choosing music on behalf of others, as meanings shared by User and Owner will be used to determine the relevance of music choices [14].

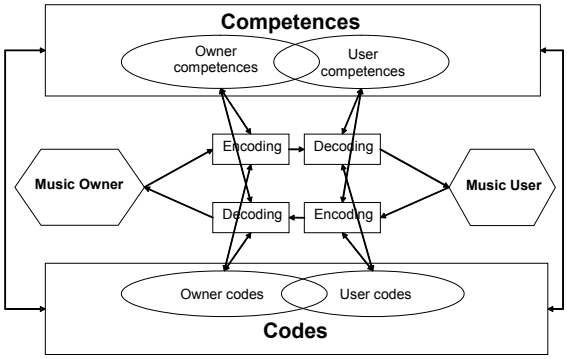

Figure 1 User centred communication model [14]

This model underpins the development of the current investigation into the musical decision making processes undertaken by creative professionals working in, or with, the film and television industries. It has also helped develop the aims, objectives and methodology of the bigger MIR project of which the current investigation is one element [14].

3. METHODOLOGY

3.1. Approach

A user-centred approach was taken for this investigation. There have been numerous examples of qualitative user-focussed research in both the text IR [2, 8, 13, 15, 29] and MIR communities [1, 7, 10, 18]. Some of this research has used surveys to investigate large populations of recreational users, while others interview users, and examine rich web queries and 'music discovery' diaries.

A breakdown of the music industry was derived from British Phonographic Industry literature [3] which showed the industry comprises two key areas: Music Owners (Artists, Record Companies, Music Publishers) on one side and Music Consumers (Recreational users) on the other. An additional layer is found between these two sides, informed by the researcher's extensive experience working in the commercial music industry. This layer, which includes advertising agencies, film companies, and broadcasters, is a 'filtering' layer of Music Users, which exploits the music in some way before it reaches the Consumer. These organisations are choosing music on behalf of Consumers rather than for their own recreational purposes. It is the aim of this research to investigate the communication process between the Music Owners and the Music Users (Fig 1).

3.2. Sample

The respondents interviewed all work with music, are mainly at a senior level and are very experienced in the use of music when accompanying moving images. Ten people were interviewed: five represented the music rights holders (publishers, record companies) and five worked with music in the film and television industries. The respondents were chosen using a snowball sampling technique [22] which, although biased owing to interviewees being inter-connected, is an accepted theoretical sampling method, which has been used to gain access to 'hidden' or constantly changing communities. It is more flexible than random sampling, allowing the researcher to follow leads or themes from one interview to the next, and a normally quite inaccessible population with expert knowledge is more readily available. The sample will be widened as research develops, in order to determine the generalisability of these results.

3.3. Interviews

Semi-structured face-to-face interviews were done at a time and place convenient to the subject, lasting up to one hour. The interviews were recorded digitally and transcribed by the researcher. Before each interview the researcher noted a number of areas to discuss, based on what had already been discovered from previous interviews and other research. This flexible approach meant that there was no need to ask redundant questions

and allowed the interviewees to talk freely about the issues that were important to them.

3.4. Topics and Questions

The exact wording of the questions varied, and sometimes the planned questions were not asked directly because answers had been given answering another question. However the researcher always had a list of areas to cover. These focussed on the process, the use, queries, communication, meaning, relevance and relationships. Areas of discussion included participant's work role and relationships with other stakeholders; the extent to which they use and search for music in their job; queries and systems they use in searching; the type of material that satisfies their requests; and whether they could recommend improvements to the process.

3.5. Ethics

University ethics guidelines were used as a framework, and all subjects were sent an explanatory statement and signed a consent form. Each interviewee was given a code to aid in anonymising their comments, and information which may identify them has been removed.

3.6. Analysis

The transcribed texts and recordings were read and listened to, focussing on discussions of the process and the issues that were felt by the respondents to be of importance. Common areas which were raised are discussed in section 4. This is a preliminary analysis designed to inform the researcher of the main issues and how the process takes place, these are subsequently compared to the reflexive communication model in Figure 1. This research is part of an investigation into the information seeking behaviour of Users, which is significantly influenced by the world they are a part of. The discussion that follows highlights some of these real world issues and relates them to Figure 1.

4. RESULTS AND DISCUSSION

4.1. Stakeholders

The interviews were started by discussing a subject the respondents knew well, their day-to-day work and their perceived role in the process. This allowed them to settle in comfortably to the interview process and relax. One of the key themes raised by this question were how on both sides there were large numbers of stakeholders involved in the process, in addition to the Consumers (Fig 2).

'You get a creative team, who write the script, who are probably in their twenties, and invariably, whatever music they're into, be it hiphop, electronica, grime, whatever, they'll want that on the ad. You get a creative director, their boss, who's probably a generation older, who wants something he's never heard on an ad before, so it stands out in the ad break and gets noticed, and then you've got the client, who's probably into classical or jazz or something like that, and he'll want that on it, and then you've got the target market, Debbie the housewife, who's 35 got three kids and lives in Sheffield who's into Madness. And you've got to find one track which all those people will buy into and say, 'Yes, we want this track on our ad'. So it is difficult at the best of times.' (003 music publisher)

Music Owners	Music Users
Synchronisation Dept, Legal / Business Affairs, Composer, Performer, Marketing & Promotions, Artists & Repertoire	Producer, Director, Film Editor, Music Editor, Music Supervisor, Client, Director, Ad Agency Creatives.

Figure 2 Stakeholders

Once a selection of pieces of music has been offered to the User, the stakeholders are brought in to reach a decision on the piece of music that will be used. Each of the stakeholders is likely to have different motivations, codes and competences, and these are not easily resolved. The criteria behind the decision making will include budget, clearance issues, aesthetic and commercial judgments. Whether the music enhances the picture seems to be the most important issue, although this will vary according to the producer (budget, market), director (picture as a whole), editor (the story), music supervisor (aesthetic and administrative) and audience (in test screenings). However this is not easily evaluated, most respondents resorting to 'gut feeling' when asked to describe what makes a great sync.

'And I think I think the hardest thing about searching for music for an advert is it's an opinion, and it's a take on a brief. And it's a take on visuals. And it's what you think works.' (007 record company)

4.2. Briefs

The query comes generally from the User to the Owner as a brief, which may take the form of a short informal email, a more widely circulated document which has been approved by internal User stakeholders, a script element, a conversation, or a moving image in the form of a clip of a completed ad.

'It's very easy to actually try stuff against picture and know whether it works and know whether a really odd idea is really worth playing somebody. (002 independent music supervisor)

The Owner may have the opportunity to clarify key issues with the User. These issues would include budget, deadlines, more detailed information on the use and context, or guidelines on the type of music that is sought.

The User frequently works with a 'temp' track of music that is not going to be used in the final product, but can be used for similarity queries. Normally a range of Owners are approached with the same or a similar query. These will be drawn from favoured or targeted contacts within the industry who are likely to make an offer that meets the aesthetic, time and budget constraints of the User.

According to the model, different musical codes are likely to have varying relevance for each query. For example, while a director may require a particular song, a music supervisor may extract the codes from that recording that are important to the meaning the director is attempting to convey, and match them to other recordings that may be more affordable or are not so obvious, uniqueness being an important decision-making factor.

'And sometimes if they want a track, for example, they can't have – let's say they want the Stones and there's no way they can afford it, they've only got thirty grand or something, you have to find something that's equally going to impress them as much as the Stones is. And I think that's by finding something unique that's going to work to picture.' (007 record company)

These codes would have increasingly granular specificity, from highly specific (different artists performing the same song in a similar style) through to different artists from other musical traditions that may have important perceived similarity, not necessarily musical but more relating to image and status. User codes are conveyed to the Owner as a brief, which may focus on the qualities of a product ('*speed, power, control, ability*', '*beauty, sophistication and intricacy of design*' *(011 ad agency supervisor*) or key words, such as '*tension, mystery, playfulness and warmth*' *(011 ad agency supervisor*). The Owner may have different competences and interpret these emotive connotations of signification differently to the User, although a sharing of codes and competences will reduce this semantic gap. A selection of songs are offered that are deemed to match the brief and the User makes a choice. There are frequent references to a 'left field' suggestion, which does not appear to meet a brief, being chosen as the 'perfect' sync. In these instances it is possible that while the User felt the song did not 'meet the brief', because of a mis-match between competences, the Owner's view of the track was that it was not at all left-field but an obvious match for this query.

4.3. Product knowledge

The Owners then attempt to match the brief to their catalogue, which may consist of up to one million pieces of music. If the brief is clear – a specific piece of music is required, or a specific genre, era, tempo or style is described, and if the Owners' catalogue is able to match those search terms to its metadata, the specific piece is found and terms negotiated, or a selection of pieces are offered to the User. These may be sent via email or ftp or on a cd, played to picture in an editing suite with the User present, or sent already matched roughly to picture using consumer software. The choice offered will vary in quantity depending on the brief. A small number of Owners offer a web-based facility to Users whereby they log in to production folders of music put together by the Owner, which include downloadable tracks as mp3 and wav (for broadcast). Then either feedback is received from the User to guide further searches, one of the songs is chosen and a license is negotiated, or the User does not contact the Owner again, choosing material elsewhere.

The search by the Owner is of particular interest. During the interviews some described how, rather than search the whole catalogue, they would refer to existing searches that they had done for similar queries, and discussed how they usually browsed their own collection, which was narrowed-down from the catalogue in terms of it's 'sync-friendly' nature. Removing material that was not likely to be used for synchronisation was very important to them, often citing narrative or offensive songs as being difficult to use owing to the lyrics obscuring the message of the picture. Their 'bespoke' collections are organized using very specific terms, such as 'instrumental – surf' and 'female vocal – 1950s' which they feel relate more to the queries they receive than the artist-album-title-genre format. An unclear brief means the Owner has to use their experience and product knowledge to 'second guess' the Users needs. Experience as a film maker helps in this area and some of the more successful practitioners have experience on 'the other side'. Most of them used iTunes as their personal media library and when doing a search referred both to their company catalogue and to the outside world, including the iTunes store and their own collections, and bringing in expert advisors from within and outside their company when specialist music was required that lay outside their own product knowledge. Great importance is placed on 'knowing the catalogue' rather than relying on a bespoke search engine and the respondents spend a large amount of time listening to music in order to familiarise themselves with it in order to maximise any opportunities for exploitation.

'A big part of my day is always every day at least fifty per cent will be listening to music. And if I can get more of the day to listen to music, I will. I mean there's just so much..' (005 independent music supervisor)

While the Owner is performing the search for the User, the User is exploring other avenues for music for use. They will not only be approaching other Owners, but will, again, search their own collections and look at their previous similar searches, use the iTunes store to listen to samples of tracks, and refer to web-based resources such as Google, All Music Guide, Pandora and last.fm. They also may have experience on 'the other side', are experienced in searching for music and are driven by a passion for music that is matched by in-depth knowledge.

4.4. Relevance

When the User is supplied with a selection of material this is listened to in order to evaluate its relevance and narrow down the selection. The choices are then matched to the visual. Other stakeholders are involved in this process, their input varying by their importance and ability to argue their case. While the relevance appears mainly to be gauged by 'gut-feeling', a term used by many respondents, probing revealed contextual factors such as target audience, genre, uniqueness, budget, availability, recognisability and chart position, and content features such as tempo, structure, mood, instrumentation, texture are all considered. It is here that the query becomes more concrete, and a further iteration can be made much more specific, when the User can refer to features of offered material as reference points.

5. IMPLICATIONS FOR MIR AND COMMUNICATION MODEL

It can clearly be seen from the analysis of the process that this is similar to the interactive information retrieval system used in online databases. The system consists of a knowledgeable and expert User, an equally knowledgeable and expert intermediary (the Owner) and a system (the record company or publisher's catalogue). It could be suggested that the expert intermediary be removed from the process (disintermediation) and the User allowed direct access to the collection [5]. This situation is discussed in a recent survey of graphic designers search for music [16] which indicated the value of online systems when looking for music to accompany presentations. This could reduce the problem with inadequately worded briefs as an interactive system would allow the User to adjust the query until a satisfactory result has been found. However it raises some problems. The Users have very limited time budgets and are reluctant to use these doing 'narrowing down' searches that can be delegated to others. Secondly, there are numerous Owners, so this approach would require the User doing searches on each Owner's bespoke search engine. However it appears that the expert knowledge of the Owners is frequently invaluable, as they know their catalogue well, and some of them have real experience in the use of music in film, and are prepared to match music to picture themselves in order to encourage the User to choose that piece of music over the others available. This experience in each other's world also means the Owner and User can encode and decode each other's meanings more accurately, they are aware of the other's codes and competences, and can accommodate these in forming their requests and results.

Mapping these results to the communications model in Figure 1, we have shown that there is a relationship between the Music Owners and the Music Users in terms of searching for and providing music. The Users have a piece of film, and are attempting to communicate meanings with that footage. These meanings are encoded into a query, in the form of an informal note or a brief, which is generated by a number of stakeholders, all with competing codes and competences. This brief can be of any level of specificity, ranging from a specific piece of music, to offering the piece of footage itself and asking for the creative input of the Owner, who attempts to match codes and competences with those of the User. These offerings are then sent back to the User, who, depending on how much s/he shares codes and competences, may agree or disagree with the choices on offer.

6. CONCLUSION

While the communication between Owners and Users has been shown to be reflexive and interactive, representing an interactive information retrieval system, the professionals in this sample were not schooled in searching through large collections. Their primary motivation seems to be to find the 'best' piece of music to communicate their meaning. Although there are some search engines available for this type of use, human involvement is central in the process. Attempting to build systems that meet the information needs of these users should be flexible, incorporate queries by example as well as by matching established metadata (including lyrics), and allow searching by content as well as contextual information. It is often the use that determines the choice, and it is rare that the initial query will name a song or an artist except as an example of what would work but is not available. The clearest briefs appear to be moving images, so matching video features to music features could be a worthwhile avenue of exploration. Using musical meaning to enhance moving images requires an understanding of the contexts and contents of both media and further user research is required to inform successful systems development if these needs are to be met.

7. FUTURE WORK

This preliminary analysis of interviews has been used to describe the process in choosing music for work purposes in the music and film industry. It is planned to analyse these texts more closely over time, to draw out nuances and detail of the discourses, and investigate the interpretive repertoires that are used by the participants in the process when talking about searching for and using music. This will provide a view of 'macrosociologically relevant cultural regularities' [28:26] which are used by participants and will inform a wide ranging research project on searching for and using music from large collections across the music industry, including games, broadcasters and DJs, which will also investigate and evaluate key systems to provide a holistic evaluation of searching for music for work purposes.

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USING AUDIO ANALYSIS AND NETWORK STRUCTURE TO IDENTIFY COMMUNITIES IN ON-LINE SOCIAL NETWORKS OF ARTISTS

ABSTRACT

Community detection methods from complex network theory are applied to a subset of the Myspace artist network to identify groups of similar artists. Methods based on the greedy optimization of modularity and random walks are used. In a second iteration, inter-artist audio-based similarity scores are used as input to enhance these community detection methods. The resulting community structures are evaluated using a collection of artist-assigned genre tags. Evidence suggesting the Myspace artist network structure is closely related to musical genre is presented and a Semantic Web service for accessing this structure is described.

1 INTRODUCTION

The dramatic increase in popularity of online social networking has led hundreds of millions of individuals to publish personal information on the Web. Music artists are no exception. Myspace [1] has become the de-facto standard for web-based music artist promotion. Although exact figures are not made public, recent blogosphere chatter suggests there are well over 7 million artist pages [2] on Myspace. Myspace artist pages typically include some streaming audio and a list of "friends" specifying social connections. This combination of media and a user-specified social network provides a unique data set that is unprecedented in both scope and scale.

However, the Myspace network is the result of hundreds of millions individuals interacting in a virtually unregulated fashion. Can this crowd-sourced tangle of social networking ties provide insights into the dynamics of popular music? Does the structure of the Myspace artist network have any relevance to music-related studies such as music recommendation or musicology?

In an effort to answer these questions, we identify communities of artists based on the Myspace network topology and attempt to relate these community structures to musical genre. To this end, we examine a sample of the Myspace social network of artists. First we review some previous work on the topics of artist networks, audio-based music analysis, and complex network community identification. We then describe our methodology including our network sampling method in Section 3.1 and our community detection approaches in Section 3.2. In Section 3.3 we describe the concept of *genre entropy* - a metric for evaluating the relevance of these community structures to music. Finally, we include a discussion of the results, suggestions for future work, and describe a Semantic Web service that can be used to access some of the data in a structured format.

[1] http://myspace.com

[2] http://scottelkin.com/archive/2007/05/11/Myspace-Statistics.aspx
∼25 million songs, ∼3.5 songs/artist, ∼7 million artists

2 BACKGROUND

2.1 Complex Networks

Complex network theory uses the tools of graph theory and statistical mechanics to deal with the structure of relationships in complex systems. A network is defined as a graph $G = (N, E)$ where N is a set of *nodes* connected by a set of *edges* E. We will refer to the number of nodes as n and the number of edges as m. The network can also be defined in terms of the *adjacency matrix* $G = A$ where the elements of A are

$$A_{ij} = \begin{cases} 1 & \text{if nodes } i \text{ and } j \text{ are connected,} \\ 0 & \text{otherwise.} \end{cases} \quad (1)$$

In this work, we restrict our analysis to the *undirected* case where edges are not considered directional and A is a symmetric matrix. For a summary of recent developments in complex networks see [7, 17].

2.2 Music Networks

Networks of musicians have been studied in the context of complex network theory - viewing the artists as nodes in the network and using either collaboration, influence, or some measure of similarity to define network edges [4, 5, 9, 19]. However the networks studied are generally constructed based on expert opinions (e.g. AllMusicGuide [3]) or proprietary algorithms based on user listening habits (e.g. Last.fm [4]). The Myspace artist network is unique in that the edges - the "friend" connections - are specified by the artists themselves. This makes the Myspace artist network a true social network. It has been shown that significantly different network topologies result from different approaches to artist

[3] http://www.allmusic.com/

[4] http://last.fm

network construction [4]. Since the Myspace artist network is of unique construction - owing its structure to the decisions and interactions of millions of individuals - we are motivated to analyze its topology and explore how this network structure relates to music.

It should be noted that networks of music listeners and bipartite networks of listeners and artists have also been studied [2, 13]. While such studies are highly interesting in the context of music recommendation, and while the Myspace network could potentially provide interesting data on networks of listeners, we restrict our current investigation to the Myspace artist network.

Previous analysis of the Myspace social network (including artists and non-artists) suggests that it conforms in many respects to the topologies commonly reported in social networks - having a power-law degree distribution and a small average distance between nodes [1]. Previous analysis of the Myspace *artist* network sample used in this work shows a multi-scaling degree distribution, a small average distance between nodes, and strong assortative mixing with respect to genre [11].

2.3 Community Detection

Recently, there has been a significant amount of interest in algorithms for detecting community structures in networks. These algorithms are meant to find dense subgraphs (communities) in a larger sparse graph. More formally, the goal is to find a partition $\mathcal{P} = \{C_1, \ldots, C_c\}$ of the nodes in graph G such that the proportion of edges inside C_k is high compared to the proportion of edges between C_k and other partitions.

Because our network sample is moderately large, we restrict our analysis to use more scalable community detection algorithms. We make use of the greedy modularity optimization algorithm [6] and the walktrap algorithm [20]. These algorithms are described in detail in Section 3.2.

2.4 Signal-based Music Analysis

A variety of methods have been developed for signal-based music analysis, characterizing a music signal by its timbre, harmony, rhythm, or structure. One of the most widely used methods is the application of Mel-frequency cepstral coefficients (MFCC) to the modeling of timbre [15]. In combination with various statistical techniques, MFCCs have been successfully applied to music similarity and genre classification tasks [18, 16, 3, 10]. A common approach for computing timbre-based similarity between two songs or collections of songs creates Gaussian mixtures models (GMM) describing the MFCCs and comparing the GMMs using a statistical distance measure. Often the earth mover's distance (EMD), a technique first used in computer vision, is the distance measure used for this purpose [21]. The EMD algorithm finds the minimum work required to transform one distribution into another. We use a set of inter-artist EMD values as a means of enhancing our community detection methods as described in Section 3.2.3.

3 METHODOLOGY

We will review our methodology beginning with a description of our network sampling method in Section 3.1. We then describe the various community detection approaches applied to the network in Section 3.2 and how we incorporate audio-based measures. Finally, we describe our metric for evaluating the relevance of the Myspace artist network structure with respect to musical genre in Section 3.3.

3.1 Sampling Myspace

The Myspace social network presents a variety of challenges. For one, the massive size prohibits analyzing the graph in its entirety, even when considering only the artist pages. Therefore we sample a small yet sufficiently large portion of the network as described in section 3.1.2. Also, the Myspace social network is filled with noisy data - plagued by spammers and orphaned accounts. We limit the scope of our sampling in a way that minimizes this noise.

3.1.1 Artist Pages

It is important to note we are only concerned with a subset of the Myspace social network - the Myspace *artist* network. Myspace artist pages are different from standard Myspace pages in that they include a distinct audio player application. We use the presence or absence of this player to determine whether or not a given page is an artist page.

A Myspace page will always include a top friends list. This is a hyperlinked list of other Myspace accounts explicitly specified by the user. The top friends list is limited in length with a maximum length of 40 friends (the default length is 16 friends). In constructing our sampled artist network, we use the top friends list to create a set of directed edges between artists. Only top friends who also have artist pages are added to the sampled network; standard Myspace pages are ignored. We also ignore the remainder of the friends list (i.e. friends that are not specified by the user as top friends), assuming these relationships are not as relevant. This reduces the amount of noise in the sampled network but also artificially limits the outdegree of each node. This approach is based on the assumption that artists specified as top friends have some meaningful musical connection for the user – whether through collaboration, stylistic similarity, friendship, or artistic influence.

Each Myspace artist page includes between zero and three genre tags. The artist selects from a list of 119 genres specified by Myspace. We include this information in our data

set.

The audio files associated with each artist page in the sampled network are also collected for feature extraction as described in Section 3.2.3.

3.1.2 Snowball Sampling

For the Myspace artist network, snowball sampling is the most appropriate method [1]. Alternative methods such as random edge sampling and random node sampling would result in many small disconnected components and not provide any insight to the structure of the entire network [14]. In snowball sampling, a first seed node (artist page) is included in the sample. Then the seed node's neighbors (top friends) are included in the sample. Then the neighbors' neighbors. This breadth-first sampling is continued until a particular sampling ratio is achieved. We randomly select one seed node [5] and perform 6 levels of sampling - such that in an undirected view of the network, no artist can have a geodesic distance greater than 6 with respect to the seed artist - to collect 15,478 nodes. If the size of the Myspace artist network is around 7 million, then this is close to the 0.25% sampling ratio suggested in [12].

3.1.3 Conversion to Undirected Graph

With the sampling method described above, the edges in our Myspace artist network are directional. If j is a top friend of i, this does not mean i is a top friend of j ($(i,j) \neq (j,i)$). However, many community detection algorithms operate on *undirected* graphs where $(i,j) = (j,i)$. For this reason we convert our directed graph to an undirected graph. Where a single directed edge exists it becomes undirected and where a reflexive pair of directed edges exist a single undirected edge replaces both edges. This process reduces the edge count from $120,487$ to $91,326$.

3.2 Community Detection

We apply two community detection algorithms to our network sample - the greedy optimization of modularity [6] and the walktrap algorithm [20]. Both of these algorithms are reasonably efficient and both algorithms can be easily adapted to incorporate audio-based similarity measures.

3.2.1 Greedy Modularity Optimization

Modularity is a network property that measures the appropriateness of a network division with respect to network structure. Modularity can be defined in several different ways [7]. In general, modularity Q is defined as the number of edges within communities minus the expected number of such edges. Let A_{ij} be an element of the network's adjacency matrix and suppose the nodes are divided into communities such that node i belongs to community C_i. We define modularity Q as the fraction of edges within communities minus the expected value of the same quantity for a random network. Then Q can be calculated as follows:

$$Q = \frac{1}{2m} \sum_{ij} \left[A_{ij} - \frac{d_i d_j}{2m} \right] \delta_{C_i C_j} \quad (2)$$

where the $\delta_{C_i C_j}$ function is 1 if $C_i = C_j$ and 0 otherwise, m is the number of edges in the graph, and d_i is the *degree* of node i - that is, the number of edges incident on node i. The sum of the term $\frac{d_i d_j}{2m}$ over all node pairs in a community represents the expected fraction of edges within that community in an equivalent random network where node degree values are preserved.

If we consider Q to be a benefit function we wish to maximize, we can then use an agglomerative approach to detect communities - starting with a community for each node such that the number of partitions $|\mathcal{P}| = n$ and building communities by amalgamation. The algorithm is greedy, finding the changes in Q that would result from the merge of each pair of communities, choosing the merge that results in the largest increase of Q, and then performing the corresponding community merge. It can be proven that if no community merge will increase Q the algorithm can be stopped because no further modularity optimization is possible [6]. Using efficient data structures based on sparse matrices, this algorithm can be performed in time $\mathcal{O}(m \log n)$.

3.2.2 Random Walk: Walktrap

The walktrap algorithm uses random walks on G to identify communities. Because communities are more densely connected, a random walk will tend to be 'trapped' inside a community - hence the name "walktrap".

At each time step in the random walk, the walker is at a node and moves to another node chosen randomly and uniformly from its neighbors. The sequence of visited nodes is a *Markov chain* where the states are the nodes of G. At each step the transition probability from node i to node j is $P_{ij} = \frac{A_{ij}}{d_i}$ which is an element of the transition matrix P for the random walk. We can also write $P = D^{-1}A$ where D is the diagonal matrix of the degrees ($\forall i, D_{ii} = d_i$ and $D_{ij} = 0$ where $i \neq j$).

The random walk process is driven by powers of P: the probability of going from i to j in a random walk of length t is $(P^t)_{ij}$ which we will denote simply as P^t_{ij}. All of the transition probabilities related to node i are contained in the i^{th} row of P^t denoted as $P^t_{i\bullet}$. We then define an inter-node distance measure:

$$r_{ij} = \sqrt{\sum_{k=1}^{n} \frac{(P^t_{ik} - P^t_{jk})^2}{d_k}} = \| D^{-\frac{1}{2}} P^t_{i\bullet} - D^{-\frac{1}{2}} P^t_{j\bullet} \| \quad (3)$$

[5] our randomly selected artist is French rapper Karna Zoo `http://www.myspace.com/karnazoo`

where $\|.\|$ is the Euclidean norm of $\Re^n$. This distance can also be generalized as a distance between communities: $r_{C_iC_j}$ or as a distance between a community and a node: r_{C_ij}.

We then use this distance measure in our algorithm. Again, the algorithm uses an agglomerative approach, beginning with one partition for each node ($|\mathcal{P}| = n$). We first compute the distances for all adjacent communities (or nodes in the first step). At each step k, two communities are chosen based on the minimization of the mean σ_k of the squared distances between each node and its community.

$$\sigma_k = \frac{1}{n} \sum_{C_i \in \mathcal{P}_k} \sum_{i \in C_i} r_{iC_i}^2 \tag{4}$$

Direct calculation of this quantity is known to be NP-hard, so instead we calculate the variations $\Delta\sigma_k$. Because the algorithm uses a Euclidean distance, we can efficiently calculate these variations as

$$\Delta\sigma(C_1, C_2) = \frac{1}{n} \frac{|C_1||C_2|}{|C_1| + |C_2|} r_{C_1C_2}^2 \tag{5}$$

The community merge that results in the lowest $\Delta\sigma$ is performed. We then update our transition probability matrix

$$P^t_{(C_1 \cup C_2)\bullet} = \frac{|C_1| P^t_{C_1\bullet} + |C_2| P^t_{C_2\bullet}}{|C_1| + |C_2|} \tag{6}$$

and repeat the process updating the values of r and $\Delta\sigma$ then performing the next merge. After $n-1$ steps, we get one partition that includes all the nodes of the network $\mathcal{P}_n = \{N\}$. The algorithm creates a sequence of partitions $(\mathcal{P}_k)_{1 \leq k \leq n}$. Finally, we use modularity to select the best partition of the network, calculating $Q_{\mathcal{P}_k}$ for each partition and selecting the partition that maximizes modularity.

Because the value of t is generally low (we use $t = 4$), this community detection algorithm is quite scalable. For most real-world networks, where the graph is sparse, this algorithm runs in time $\mathcal{O}(n^2 \log n)$ [20].

3.2.3 Audio-based Community Detection

Both algorithms described above are based on the adjacency matrix A of the graph. This allows us to easily extend these algorithms to include audio-based similarity measures. We simply insert an inter-node similarity value for each non-zero entry in A. We calculate these similarity values using audio-based analysis.

For the audio analysis, MFCCs are extracted resulting in 100ms non-overlapping frames. For each artist node a GMM is built from the concatenation of MFCC frames for all songs found on each artist's Myspace page (generally between 1 and 4 songs although some artists have more). For each non-zero value in the adjacency matrix A_{ij} a dissimilarity value is calculated using the earth mover's distance λ_{ij} between the GMMs corresponding to nodes i and j.

These dissimilarity values must be converted to similarity values to be successfully applied to the community detection algorithms. This is achieved by setting a maximum value for λ and subtracting each distance from the maximum value.

$$A_{ij} = \begin{cases} \lambda_{max} - \lambda_{ij} & \text{if nodes } i \text{ and } j \text{ are connected,} \\ 0 & \text{otherwise.} \end{cases} \tag{7}$$

Based on the distribution of λ for our data set we select $\lambda_{max} = 150$.

3.3 Genre Entropy

Now that we have several methods for detecting community structures in our network, we need a means of evaluating the relevance of these structures in the context of music. Traditionally, music and music artists are classified in terms of *genre*. If the structure of the Myspace artist network is relevant to music, we would expect the communities identified within the network to be correlated with musical genres. That is, communities should contain nodes with a more homogenous set of genre associations than the network as a whole.

As mentioned in Section 3.1, we have collected genre tags that are associated with each artist. In order to measure the diversity of each community with respect to genre we use a variant of Shannon entropy we call *genre entropy* S. This approach is similar to that of Lambiotte [13]. For a given community C_k we calculate genre entropy as:

$$S_{C_k} = -\sum_{\gamma \in C_k} P_{\gamma|C_k} \log P_{\gamma|C_k} \tag{8}$$

where $P_{\gamma|C_k}$ is the probability of finding genre tag γ in community C_k. As the diversity of genre tags in a community C_k increases, the genre entropy S_{C_k} increases. As the genre tags become more homogenous, the value of S_{C_k} decreases. If community C_k is described entirely by one genre tag then $S_{C_k} = 0$. We can calculate an overall genre entropy S_G by including the entire network sample. In this way, we can evaluate each community identified by comparing S_{C_k} to S_G. If the community structures in the network are related to musical genre, we would expect the communities to contain more homogenous mixtures of genre tags. That is, in general, we would expect $S_{C_k} \leq S_G$. However, as community size decreases so will the genre entropy because fewer tags are available. To account for this, we create a random partitioning of the graph that results in the same number of communities and calculate the corresponding genre entropies S_{rand} to provide a baseline.

If an artist specified no genre tags, this node is ignored and makes no contribution to the genre entropy score. In our data set, 2.6% of artists specified no genre tags.

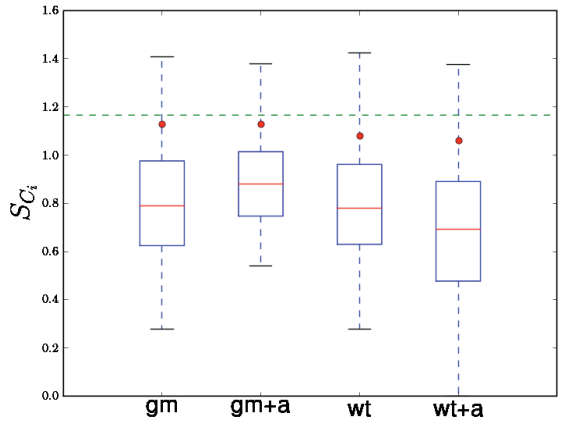

Figure 1. Box and whisker plot showing the spread of community genre entropies for each graph partition method where gm is greedy modularity, gm+a is greedy modularity with audio weights, wt is walktrap, and wt+a is walktrap with audio weights. The horizontal line represents the genre entropy of the entire sample. The circles represent the average value of genre entropy for a random partition of the network into an equivalent number of communities.

algorithm	c	$\langle S_C \rangle$	$\langle S_{rand} \rangle$	Q
none	1	1.16	-	-
gm	42	0.81	1.13	0.61
gm+a	33	0.90	1.13	0.64
wt	195	0.80	1.08	0.61
wt+a	271	0.70	1.06	0.62

Table 1. Results of the community detection algorithms where c is the number of communities detected, $\langle S_C \rangle$ is the average genre entropy for all communities, $\langle S_{rand} \rangle$ is the average genre entropy for a random partition of the network into an equal number of communities, and Q is the modularity for the given partition.

4 RESULTS

The results of the various community detection algorithms are summarized in Figure 1 and Table 1. When the genre entropies are averaged across all the detected communities, we see that for every community detection method the average genre entropy is lower than S_G as well as lower than the average genre entropy for a random partition of the graph into an equal number of communities. This is strong evidence that the community structure of the network *is* related to musical genre.

It should be noted that even a very simple examination of the genre distributions for the entire network sample suggests a network structure that is closely related to musical genre. Of all the genre associations collected for our data set, 50.3% of the tags were either "Hip-Hop" or "Rap" while 11.4% of tags were "R&B". Smaller informal network samples, independent of our main data set, were also dominated by a handful of similar genre tags (i.e. "Alternative", "Indie", "Punk"). In context, this suggests our sample was essentially "stuck" in a community of Myspace artists associated with these particular genre inclinations. However, it is possible that these genre distributions are indicative of the entire Myspace artist network. Regardless, given that the genre entropy of our entire set is so low to begin with it is an encouraging result that we could efficiently identify communities of artists with even lower genre entropies.

Without audio-based similarity weighting, the greedy modularity algorithm (gm) and the walktrap algorithm (wt) result in genre entropy distributions with no statistically significant differences. However the walktrap algorithm results in almost five times as many communities which we would expect to result in a lower genre entropies because of smaller community size. Also note that the optimized greedy modularity algorithm is considerably faster than the walktrap algorithm - $\mathcal{O}(m \log n)$ versus $\mathcal{O}(n^2 \log n)$.

With audio-based similarity weighting, we see mixed results. Applying audio weights to the greedy modularity algorithm (fg+a) actually increased genre entropies but the differences between fg and fg+a genre entropy distributions are not statistically significant. Audio-based weighting applied to the walktrap algorithm (wt+a) results in a statistically significant decrease in genre entropies compared to the un-weighted walktrap algorithm ($p = 4.2 \times 10^{-4}$). It should be noted that our approach to audio-based similarity results in dissimilarity measures that are mostly orthogonal to network structure [8]. Future work will include the application of different approaches to audio-based similarity.

5 MYSPACE AND THE SEMANTIC WEB

Since our results indicate that the Myspace artist network is of interest in the context of music-related studies, we have made an effort to convert this data to a more structured format. We have created a Web service [6] that describes any Myspace page in a machine-readable Semantic Web format. Using FOAF [7] and the Music Ontology [8], the service describes a Myspace page in XML RDF. This will allow future applications to easily make use of Myspace network data (i.e. for music recommendation).

[6] available at `http://dbtune.org/myspace`
[7] `http://www.foaf-project.org/`
[8] `http://musicontology.com/`

6 CONCLUSIONS

We have presented an analysis of the community structures found in a sample of the Myspace artist network and shown that these community structures are related to musical genre. We have applied two efficient algorithms to the task of partitioning the Myspace artist network sample into communities and we have shown how to include audio-based similarity measures in the community detection process. We have evaluated our results in terms of genre entropy - a measure of genre tag distributions - and shown the community structures in the Myspace artist network are related to musical genre.

In future work we plan to examine community detection methods that operate locally, without knowledge of the entire network. We also plan to address directed artist graph analysis, bipartite networks of artists and listeners, different audio analysis methods, and the application of these methods to music recommendation.

7 ACKNOWLEDGEMENTS

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KERNEL EXPANSION FOR ONLINE PREFERENCE TRACKING

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ABSTRACT

User preferences of music genres can significantly changes over time depending on fashions and the personal situation of music consumers. We propose a model to learn user preferences and their changes in an adaptive way. Our approach refines a model for user preferences by explicitly considering two plausible constraints of computational costs and limited storage space. The model is required to adapt itself to changing data distributions, and yet be able to compress "historical" data. We exploit the success of kernel SVM, and we consider an online expansion of the induced space as a preprocessing step to a simple linear online learner that updates with maximal agreement to previously seen data.

1 INTRODUCTION

Adaptivity is an indispensable prerequisite for personalization of acoustic devices like hearing aids. Hearing instruments, for instance, are often expected to suppress distracting or even annoying acoustic background sounds, whereas enjoyable acoustic scenes should be enhanced.

Such acoustic scenes can often be differentiated in a fairly fine-grained way, e.g. users might like some specific pop artists, but dislikes other pop entertainers. We currently live in a constantly changing world, and the amount of personalization that can be predicted and implemented in advance is quite limited. This pre-training of device parameters cannot cover all possible acoustic scenes (databases) to provide user specific settings. Furthermore, new acoustic scenes might evolve (e.g. new music types), and the user might change his preference over time. Hence, we need to look at real-time online adaptive algorithms.

Many classification results are based on batch learning, where the training set with complete or partial labeling is static and given at the training time. State-of-the-art methods for classification are Support Vector Machines (SVM) [1] which have been successfully employed in genre/artist classification [2] as well as in active learning [3]. A SVM classifier is a kernel method which finds an optimal separating hyperplane of the data points x in a high-dimensional projection space $\Phi(x)$, without having to explicitly expand the data representation in this high-dimensional space. This computational advantage is achieved via the kernel trick, i.e., the scalar product in this high dimensional space assumes a functional form defined by the kernel. The large margin discriminatory hyperplane in this high dimensional space accounts for the success of SVM in batch classifiers.

In our scenario we deal with the user's preference. Here, batch algorithms are no longer applicable since data are not available in advance, nor can the system select data points from a known database for querying (e.g. in active learning). Furthermore, temporal preference changes can not be modeled via batch learning, since the time information is discarded.

Hence we need to consider a sequential online learning paradigm. In sequential online learning [4], the data are processed as a stream, and the algorithm updates the classifier parameters online. Here, we use two online-learning algorithms: an incremental SVM (LASVM) [5] and online passive-aggressive algorithms (PA) [6]. Both algorithms are discriminative kernel algorithms that learn sequentially. A drawback of kernelized online large margin methods is the tendency to invoke more and more support vectors which inevitably will violate the constraints on computational costs and space.

To maintain the advantages of kernelization without the space/computational violations, we introduce a infinite memory extension of kernel machines (specifically to PA) which integrates the advantages of kernelization with the memory space limitations.

2 BACKGROUND

In sequential learning, the data is accessible at times $t = 1, ..., T$, where the horizon T can be arbitrarily large. At each time step t, a new observation x_t is presented to the system, which predicts a label $f_t(x_t)$. After this prediction, the correct label y_t is revealed to the system which uses this information to update the classifier f_{t+1}. The new classifier f_{t+1} is used for the next time step $t + 1$.

This paper presents an extension of the passive-aggressive algorithm (PA) [6] for online preference learning and it compares it with an incremental SVM learning system (LASVM) [5]. Both algorithms learn discriminative classifiers in a sequential manner, i.e., LASVM optimizes a quadratic cost

function, whereas PA implements gradient descent. Our variation of PA replaces the scalar product in the linear form of PA by a kernel expansion.

2.1 Incremental SVM

For comparison purposes, we consider an incremental version of SVM, the LASVM [5]. SVM optimization solves a quadratic programming problem for the dual objective function of large margin classifiers. One commonly used solver is the sequential minimal optimization (SMO) [7] which computes a τ-approximate solution. The LASVM implementation reorganizes the SMO sequential direction searches by considering pairs of examples (the new data point and existing support vectors (SVs)). In a second step, it repeats the search between pairs of SVs and eliminates redundant SVs.

We modified the code provided by the authors [1] to process the data in order of input. Simultaneously, we compute the cumulative error of the training in one epoch.

2.2 Online Passive-Aggressive Algorithms

Online passive-aggressive algorithms (PA) define a family of margin-based online learning algorithms [6] which are based on the hinge-loss for large margin classification

$$l(w;(x,y)) = \begin{cases} 0 & y(w \cdot x) \geq 1 \\ 1 - y(w \cdot x) & \text{otherwise.} \end{cases} \quad (1)$$

This constrained optimization is formulated as

$$w_{t+1} = \arg \min_{w \in \mathbb{R}^D} \frac{1}{2}||w - w_t||^2 \text{ s.t. } l(w;(x_t,y_t)) = 0. \quad (2)$$

This procedure can be interpreted as follows: At each update step, if the current hyperplane misclassifies the current sample x (within a margin), then the hyperplane is adjusted to the next nearest hyperplane that resolves this problem. The nearest hyperplane is the one that deviates the least from the original hyperplane.

Algorithm 1 PA

Require: Input: aggressive parameter $C > 0$, initial classifier w_1

1: **for** t = 1, 2, ... **do**
2: Predict: $\hat{y}_t = \text{sign}(w_t' x_t)$ for current data x_t
3: Obtain y_t: loss $l_t = \max\{0, 1 - y_t(w_t' x_t)\}$
4: Update:

$$\begin{aligned} \alpha_t &= \min\left\{C, \frac{l_t}{||x_t||^2}\right\} \\ w_{t+1} &= w_t + \alpha_t y_t x_t \end{aligned}$$

5: **end for**

[1] http://leon.bottou.org/projects/lasvm/

The PA algorithm is outlined above. Its kernel counterpart is obtained by replacing the scalar product with a kernel function. The update step

$$w_{t+1} = w_t + \alpha_t y_t x_t \quad (3)$$

of the classifier function is replaced by

$$f_{t+1}(x) = \sum_{i=1}^{t} \alpha_i y_i K(x_i, x). \quad (4)$$

Comparing the Eq 3 and Eq 4, we see that the kernel formulation requires an explicit storage of the data points with $\alpha_i \neq 0$ ("SVs"). These SVs are accumulated during the learning phase and their number may grow arbitrarily large. Apart from violating storage constraints, a large number of SVs also leads to an increased computational cost for the prediction step, when the kernel function has to be evaluated for each of the many "SVs". Contrary to this growth of the computational complexity for kernel machines, the linear support vector machines (Eq 3) implicitly encode the SVs with the normal vector w_{t+1}, which leaves the storage requirements and the computational costs unchanged during update.

2.3 Limiting the Kernel PA

When the cache for the kernel storage is limited, two alternatives are available for discarding exemplars: removal of the oldest "SV", or removal of the worst performing "SV". By removing the oldest sample, we penalize the system where memory is needed. If the data shows cyclical behavior, then the "SVs" used to model a certain phase might have been discarded by the time this phase is revisited. In expulsion of the worst performing "SV", we discard "SVs", as advocated in [8]. If a loss is incurred, we find the "SV" for which the removal decreases the loss maximally:

$$\begin{aligned} &\arg\max_j y_t(f_t(x_t) - \alpha_j y_j K(x_j, x_t)) = \\ &\arg\min_j y_t \alpha_j y_j K(x_j, x_t) \end{aligned} \quad (5)$$

3 KERNEL EXPANSION FOR THE PASSIVE AGGRESSIVE ALGORITHM

Mercer's theorem states that a continuous, symmetric, positive semi-definite kernel function represents a scalar product in an appropriately high-dimensional space. Via the kernel trick [9], observations x_i of a non-linearly separable problem are mapped to a high-dimensional space $\Phi(x_i)$, such that these data becomes linearly separable. The scalar product in the linear function is replaced with the kernel function, such that it is not necessary to explicitly compute $\Phi(x)$:

$$K(x,z) = \Phi(x) \cdot \Phi(z). \quad (6)$$

Two commonly used kernels are the radial basis function (RBF) kernel

$$K(x,z) = e^{-\gamma||x-z||^2}, \quad (7)$$

where γ is the kernel width, and the polynomial kernel

$$K(x,z) = (\gamma x'z + c)^{\delta}, \quad (8)$$

for some scale factor γ, bias c and degree δ. For instance, the polynomial kernel emulates the scalar product of a finite induced feature space. Consider the polynomial kernel of degree 2, $K(x,z) = (x \cdot z')^2$. We obtain

$$\Phi(x) = (x_1^2, ..., x_D^2, \sqrt{2}x_1x_2, ..., \sqrt{2}x_dx_{d+j}, ..., \sqrt{2}x_{D-1}x_D) \quad (9)$$

where $d = 1, ..., D$, and $j = 1, ..., D-d$ indexes an exhaustive pairing of all features. This transformation yields a $D(D+1)/2$ dimensional space, i.e. $\Phi(x) \in \mathbb{R}^{D(D+1)/2}$.

We can now modify the algorithm PA to derive the algorithm passive-aggressive linear (expanded) PA-L-EX (Table3) which requires an additional input: function Φ which maps the input data to a different (higher) dimensional space induced by the kernel. Whilst this high dimensional representation is infeasible for batch-learning algorithms (dataset representation has to be stored for ALL data points which may be infeasible for large datasets), it is readily integrated in the online scenario. Since the data points are processed individually, this high dimensional expansion is calculated on the fly and, therefore, it is practical. In general, this extension holds for any kernel with a finite expansion for the induced feature space, or which admits some approximation of this induced feature space.

Algorithm 2 PA-L-EX

Require: Input: aggressive parameter $C > 0$, Expansion function $\Phi : \mathbb{R}^D \rightarrow \mathbb{R}^M, M \gg D$, initial classifier $w_1 \in \mathbb{R}^M$

1: **for** t = 1, 2, ... **do**
2: Expand data $x_t \in \mathbb{R}^D : \Phi(x_t) \in \mathbb{R}^M$
3: Predict: $\hat{y}_t = \text{sign}(w_t'\Phi(x_t))$ for current data x_t
4: Obtain y_t: loss $l_t = \max\{0, 1 - y_t(w_t'\Phi(x_t))\}$
5: Update:

$$\alpha_t = \min\left\{C, \frac{l_t}{||\Phi(x_t)||^2}\right\}$$
$$w_{t+1} = w_t + \alpha_t y_t \Phi(x_t)$$

6: **end for**

4 EXPERIMENTAL SETUP

Our experiments use a subset of the uspop2002 [2] dataset. This subset of 18 pop artists 1 each with 5 or 6 albums is described in [3]. We use these authors' training and test set partition definition, which takes into account the producer's effect. The validation set has not been used during training. In one experiment, we combined the validation and training set to created a large 2-phase-training set to simulate user-preference change. Table 2 shows the statistics for the various database partitions.

Rolling Stones	Aerosmith	Beatles
Dave Matthews Band	Metallica	Queen
Fleetwood Mac	Garth Brooks	Madonna
Depeche Mode	Green Day	Roxette
Bryan Adams	Pink Floyd	Genesis
Creedence Clearwater Revival	Tina Turner	U2

Table 1. Artists in the 18-artist subset of uspop2002.

Set	Artist	Album	Song
`Train`	18	3×18	650
`Test`	18	2×18	448
`Valid`	9	1×9	102
`Phase1` $\subset$ Train	18	2×18	431
`Phase2` $=$ Valid $\cup$ Train$\setminus$Phase1	18	$2 \times 9 + 1 \times 9$	321

Table 2. Number of artists, albums and songs covered in each subset of the dataset. `Phase1` and `Phase2` data is a repartition of the combined `Train` and `Valid` sets.

4.1 Feature Extraction

We employed the 20 dimensional MFCC features distributed with the database (some songs were corrupt and discarded from the database), and we generated song level features. This grouping is achieved by vectorizing the sufficient statistics of the MFCC features per song, yielding a 230 dimensional input feature vector per song. We normalized [3] each feature dimension using the training data. High level features were not considered but can be readily integrated.

4.2 Verification and Simulation

To verify our dataset, we repeated the artist classification task outlined in [3] and we obtained the same accuracy rates.

For our experiments, we simulated random user profiles by splitting the 18 artists into two categories (like/dislike). The user likes a random subset of 3 to 15 artists, and dislikes the remaining artists. In this manner, we cover both balanced (same number of artists in each class) and unbalanced (one class having more artists) scenarios. For each

[2] http://labrosa.ee.columbia.edu/projects/musicsim/uspop2002.html

[3] For real-life scenario, approximate normalization factors can be estimated for a small dataset.

Kernel	$K(x,z)$	Test Acc
Linear	$x'z$	77.8 ± 6.6
Poly2	$(x'z)^2$	81.4 ± 5.2
RBF	$e^{-0.01*\|\|x-z\|\|^2}$	82.7 ± 4.7

Table 3. Results for batch SVM models. Hyper-parameters are optimized.

profile, we run 10 fold cross validation by using 10 permutations of the order in which the data is presented to the user. While the assumption of random selection might not correctly model the preferences of most users (preferences may occur in more structured groupings, e.g. mood or style), the random choice is certainly a difficult setting, since the classifier might be required to learn potential quirks.

4.3 Evaluation Measure

We consider two evaluation measure. For testing, we report accuracy rates $\frac{1}{n}\sum_{i=1}^{n}\mathbf{1}(f(x_i), y_i) * 100\%$ where $\mathbf{1}$ denotes the indicator function. For online learning scenarios, we report the cumulative accuracy

$$CumAcc = \frac{1}{T}\sum_{t=1}^{T}\mathbf{1}(f_t(x_{t-1}), y_t) * 100\%, \qquad (10)$$

which is the accuracy during the learning phase. The incurred error $(100\% - CumAcc)$ indicates how often the user was required to correct the system by responding with the correct label.

5 EXPERIMENTAL RESULTS

We first consider static random user preferences by simulating 100 randomly sampled user profiles. The results for batch learning algorithms (Sec 5.1) are analyzed for comparison purposes, then we consider sequential learning with independent and identically-distributed (i.i.d.) drawing of the data (Sec 5.2). In Sec 5.3 we further analyze the situation when the data are no longer presented randomly, rather, they occur in cycles. Here, the simulations considered balanced preferences (liking 50% of the artists). Finally, in Sec 5.4 the users change their preferences after an initial training phase. During the second training phase some preferences are flipped, and the classifier should adapt for such opinion changes.

5.1 Batch Conditions

We first test the "best" case when all training data are known in advance. For this batch learning, we use a SVM classifier using libsvm [4]. Comparing the different kernel types, the linear kernel performs poorly, whereas the RBF kernel shows the best performance, as can be seen in Table 3. The less frequently evaluated polynomial kernel, here with degree 2, almost matches the performance of RBF kernels.

Observing the models, we note that a large proportion of the training data was almost always retained within the model as support vectors, with almost all SVs located within the margin. For the RBF model, this $84.5 \pm 6.5\%$ of the training data were retained as SVs. This high proportion of SVs can be visualized as in Figure 1. New subcategories of data mark out new regions. As the amount of training data increases, it is possible that this growth in number of SV will level off. However, it is likely that new preferences (sub-categories) will require more SVs. Hence, for a fixed, batch training scenario, SVM may perform well, but for an equivalent online scenario, the algorithm may face resource problems such as infinitely large storage of SVs.

[4] http://www.csie.ntu.edu.tw/cjlin/libsvm

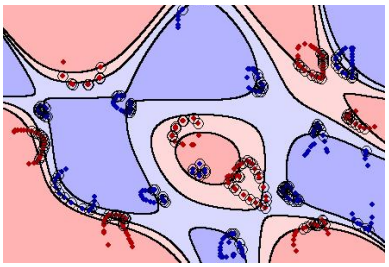

Figure 1. Illustration of SVM boundaries for RBF kernels. Circled data points are SVs. A high proportion of the observations serve as SVs.

Model	Kernel, Details	CumAcc	Test Acc
PA-L	Linear, D=230	75.0 ± 3.9	77.5 ± 5.7
PA-L-EX	Linear, Expand	80.8 ± 5.9	81.8 ± 5.7
PA-R200	RBF, 200 SV	77.5 ± 6.8	76.6 ± 7.6
PA-R	RBF, ∞ SV	81.6 ± 5.2	83.3 ± 5.1
LASVM	RBF, ∞ SV	80.8 ± 4.3	81.7 ± 4.6

Table 4. Descriptions of different incremental models and their performances during training (CumAcc) and for a hold out set (Test Acc). Sequential online learning on i.i.d. data. First three rows implement limited memory space. The last two rows are not truly online learners since no memory limitations is imposed on them.

5.2 Sequential i.i.d. Conditions

To test the online learning under sequential i.i.d. conditions, we present the training data in a randomly permuted order. To ensure strict comparison, permutations were preserved across all systems, and 10 permutations were simulated for each different random preference.

Table 4 shows the descriptions and results for the various

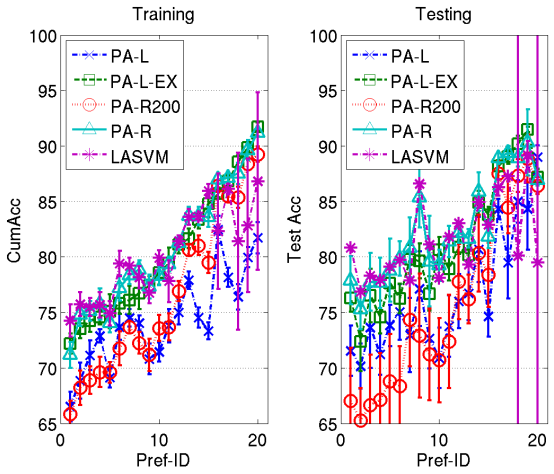

Figure 2. Detailed results on 20 randomly selected user profiles. Left: Train cumulative results. Right: Test accuracy on hold-out test set. Models are explained in Table 4.

models tested. The first three models are online sequential models which operate on a limited memory-storage, while the last two models (with ∞ SV) assume infinite storage space, and are not practical, and serve as a reference.

Comparing the performance sequential algorithms with that of the batch SVM classifiers (Table 3), we see that the performances are similar. LASVM attains a 81.7% performance which is 1.0% poorer than that of the SVM-RBF counterpart. PA-R even shows better performance at 83.3%. Similarly for the linear classifiers, PA-L performs 0.3% worse than that of the linear SVM.

While RBF-based algorithms are superior over the linear models for batch models, and also under impractical conditions of infinite memory, we see that this advantage no longer holds when memory is limited. Limiting the number of SV to 200SV (30% of the training set) as in PA-R200, we see a significant drop in performance, Instead, we observe that our approach of space expansion (PA-L-EX) combined with a linear classifier achieves similar performance levels to the kernel based classifiers with infinite storage.

Figure 2 demonstrates detailed results for 20 of the 100 random user profiles, both during training and on the hold-out test set. Note that while LASVM show slightly better results in the more difficult cases where accuracy is lower, it starts to show instable behavior when the sets are easy, resulting in large error bars. A detailed analysis shows that LASVM's stability depends on the sequence in which the data is presented. On the other hand, PA-algorithms are more stable for the same "unfavorable" sequences.

5.3 Sequential Non-i.i.d. Conditions

For common real world scenarios, data are no longer generated under i.i.d. conditions, rather, they show some patterns.

Model	Train CumAcc	Test Acc
PA-L	84.0 ± 1.6	65.7 ± 4.3
PA-L-EX	76.5 ± 1.5	73.1 ± 2.4
PA-R200	73.3 ± 2.6	59.1 ± 4.4
PA-R	79.7 ± 1.8	73.4 ± 4.2
LASVM	78.5 ± 2.0	78.5 ± 1.7
SVM (rbf)	batch	78.7 ± 1.7

Table 5. Performance when learning occur in an ordered manner (non-i.i.d.). Classes are balanced.

Artist ID	Phase 1	Phase 2	Test
a_1	likes	likes	likes
a_2	likes	likes	likes
a_3	likes	**dislikes**	**dislikes**
a_4	dislikes	**likes**	**likes**
a_5	dislikes	**likes**	**likes**
a_6	dislikes	dislikes	dislikes

Table 6. Simulation example for user preference change

For example, Christmas songs are usually popular only over Christmas, or some artist are more frequently played when they have released a new album. We have a learning phase that records the behavior of the user over a cycle, and hope that this preference is still represented during the evaluation cycle, where user feedback is no longer provided.

In this experiment, we have generated 10 random user profiles. Each user likes a random subset of 9 artists, resulting in a balanced class problem. The data is presented in phases, at each phase, only data from one "like" artist and one "dislike" artist is exhaustively presented, yielding a total of 9 phases during the training simulation.

Table 5 summarizes the results. Again, we show results for two online systems PA-L-EX and PA-R200. Results are also reported for PA-R and LASVM, but since they require infinite memory, they are impractical and serve as bounds.

Comparing the PA-L-EX and PA-R200 (limited storage), we see that the kernelized version does not perform well when data from previous phases are presented to the system during the evaluation phase, resulting in a low accuracy of 59.1%. PA-L-EX, on the other hand, still retains good classification accuracy, performing slightly worse at 73.1%. However, this almost equals the performance of a PA-RBF with infinite memory. We see that LASVM performs better, but is not feasible due to its requirement of large SV storage.

5.4 Adaptation to Preference Change over Time

In this experiment, we simulate a very adverse scenario. There are two phases during learning. During phase one, data from the set Phase1 are streamed i.i.d. to the system,

Model	Train CumAcc	Test Acc
PA-L-EX	75.8 ± 3.3	68.9 ± 3.6
LASVM	71.5 ± 1.6	61.1 ± 4.7

Table 7. Simulation for adverse (50%) preference change.

with the same 100 user profiles generated in the first experiment (i.i.d. data). Upon completion, data from set Phase2 are streamed with a 50% preference change. The 9 artists with 2 albums in the set Phase2 now have their labels (user preference) flipped. At the end of phase two, we test the final model on the test set, where the artist preference reflects that of phase two. Table 6 shows an example.

The performance of the two models LASVM and PA-L-EX are presented in Table 7. Recall that LASVM stores all the SVs that it accumulates. Its learning strength is greatly undermined by the change in user preference, which lead to conflicting SVs that greatly weakens the performance. As such, it breaks down and shows poor behavior (61.1% test accuracy). Even during the learning phase, it is already experiencing difficulties, degrading significantly from previous scenarios. PA-L-EX tries to accommodate for the new data by adjusting the hyperplane while trying to compromise for the previously learnt information. It shows weakened performance, but still retains information at 68.9%. This simulation models a severe 50% change in the classification classes, which takes its toll. Nevertheless, we notice that PA-L-EX does not break down which is in contrast to the complete failure of LASVM.

6 CONCLUSION

We have presented a user preference tracking system that exploits the advantages of using kernels for learning, without having to store potentially infinite numbers of support vectors. Our preference tracking system uses a simple algorithm PA-L-EX which can be efficiently implemented into small devices with limited storage and computational power. Despite these constraints, the algorithm has shown to be capable of recalling historically learnt events, and furthermore, adapt well to changes in concepts. These are necessary ingredients in adjusting and learning user preferences over time.

Our focus was primarily on song-level features, based on low level MFCC features. We experimented with 30 second music segments, which had slightly degraded performance (76.9% versus 81.8% test accuracy for i.i.d. case). Nevertheless, the results are encouraging given the simplicity of the features. There is potential for this algorithm to be used in combination with more sophisticated higher level features, and also online features.

We have addressed the case where the system learns and adjusts to the true label after each prediction. Future work could encompass the consideration of stream-based active learning ([10], [11]) where the data arrives in a stream, but labels must be procured. Other scenarios can include sparse feedback and inconsistent feedback.

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A REAL-TIME EQUALIZER OF HARMONIC AND PERCUSSIVE COMPONENTS IN MUSIC SIGNALS

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ABSTRACT

In this paper, we present a real-time equalizer to control a volume balance of harmonic and percussive components in music signals without a priori knowledge of scores or included instruments. The harmonic and percussive components of music signals have much different structures in the power spectrogram domain, the former is horizontal, while the latter is vertical. Exploiting the anisotropy, our methods separate input music signals into them based on the MAP estimation framework. We derive two kind of algorithm based on a I-divergence-based mixing model and a hard mixing model. Although they include iterative update equations, we realized the real-time processing by a sliding analysis technique. The separated harmonic and percussive components are finally remixed in an arbitrary volume balance and played. We show the prototype system implemented on Windows environment.

1 INTRODUCTION

A graphic equalizer is one of the most popular tools on an audio player, which allows an user to control the volume balance between frequency bands as its preference by separating an input audio signal by several band-pass filters and remixing them with different gains. Recently, based on other kinds of separation, more advanced audio equalizations have been discussed and developed [1, 2, 3], which increase the variety of modifying audio sounds and enrich functions of audio players.

In this paper, focusing on two different components included in music signals: harmonic and percussive ones, we present a technique to equalize them in real-time without a priori knowledge of the scores or the included instruments. Not only as an extended audio equalizer, the technique should yield the useful pre-processing for various tasks related to music information retrieval from audio signals [4]. It can suppress percussive tones, which often interfere multipitch analysis, while, suppression of harmonic component will facilitate drum detection or rhythm analysis. We have currently applied this technique to automatic chord detection based on emphasized chroma features [5], rhythm pattern extraction and rhythm structure analysis [6], and melody extraction.

For independently equalizing the harmonic and percussive components, it is required to separate them. This kind of separation problem has been widely discussed in the literature. Uhle *et al*. applied Independent Component Analysis (ICA) to the magnitude spectrogram, and classified the extracted independent components into a harmonic and a percussive groups based on the several features like percussiveness, noise-likeness, etc [7]. Helen *et al*. utilized Nonnegative Matrix Factorization (NMF) for decomposing the spectrogram into elementary patterns and classified them by pre-trained Support Vector Machine (SVM) [8]. Through modeling harmonic and inharmonic tones on spectrogram, Itoyama *et al*. aimed to an instrument equalizer and proposed separation of an audio signal to each track based on the MIDI information synchronized to the input audio signal [1].

The contribution of this paper is to derive a simple and real-time algorithm specifically for the harmonic/percussive separation without any pre-learning or a priori knowledge of score or included instruments of the input audio signals. We present the formulation of the separation in Maximum A Priori (MAP) estimation framework, derive the fast iterative solution to it by auxiliary function approach, implement it with sliding update technique for real-time processing, and examine the performance by experiments to popular and jazz music songs.

2 FORMULATION OF HARMONIC/PERCUSSIVE SEPARATION

2.1 MAP Estimation Approach

Let $F_{\omega,\tau}$ be a Short Time Fourier Transform (STFT) of a monaural audio signal $f(t)$, and $W_{\omega,\tau} = |F_{\omega,\tau}|^2$ be a short time power spectrum, where ω and τ represent frequency and time bins. Let $H_{\omega,\tau}$ and $P_{\omega,\tau}$ be a harmonic and a percussive component of $W_{\omega,\tau}$, respectively. The variables $\boldsymbol{W}$, $\boldsymbol{H}$, and $\boldsymbol{P}$ denote a set of $W_{\omega,\tau}$, $H_{\omega,\tau}$, and $P_{\omega,\tau}$, respectively.

The separation of W into H and P is a kind of underdetermined blind source separation problem. One way to mathematically formulate this kind of problems is putting it on MAP (Maximum A Posteriori) estimation framework through representing desired source properties as a priori probabilities. Assuming that H and P are independent, the objective function of MAP estimation in our problem can be written as

$$\begin{aligned} & J(\boldsymbol{H}, \boldsymbol{P}) \\ = \; & \log p(\boldsymbol{H}, \boldsymbol{P}|\boldsymbol{W}) \\ = \; & \log p(\boldsymbol{W}|\boldsymbol{H}, \boldsymbol{P}) + \log p(\boldsymbol{H}, \boldsymbol{P}) + C \\ = \; & \log p(\boldsymbol{W}|\boldsymbol{H}, \boldsymbol{P}) + \log p(\boldsymbol{H}) + \log p(\boldsymbol{P}) + C, \end{aligned} \quad (1)$$

where the first term represents the log-likelihood, the second and the third terms represent the prior probabilities, and C is a constant term not including H and P, hereafter, we will omit it since it is not used for MAP estimation.

A harmonic component on the spectrogram usually has a stable pitch and form parallel ridges with smooth temporal envelopes, while the energy of a percussive tone is concentrated in a short time frame, which forms a vertical ridge with wideband spectral envelopes. Then typically, the vertical and horizontal structure emerges in the spectrogram of audio signals shown in the top of Fig. 3.

Focusing on the horizontal and vertical smoothed envelope of $H_{\omega,\tau}$ and $P_{\omega,\tau}$, we model their a priori distribution as functions of spectrogram gradients as:

$$p(\boldsymbol{H}) \propto \prod_{\omega,\tau} \frac{1}{\sqrt{2\pi}\sigma_H} \exp\left(-\frac{(H^{\gamma}_{\omega,\tau-1} - H^{\gamma}_{\omega,\tau})^2}{2\sigma_H^2}\right), \quad (2)$$

$$p(\boldsymbol{P}) \propto \prod_{\omega,\tau} \frac{1}{\sqrt{2\pi}\sigma_P} \exp\left(-\frac{(P^{\gamma}_{\omega-1,\tau} - P^{\gamma}_{\omega,\tau})^2}{2\sigma_P^2}\right), \quad (3)$$

where σ_H^2 and σ_P^2 are the variance of the spectrogram gradients, probably depending on the frame length or frame shift of STFT, and γ represents a range-compression factor such that $(0 < \gamma \leq 1)$, which we introduced for increasing the degree of freedom of our model with holding the assumption of the Gaussian distribution.

2.2 Method 1: I-divergence-based mixing model

Although $H_{\omega,\tau}$ and $P_{\omega,\tau}$ are the power spectrograms the additivity of them is not rigorously hold, $H_{\omega,\tau} + P_{\omega,\tau}$ should be close to the observation $W_{\omega,\tau}$. In several power-spectrogram-based signal processing methods NMF [9, 10, 11], the distance between power spectrograms $A_{\omega,\tau}$ and $B_{\omega,\tau}$ can be measured by I-divergence:

$$I(\boldsymbol{A}, \boldsymbol{B}) = \sum_{\omega,\tau} \left(A_{\omega,\tau} \log \frac{A_{\omega,\tau}}{B_{\omega,\tau}} - A_{\omega,\tau} + B_{\omega,\tau}\right), \quad (4)$$

which is almost equivalent to the assumption that $p(\boldsymbol{W}|\boldsymbol{H}, \boldsymbol{P})$ is Poisson distribution [11]. Assuming that observation at each time-frequency is independent, the log-likelihood term can be written as

$$\begin{aligned} & \log p(\boldsymbol{W}|\boldsymbol{H}, \boldsymbol{P}) - C \\ = \; & -\sum_{\omega,\tau} \left\{ W_{\omega,\tau} \log \frac{W_{\omega,\tau}}{H_{\omega,\tau} + P_{\omega,\tau}} - W_{\omega,\tau} + H_{\omega,\tau} + P_{\omega,\tau} \right\}, \end{aligned} \quad (5)$$

where C is a constant term for normalization.

In the MAP estimation, the balance between a log-likelihood term and a prior distribution term is significant. Specifically in our problem, the relationship between them should be invariant for scaling. The property is satisfied by setting the range-compression factor as $\gamma = 0.5$. Then, the objective function can be written as

$$\begin{aligned} & J_1(\boldsymbol{H}, \boldsymbol{P}) \\ = \; & -\sum_{\omega,\tau} \left\{ W_{\omega,\tau} \log \frac{W_{\omega,\tau}}{H_{\omega,\tau} + P_{\omega,\tau}} - W_{\omega,\tau} + H_{\omega,\tau} + P_{\omega,\tau} \right\} \\ & -\frac{1}{\sigma_H^2}(\sqrt{H_{\omega,\tau-1}} - \sqrt{H_{\omega,\tau}})^2 \\ & -\frac{1}{\sigma_P^2}(\sqrt{P_{\omega-1,\tau}} - \sqrt{P_{\omega,\tau}})^2). \end{aligned} \quad (6)$$

Note that, when $H_{\omega,\tau}$, $P_{\omega,\tau}$, and $W_{\omega,\tau}$ are multiplied by a scale parameter A, the objective function is also just multiplied by A and the function form is invariant.

2.3 Method 2: hard mixing model

Since the intersection of the horizontal and vertical ridges is small, we can make a more strong assumption that they are approximately disjoint. In the case, $W_{\omega,\tau} = H_{\omega,\tau}$ or $W_{\omega,\tau} = P_{\omega,\tau}$ are exclusively satisfied at each (ω, τ). However, the sparse mixing model leads us to a large number of combination problem. For avoiding it and obtaining an approximative solution, we cast it to a hard mixing model on the range-compressed power spectrum as

$$\hat{W}_{\omega,\tau} = \hat{H}_{\omega,\tau} + \hat{P}_{\omega,\tau}, \quad (7)$$

where

$$\hat{W}_{\omega,\tau} = W^{\gamma}_{\omega,\tau}, \;\; \hat{H}_{\omega,\tau} = H^{\gamma}_{\omega,\tau}, \;\; \hat{P}_{\omega,\tau} = P^{\gamma}_{\omega,\tau}. \quad (8)$$

Eq. (7) is hold if $H_{\omega,\tau}$ and $P_{\omega,\tau}$ are actually disjoint. Although the model is rough, this assumption leads us to simple formulation and solution. Since the deterministic mixing model of eq. (7) vanishes the log-likelihood term, the objective function is given by

$$\begin{aligned} J_2(\hat{\boldsymbol{H}}, \hat{\boldsymbol{P}}) \; = \; & -\frac{1}{2\sigma_H^2} \sum_{\omega,\tau} (\hat{H}_{\omega,\tau-1} - \hat{H}_{\omega,\tau})^2 \\ & -\frac{1}{2\sigma_P^2} \sum_{\omega,\tau} (\hat{P}_{\omega-1,\tau} - \hat{P}_{\omega,\tau})^2, \end{aligned} \quad (9)$$

under the constraint of eq. (7).

3 DERIVATION OF UPDATE EQUATIONS THROUGH AUXILIARY FUNCTION

3.1 Method 1

Maximizing eq. (6) is a nonlinear optimization problem. In order to derive an effective iterative algorithm, we introduce an auxiliary function approach, which has been recently utilized in several signal processing techniques such as NMF [9] and HTC (Harmonic Temporal Clustering) [10].

Note that the following auxiliary function:

$$\begin{aligned} & Q_1(\boldsymbol{H}, \boldsymbol{P}, \boldsymbol{m_P}, \boldsymbol{m_H}) \\ = & -\sum_{\omega,\tau} m_{P\omega,\tau} W_{\omega,\tau} \log\left(\frac{m_{P\omega,\tau} W_{\omega,\tau}}{P_{\omega,\tau}}\right) \\ & -\sum_{\omega,\tau} m_{H\omega,\tau} W_{\omega,\tau} \log\left(\frac{m_{H\omega,\tau} W_{\omega,\tau}}{H_{\omega,\tau}}\right) \\ & -\frac{1}{\sigma_H^2}(\sqrt{H_{\omega,\tau-1}} - \sqrt{H_{\omega,\tau}})^2 \\ & -\frac{1}{\sigma_P^2}(\sqrt{P_{\omega-1,\tau}} - \sqrt{P_{\omega,\tau}})^2) \end{aligned} \quad (10)$$

holds

$$J_1(\boldsymbol{H}, \boldsymbol{P}) \geq Q_1(\boldsymbol{H}, \boldsymbol{P}, \boldsymbol{m_P}, \boldsymbol{m_H}), \quad (11)$$

for any $\boldsymbol{H}$, $\boldsymbol{P}$, $\boldsymbol{m_P}$, and $\boldsymbol{m_H}$ under the condition that

$$m_{P\omega,\tau} + m_{H\omega,\tau} = 1, \quad (12)$$

where $m_{P\omega,\tau}$ and $m_{H\omega,\tau}$ are auxiliary variables and $\boldsymbol{m_P}$ and $\boldsymbol{m_H}$ are sets of $m_{P\omega,\tau}$ and $m_{H\omega,\tau}$, respectively. The equality of eq. (10) is satisfied for

$$m_{X\omega,\tau} = \frac{X_{\omega,\tau}}{H_{\omega,\tau} + P_{\omega,\tau}}, \quad (13)$$

for $X = H$ or $X = P$. Then, updating $\boldsymbol{m_H}$ and $\boldsymbol{m_P}$ by eq. (13) increases the auxiliary function Q_1 and it achieves to J. After that, updating $\boldsymbol{H}$ and $\boldsymbol{P}$ by solving $\partial Q_1/\partial P_{\omega,\tau} = 0$ and $\partial Q_1/\partial H_{\omega,\tau} = 0$ increases Q_1 again and J_1 increases together because of the inequality of eq. (10). Hence, the iterations increases J_1 monotonically.

From $\partial Q_1/\partial P_{\omega,\tau} = 0$, $\partial Q_1/\partial H_{\omega,\tau} = 0$, and eq. (13), we have the following update equations:

$$H_{\omega,\tau} \leftarrow \left(\frac{b_{H\omega,\tau} + \sqrt{b_{H\omega,\tau}^2 + 4a_{H\omega,\tau}c_{H\omega,\tau}}}{2a_{H\omega,\tau}}\right)^2 \quad (14)$$

$$P_{\omega,\tau} \leftarrow \left(\frac{b_{P\omega,\tau} + \sqrt{b_{P\omega,\tau}^2 + 4a_{P\omega,\tau}c_{P\omega,\tau}}}{2a_{P\omega,\tau}}\right)^2 \quad (15)$$

$$m_{H\omega,\tau} \leftarrow \frac{H_{\omega,\tau}}{H_{\omega,\tau} + P_{\omega,\tau}} \quad (16)$$

$$m_{P\omega,\tau} \leftarrow \frac{P_{\omega,\tau}}{H_{\omega,\tau} + P_{\omega,\tau}} \quad (17)$$

where

$$a_{H\omega,\tau} = \frac{2}{\sigma_H^2} + 2, \quad c_{H\omega,\tau} = 2m_{H\omega,\tau}W_{\omega,\tau}, \quad (18)$$

$$b_{H\omega,\tau} = \frac{(\sqrt{H_{\omega,\tau-1}} + \sqrt{H_{\omega,\tau+1}})}{\sigma_H^2}, \quad (19)$$

$$a_{P\omega,\tau} = \frac{2}{\sigma_P^2} + 2, \quad c_{P\omega,\tau} = 2m_{P\omega,\tau}W_{\omega,\tau}, \quad (20)$$

$$b_{P\omega,\tau} = \frac{(\sqrt{P_{\omega-1,\tau}} + \sqrt{P_{\omega+1,\tau}})}{\sigma_P^2}. \quad (21)$$

3.2 Method 2

Since eq. (9) is a quadrature form of $H_{\omega,\tau}$ and $P_{\omega,\tau}$ with a linear constraint, the optimal $\boldsymbol{H}$ and $\boldsymbol{P}$, $\boldsymbol{m}$ can be obtained by solving a simultaneous equation but it includes a large number of variables equal to the number of time-frequency bins. To avoid it and derive a simple iterative solution, we derived the following auxiliary function:

$$\begin{aligned} & Q_2(\boldsymbol{H}, \boldsymbol{P}, \boldsymbol{U}, \boldsymbol{V}) \\ = & -\frac{1}{\sigma_H^2}\sum_{\omega,\tau}\left\{(\hat{H}_{\omega,\tau-1} - U_{\omega,\tau})^2 + (\hat{H}_{\omega,\tau} - U_{\omega,\tau})^2\right\} \\ & -\frac{1}{\sigma_P^2}\sum_{\omega,\tau}\left\{(\hat{P}_{\omega-1,\tau} - V_{\omega,\tau})^2 + (\hat{P}_{\omega,\tau} - V_{\omega,\tau})^2\right\} \end{aligned} \quad (22)$$

satisfies

$$J_2(\boldsymbol{H}, \boldsymbol{P}) \geq Q_2(\boldsymbol{H}, \boldsymbol{P}, \boldsymbol{U}, \boldsymbol{V}), \quad (23)$$

where $U_{\omega,\tau}$ and $V_{\omega,\tau}$ are auxiliary variables and $\boldsymbol{U}$ and $\boldsymbol{V}$ are sets of $U_{\omega,\tau}$ and $V_{\omega,\tau}$, respectively. The equality of eq. (10) is satisfied for $U_{\omega,\tau} = (\hat{H}_{\omega,\tau-1} + \hat{H}_{\omega,\tau})/2$ and $V_{\omega,\tau} = (\hat{P}_{\omega-1,\tau} + \hat{P}_{\omega,\tau})/2$. By taking the constraint of eq. (7) into consideration and organizing variables, we have the following update rules, which guarantees to monotonically increase the objective function J_2. The detailed derivation is presented in [12].

$$\begin{aligned} \Delta_{\omega,\tau} \leftarrow & \ \alpha\left(\frac{\hat{H}_{\omega,\tau-1} - 2\hat{H}_{\omega,\tau} + \hat{H}_{\omega,\tau+1}}{4}\right) \\ & -(1-\alpha)\left(\frac{\hat{P}_{\omega-1,\tau} - 2\hat{P}_{\omega,\tau} + \hat{P}_{\omega+1,\tau}}{4}\right) \end{aligned} \quad (24)$$

$$\hat{H}_{\omega,\tau} \leftarrow \min(\hat{W}_{\omega,\tau}, \max(\hat{H}_{\omega,\tau} + \Delta_{\omega,\tau}, 0)), \quad (25)$$

$$\hat{P}_{\omega,\tau} \leftarrow \hat{W}_{\omega,\tau} - \hat{H}_{\omega,\tau}, \quad (26)$$

where

$$\alpha = \frac{\sigma_P^2}{\sigma_H^2 + \sigma_P^2}. \quad (27)$$

In method 2, any γ is allowable. According to our experiments, setting γ to be about 0.3 gives a good performance.

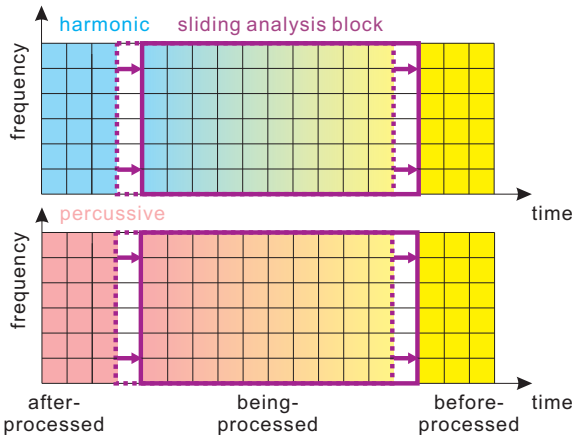

Figure 1. The process of the sliding block analysis

4 REAL-TIME PROCESSING BY SLIDING BLOCK ANALYSIS

Although the objective functions eq. (6) and eq. (9) should include all time-frequency bins, the iterative updates for the whole bins are much time-consuming. In order to obtain an approximate solution in real-time, we propose a sliding update algorithm. Based on the assumption that the separation of a certain time-frequency bin is weakly affected by far bins, we limit the processed frames to $n \leq \tau \leq n + B - 1$, where B is the size of the analysis block, and slide n iteratively. The real-time version of the Method 1 is summarized as follows.

1. Set the new frame as $H_{\omega,n+B-1} = P_{\omega,n+B-1} = W_{\omega,n+B-1}/2$.
2. Update variables by eq. (14), eq. (15), eq. (16), and eq. (17) for $n \leq \tau \leq n + B - 1$.
3. Convert the nth frame to a waveform by the inverse-STFT.
4. Increment n to slide the analysis block.

The real-time version of the Method 2 is in the same way. In step 3, the original phase is used for converting the STFT domain to the time domain. Note that the overlap of the frame shift should actually be considered for the conversion.

Each time-frequency bin is updated only once at step 2. Then, it is totally updated B times after passing through the analysis block shown in Fig. 1. Although the larger block size B shows better performance, the processing time from step 1 to step 4 must be less than the length of the frame shift for real-time processing.

5 IMPLEMENTATION AND EVALUATIONS

We implemented our algorithms in VC++ on Microsoft Windows environment. The GUI of the prototype system is shown in Fig. 2. After clicked a start button, the separation process begins. The processing steps are as folllows.

1. Loading a frame-shift-length fragment of the input audio signal from a WAV-formated file.
2. Calculating FFT for a new frame.
3. Updating stored frames as described in the previous section.
4. Calculating inverse FFT for the oldest frame.
5. Overlap-adding the waveform and playing it.
6. Go to Step 1.

The two bar graphs shown in Fig. 2 represent the power spectra of the separated harmonic and percussive component. The sliding bar named "P-H Balance" enables an user to change the volume balance between the harmonic and percussive components on play. The examples of the separated two spectrogram sequences are shown in Fig. 3. We can see that the input power spectrogram is sequentially separating in passing through the analysis block. In auditory evaluation, we observed:

- The pitched instrument tracks and the percussion tracks are well separated in both of method 1 and 2.
- Under the same analysis block size, the method 1 gives a little better performance than method 2.
- The method 1 requires about $1.5 \sim 2$ times computational time than the method 2 because of the calculation of square root. Thus, the method 2 allows the larger block size.
- The separation results depend on several parameters as σ_H, σ_P, the frame length, and the frame shift. But the dependency is not so large.

In order to quantitatively evaluate the performance of the harmonic/percussive separation and the relationship to the block size, we prepared each track data of two music pieces (RWC-MDB-J-2001 No.16 and RWC-MDB-P-2001 No.18 in [13]) by MIDI-to-WAV conversion and inputed the summation of all tracks to our algorithms. As a criterion of the performance, the energy ratio of the harmonic component $h(t)$ and the percussive component $p(t)$ included in each track was calculated as

$$r_h = \frac{E_h}{E_h + E_p}, \quad r_p = \frac{E_p}{E_h + E_p}, \tag{28}$$

where

$$E_h =< f_i(t), h(t) >^2, \quad E_p =< f_i(t), p(t) >^2, \tag{29}$$

and $<>$ represents the cross correlation operation and $f_i(t)$ represents a normalized signal of each track. The results are shown in Fig. 4. The pitched instrument tracks and the percussion tracks are represented by solid and dotted lines,

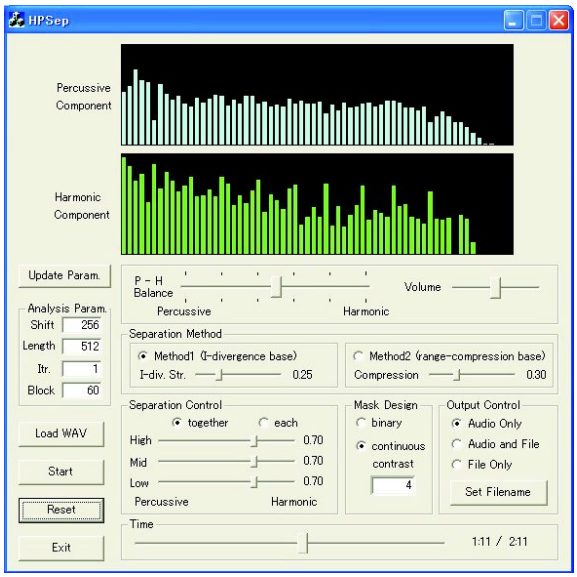

Figure 2. The GUI of the harmonic/percussive equalizer

Table 1. Experimental conditions

signal length	10s
sampling rate	16kHz
frame size	512
frame shift	256
range-compression factor (method 1)	$\gamma = 0.5$
range-compression factor (method 2)	$\gamma = 0.3$
gradient variance	$\sigma_P = \sigma_H = 0.3$

respectively. We can see that the separation was almost well performed. Only the bass drum track has a tendency to belong to the harmonic component, which can be considered due to the long duration. Fig. 4 also shows that a large block size is not required and the separation performance converges at the block size of 30 or 40 frames in this condition.

6 CONCLUSION

In this paper, we presented a real-time equalizer of harmonic and percussive components in music signals without any a priori knowledge of score and included instruments. In auditory evaluation and experiments, we confirmed the good performance. Based on our equalizer, applying existing audio modification technique as conventional equalizing, reverb, pitch-shift, etc., to harmonic/percussive components independently will yield more interesting effect. Applying it as pre-processing for multi-pitch analysis, chord detection, rhythm pattern extraction, is another interesting future work.

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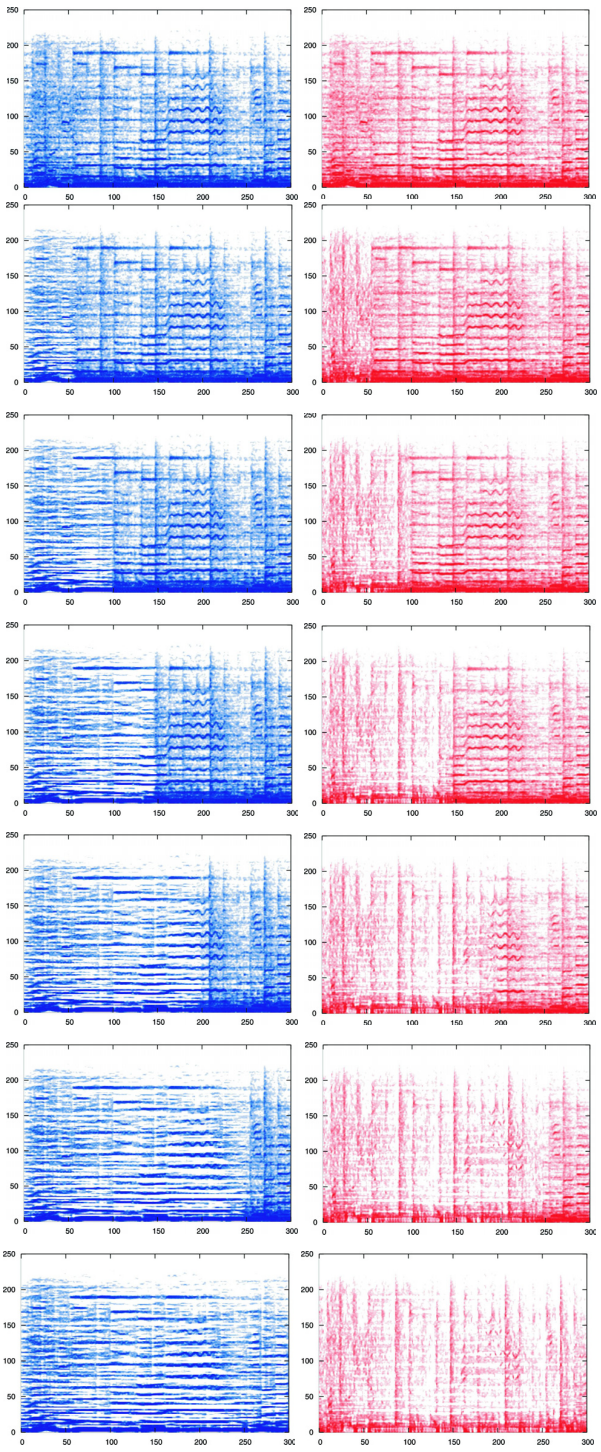

Figure 3. The spectrograms of separated harmonic component (left) and percussive component (right) by sliding block analysis. The first frame of the analysis block is 0, 10, 50, 100, 150, 200, and 250 from top to bottom, respectively.

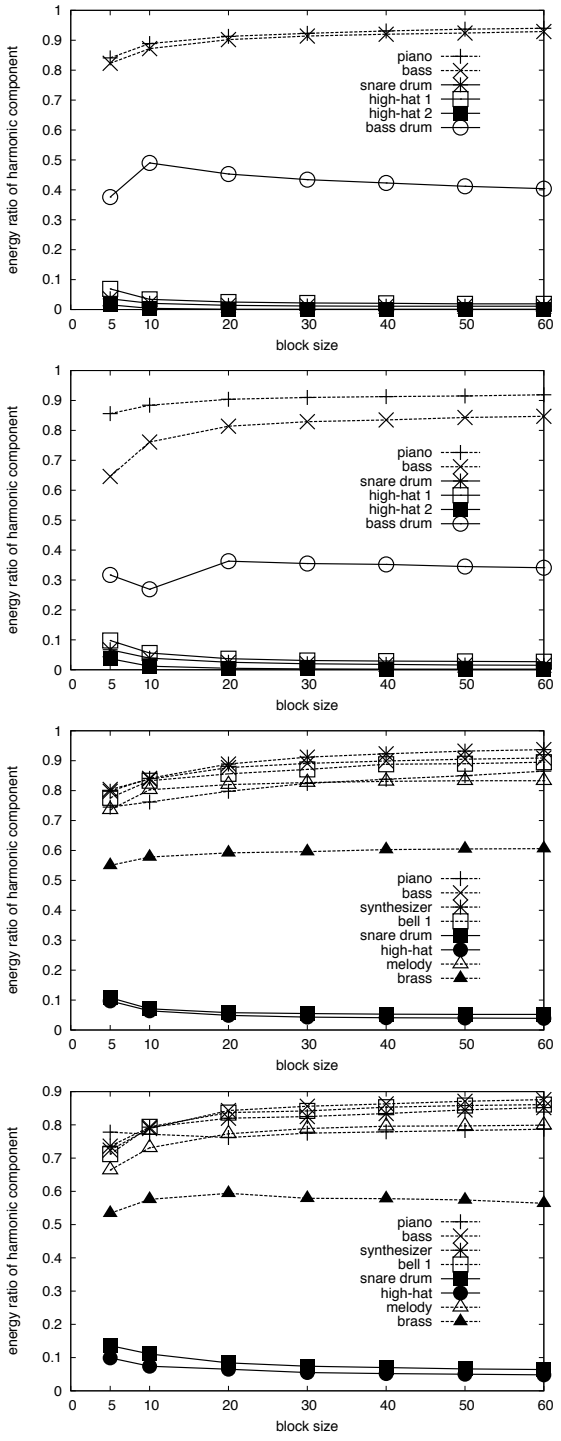

Figure 4. The energy ratio of the separated harmonic component in each track (r_h) for different block sizes. Their results from top to bottom are obtained by method 1 for RWC-MDB-J-2001 No.16, by method 2 for RWC-MDB-J-2001 No.16, by method 1 for RWC-MDB-P-2001 No.18, and by method 2 for RWC-MDB-P-2001 No.18, respectively.

SEGMENTATION-BASED LYRICS-AUDIO ALIGNMENT USING DYNAMIC PROGRAMMING

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ABSTRACT

In this paper, we present a system for automatic alignment of textual lyrics with musical audio. Given an input audio signal, structural segmentation is first performed and similar segments are assigned a label by computing the distance between the segment pairs. Using the results of segmentation and hand-labeled paragraphs in lyrics as a pair of input strings, we apply a dynamic programming (DP) algorithm to find the best alignment path between the two strings, achieving segment-to-paragraph synchronization. We demonstrate that the proposed algorithm performs well for various kinds of musical audio.

1 INTRODUCTION

As the market for portable media players increases rapidly, more and more manufacturers and multimedia content providers are searching for outstanding features that will set their offering apart from the rest. At the same time the rush towards online distribution of digital assets has stripped most content of rich metadata such as album artwork and lyrics. While artwork is gaining a lot of traction with the introduction of higher resolution color displays in devices, only few high end devices offer the ability to display lyrics in a readable form along with music. As devices tend to get smaller and smaller with advancing technology, there is not much hope for displays to become large enough for the user to comfortably follow song lyrics without some form of synchronized presentation. Some devices are already capable of displaying lyrics synchronized to the playback of music using manually inserted time stamps into the lyrics file.

However, this approach does not scale well for large collections, so an automated approach to align lyrics with music is the strongly preferable approach. It is notable that word-by-word level synchronization for most applications is not necessary. A paragraph or line-by-line synchronization, whatever is more appropriate for the device display resolution, will be sufficient. As this promises to be a fairly solvable problem, to which a scalable automated solution could be provided, it has recently attracted a number of researchers in the music information retrieval (MIR) community.

Wang *et al*., for example, have proposed a hierarchical approach for automatic alignment of acoustic musical signals with textual lyrics [12, 9]. They decompose the problem into two separate tasks — a higher-level section-based alignment followed by lower-level per-line alignment. To this end, they first process audio to obtain high-level structural information such as Measure, Chorus and Singing Voice Section. In parallel, textural lyrics are analyzed and each section is labeled with one of the pre-defined section types. In this text processing stage, they also compute approximate durations of each section and line. However, their algorithm is limited by strong assumptions about the song structure as well as the fixed rhythmic structure.

A different approach has been taken by Chen *et al*. who have presented an automatic lyrics-audio synchronization system using the low-level acoustic features only [3]. Their algorithm has two main components: 1) vocal/non-vocal detector and 2) alignment of the audio signal with its lyrics at multiple levels using acoustic models. Given a musical audio signal, a vocal/non-vocal classifier detects candidates for the singing voice sections. In parallel, they construct the grammar net from the lyrics and force an alignment utilizing the previously obtained acoustic model units with a maximum likelihood linear regression technique. They have tested their algorithm on a small set of Chinese song segments and achieve a boundary accuracy of 81.5% at the phrase level.

Fujihara *et al*. tackle the lyrics-audio alignment challenge by solving three sub-problems in series [8]: *i.e*., 1) separation of singing voice, 2) singing voice detection, and 3) alignment of segregated vocal signals with lyrics using a *Viterbi*-based matching technique. At the final alignment stage, they first build a language model from the lyrics using only vowel phonemes and short pauses between word, sentence or phrase boundaries. They also employ an adaptation of a phone model to the specific singer of the input audio signal to improve performance. Using 10 Japanese popular songs as test bed, they have achieved over 90% accuracy for eight songs.

The ultimate goal of the lyrics-audio alignment systems described so far is to automate karaoke-style synchronization at a line or word level. Although a word-level or even

syllable-level synchronization may appear to be ideal, it is extremely challenging to achieve and usually involves solving other difficult problems such as singing voice separation or constructing proper speech models.

In this paper, we deviate from this goal and propose a solution to a simpler, more basic problem: we aim to align song lyrics to the corresponding audio signal at a segment-to-paragraph level. Paragraph-level alignment may not be sufficient for karaoke applications, but as stated initially we believe this will help users follow the lyrics as they listen to music by providing them with referential points over time. Furthermore, as reported by Wang *et al*. [12, 9], section- or segment-level alignment provides an initial solution that makes lower-level alignment such as line- or word-level alignment easier and more robust.

This publication is organized as follows. We describe our proposed method for automatic lyrics-audio alignment in detail in the next Section. Thereafter, in Section 3, we present experimental results with several real examples. Finally, in Section 4, we draw conclusions from the previously presented results, and give an outlook on future work in this domain.

2 METHOD

As mentioned in Section 1, our system for lyrics-audio alignment strives to achieve paragraph-to-segment level synchronization. The motivations for this are as follows. First, we find that the segment structure in musical audio corresponds approximately to the paragraphs in lyrics. For example, a `verse` and/or `chorus` section is found in both audio and lyrics for most popular music, even when investigating one without knowledge of the other. Therefore, if we can divide an entire song into the appropriate segments, we can search for the corresponding paragraphs in lyrics.

Second, paragraph-to-segment level alignment is far easier and more robust than word-level or syllable-level alignment because the latter usually depends on other complex algorithms such as singing voice separation and/or speech recognition, which are very challenging problems by themselves. On the other hand, structural music segmentation has achieved fairly good performance with relatively simple and straightforward techniques, as previously discussed by many other researchers [7, 5, 1, 10].

Therefore, the first stage in our system consists of a segmentation of musical audio, and is described in the next paragraph.

2.1 Structural Segmentation

Structural music segmentation is one of the major research topics in the current field of MIR. We do not intend to solve this problem in this paper, however, and thus we briefly describe the algorithm we use for segmentation as an example.

Most contemporary and classical music has a certain structure or pattern with regard to its temporal evolution that contains systematic change and repetition; for example, a pattern like {`intro-verse-chorus-verse-chorus-bridge-solo-chorus-outro`} is observed to be very common in rock or pop music. Structural music segmentation is referred to as finding the boundaries between these sections. Many algorithms have been proposed so far to find the section boundaries in musical audio. We use an approach that is based on the self-similarity matrix of low-level features derived from the audio signal [6, 7].

In this algorithm, the entire song or musical piece is processed using common spectral analysis methods, and a sequence of feature frames is obtained. A 2-dimensional self-similarity matrix S is then computed, where an element $S(i, j)$ represents a similarity measure between the feature-frame pair (f_i, f_j). A cosine distance is used as similarity metric. Then a kernel correlation is performed on the similarity matrix S to generate a 1-dimensional novelty score, whereby peaks indicate significant changes in musical audio. The novelty score is smoothed in order to suppress spurious peaks. Subsequently, the final segment boundaries are obtained by choosing the peaks whose values are above a certain heuristically determined threshold. Foote and Cooper also use the self-similarity matrix and the segmentation results to cluster the segments and summarize music [4, 5], where similar segments are grouped as a *cluster* like `chorus` or `verse`.

The final output of the segmentation algorithm results therefore in a sequence of segment labels and their corresponding time boundaries. For example, using the algorithm for segmentation outlined above, the final output of *"Super Trouper"* by *A*Teens* is {G-B-D-C-A-B-D-C-A-C-A}, where each label or letter represents a segment, and the segments with the same label represent a cluster. At the clustering stage, we locate the cluster within which the segments are closest to each other, and label it as `chorus`, which is denoted as 'A' in the above example. This simplified heuristic is based on the observation that in by large most popular music the `chorus` section remains mostly unaltered throughout the entire song, as it represents the main theme or hook of the song which is emphasized by repetition as much as other musical techniques.

In the following sections, we explain how we align the paragraphs found in lyrics with the segments that our segmentation algorithm was able to retrieve.

2.2 Paragraph-to-Segment Alignment

As is mentioned above, the segmentation algorithm outputs a sequence of labels with additional clustering information. This and the location of `chorus` sections in particular, yield partial but useful information about how to coarsely align lyrics with audio, provided that we have lyrics that have

been hand-labeled at a paragraph or section level. That is, we can synchronize lyrics with audio using one or multiple `chorus` sections as *anchoring* points. For instance, the structure of lyrics in the above example by *A*Teens* is {`intro-verse-verse-chorus-chorus-verse-verse-chorus-chorus-bridge-chorus-chorus-chorus`}. Although there are seven `chorus` paragraphs in lyrics, we can group the consecutive `chorus` paragraphs to obtain three of them, as many as have been derived from the audio segmentation.

However, there are usually many other segments in the audio signal that we don't have any knowledge of as to their correspondence with the appropriate paragraphs within the lyrics. For example, using the above mentioned example again, a segment 'G' could be an instrumental `intro` or an `intro` with some lyrics content. For repetitive segments such as 'B', 'C', or 'D', we can assume them to be a `verse` section because the `verse` theme or themes usually repeat in most popular music. However, 'B', 'C', or 'D' may be a repetitive instrumental section, as well. Therefore, even if we achieve some basic alignment using `chorus` sections as points of reference we still will have many other segments unaligned, about which we have too little information as to their contents. We propose in the next section a simple solution to this alignment problem using a dynamic programming (DP) algorithm.

2.3 Dynamic Programming

In a dynamic programming (DP) algorithm an optimization problem is first divided into smaller sub-problems, and the optimal scores for the solution of every sub-problem are stored and used to solve an entire problem, without resolving the sub-problems over and over again [2]. DP algorithms are widely used in applications such as DNA sequence matching to efficiently and optimally align two strings of different length, and compute the distance between them.

We can use a DP algorithm for our task because, after structural segmentation of audio, we have obtained two sequences of input strings — one derived directly from the audio signal and the other extracted from the structural clues within the lyrics text — whereby both sequences are of different length. In order to find the minimum-cost alignment path using DP, however, we first need to compute an *error matrix* E from the two input sequences, where an element $E(i, j)$ represents the distance between the ith element of a sequence S_1 and the jth element of a sequence S_2. Hence, it is critical to appropriately define the pairwise distance between an audio segment and a lyrics paragraph so as to achieve meaningful alignment results.

Therefore, we first simplify the input string describing the audio structure by reducing the number of label types. That is, we re-label the audio segments so that there are only three labels — 'C' for `chorus`, 'V' for `verse` and 'O' for all others. However, as mentioned above, we don't know for certain which segment within the audio structure corresponds to which in the lyrics except for the `chorus`. Thus we label the segments that repeat twice or more as `verse` or 'V', and those that appear only once as 'others' or 'O'. As to labeling lyrics, we define four different labels — 'C' for `chorus`, 'V' for `verse`, 'I' for `intro` and 'B' for `bridge` — which cover all of the lyrics in our experiment data set.

The next step consists of defining a distance measure between every paragraph-segment pair. First, we determine the minimum distance between a `chorus` pair to use it as an anchor in the alignment process. This is essential because a `chorus` section is the most representative part in most popular music and therefore we would like to avoid misalignment there. Secondly, we assign the second smallest distance to a `verse` pair, but with less confidence than `chorus` pairs, because we are less certain about it being a true `verse` segment from our segmentation results than with the `chorus` segments — these segments can more often be confused with introductory solo parts, bridges or other sections aligned with the verses in one fashion or another. We define the distance between every further pair in a similar manner. Table 1 shows a distance matrix that has been generated as part of our experiments for every possible lyrics-audio pair.

Table 1. Distance matrix between paragraph-segment pairs

audio segments \ lyrics paragraph	C	V	I	B
C	0.0	0.5	0.3	0.5
V	0.5	0.2	0.5	0.3
O	0.7	0.5	0.3	0.3

After the error matrix E is computed using the pairwise distance matrix in Table 1, an accumulated error matrix is calculated by storing the minimum cost of the three possible directions from the previous step (*i.e.*, $\rightarrow, \downarrow, \searrow$). By backtracking, the DP algorithm then retrieves the optimal alignment path with minimum accumulated error.

3 EXPERIMENTS

Using the paragraph-to-segment level alignment algorithm described so far, we have performed experiments on a number of musical items representing a variety of genres and styles of popular music. Figure 1 shows the alignment results of *"Super Trouper"* by *A*Teens*. Shown is the error matrix between the alignment of lyrics and audio at a paragraph-to-segment level, which is computed using the distance measure in Table 1. The minimum-cost alignment path found by the DP algorithm is also displayed in thick, solid lines.

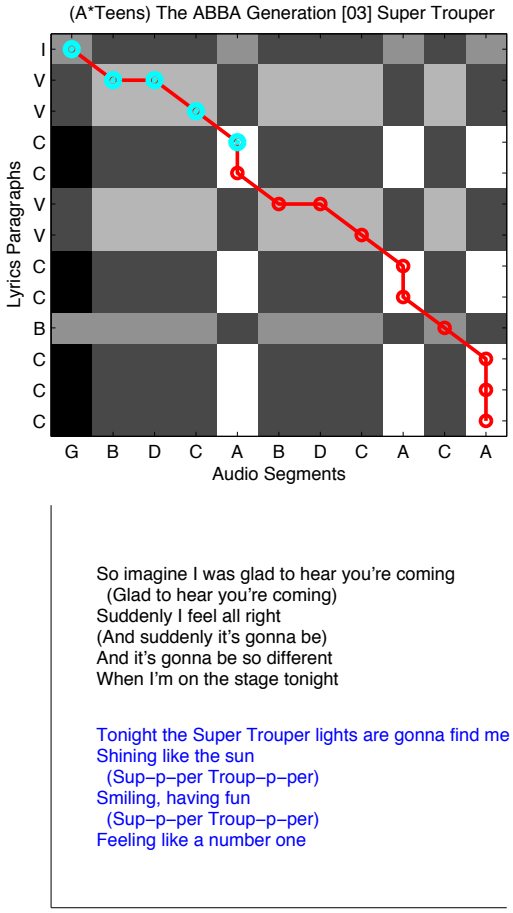

Figure 1. Alignment results of *"Super Trouper"* by *A*Teens*. At the top is shown the error matrix along with the optimal alignment path. At the bottom is displayed lyrics paragraphs corresponding to previous and current audio segments being played.

As shown in Figure 1, `chorus` sections ('C' for lyrics and 'A' for audio) are always aligned since the distance between the two is 0 by definition. Thereby anchoring points are provided which will help to align the entire song subsequently. This also allows for automatic grouping of consecutive `chorus` paragraphs in lyrics and mapping them to corresponding `chorus` segments in audio.

However, some issues can be observed in these alignment results. First of all, we can never guarantee a one-to-one mapping between lyrics paragraphs and audio segments because the number of paragraphs in lyrics seldom match the number of segments in audio. Therefore, there are cases where two or more paragraphs are assigned to one audio segment or vice versa. We approximately resolve the first case by equally dividing the audio segment by the number of lyrics paragraphs so that both lyrics and audio have the same number of paragraphs/segments. For example, we observe in Figure 1 that two `chorus` ('C') paragraphs are assigned to the first `chorus` ('A') segment. Here, we display the first 'C' in the beginning of 'A' and the second 'C' at the half the duration of 'A'. This approach results in a more realistic though still somewhat inaccurate alignment.

The more serious issue occurs when there are more than two audio segments assigned to one lyrics paragraph, such as 'B' and 'D' assigned to the first 'V', as shown in Figure 1. The same solution we use for the first case — *i.e.*, subdividing 'V' into two equal paragraphs and mapping them to 'B' and 'D' — won't work because 'B' is an *instrumental* segment in this example. Therefore, correct mapping would result in {V-V}–{D-C}, and in this case by skipping the instrumental 'B' segment. However, we are unable to conclude from the segmentation results which segment contains the singing voice and which is purely instrumental, or if indeed both contain part of the sung lyrics.

We therefore increase the alignment accuracy by first classifying the audio segments into vocal and non-vocal parts, and then applying the alignment algorithm only on the audio segments that have a higher likelihood of containing vocals. We perform segment-level vocal/non-vocal discrimination based on the classifier proposed by [11]. Table 2 shows the vocal/non-vocal classification results on *"Super Trouper"* by *A*Teens*.

Table 2. Vocal/non-vocal classification results

Segment	G	B	D	C	A	B	D	C	A	C	A
V/NV classification	V	NV	V	V	V	NV	V	V	V	V	V

Using the above referenced vocal/non-vocal classification results, we can simplify the problem significantly by ignoring the non-vocal audio segments in the alignment process. That is, we replace the original segmentation with {G-D-C-A-D-C-A-C-A} and apply the DP algorithm to obtain the final alignment results as shown in Figure 2.

As shown in Figure 2, two instrumental audio segments (denoted by 'B') are not assigned any lyrics, and this provides a much increased accuracy for following the lyrics along the audio during playback. We also notice that dividing 'A' segments into sub-segments of equal length similarly leads to more precise alignment (indicated in dash-dot lines).

Our alignment algorithm has been tested on 15 popular music items of different musical styles, including pop, rock, R&B, punk and so on. The number of paragraphs in the lyrics for each song of this set varies from 5 to 27. Table 3 displays more specific information about the songs in the test bed. Because the algorithm proposed in this paper has been designed to obtain paragraph-to-segment alignment, we have evaluated the alignment accuracy by compar-

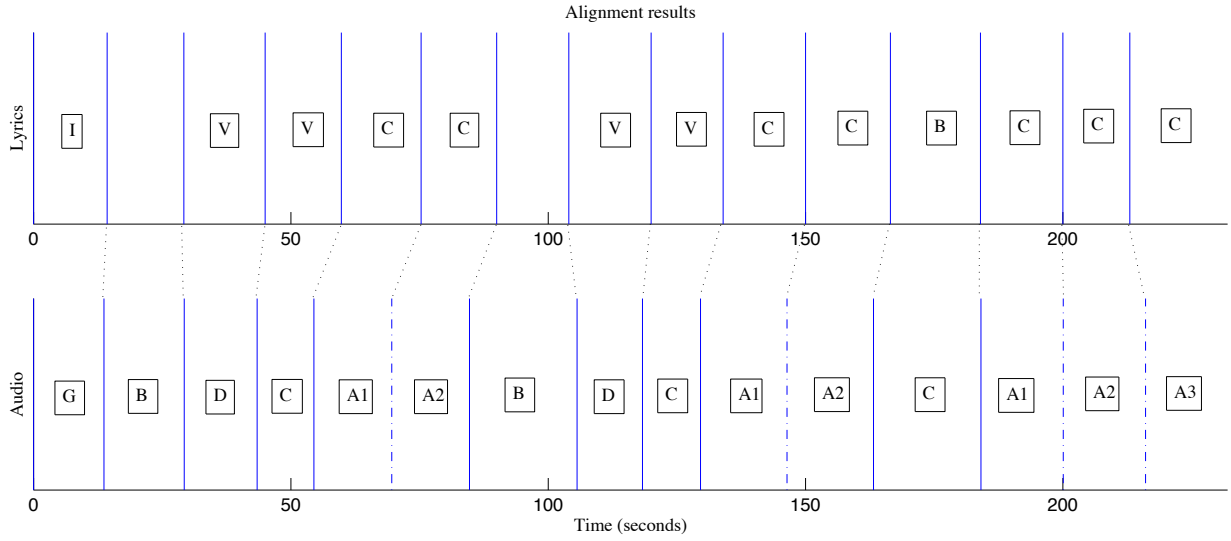

Figure 2. Final alignment results of *"Super Trouper"* by *A*Teens*. `chorus` sections in audio ('A') are sub-divided by dash-dot lines as explained in the text.

ing the manually-marked starting point of each paragraph in the corresponding audio file to that of the automatically selected audio segment.

Using 174 paragraphs in total in the test bed detailed in Table 3, overall we have obtained an average error of 3.50 seconds and a standard deviation of 6.76 seconds. These results reflect the robustness of our alignment algorithm, even though the test songs' styles and lyrical/musical structures vary to a great degree, and the segmentation/clustering results are noticeably different from song to song. The song yielding the best result is *"Hand In My Pocket"* by *Alanis Morissette*, which achieved an average error of 0.91 seconds with a standard deviation of 1.19 seconds. *"I Ran (So Far Away)"* by *A Flock Of Seagulls* produced the least satisfying outcome of 11.25 and 10.48 seconds in average error and standard deviation, respectively.

4 CONCLUSIONS

In this publication, we have presented an algorithm to align song lyrics with a corresponding musical audio recording at a paragraph-to-segment level, as we believe that multiple applications in the entertainment market can benefit from such an alignment. To this avail, we first perform a structural segmentation and clustering of the audio signal, which outputs a sequence of labels where similar segments have been assigned the same label. Lyrics has been hand-labeled at a paragraph level, and we use this pair of label strings as an input to a dynamic programming algorithm. Based on the paragraph-segment distance measure that has been constructed to precisely fit the requirements, the DP algorithm finds the minimum-cost alignment path between lyrics and audio. In addition, we improve the alignment performance by detecting the vocal segments in audio and discarding the non-vocal segments in the alignment process.

Experiments on various kinds of popular music show that the proposed algorithm successfully synchronizes lyrics to audio with all the unintended variations caused by the inaccuracy of the segmentation and clustering of the audio signal.

In the near future, we consider a hierarchical system for lyrics-audio alignment as suggested by Wang *et al*. [12, 9]. In other words, we plan to develop an algorithm for lower-level per-line alignment based on the paragraph-level alignment we achieved in this paper. We believe this hierarchical approach, or an equivalent "divide-and-conquer" method, will be more robust and accurate than performing line-level alignment for an entire song without additional intermediate steps. In addition, a vocal/non-vocal classifier which operates on a finer time grid will allow to more precisely locate the singing voice sections, which we believe will further correct the mis-alignment caused by false segmentation boundaries.

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Table 3. Songs in the test bed

Song ID	Title	Artist	Genre	# of paragraphs	# of segments
1	Hold On Loosely	.38 Special	Rock	9	13
2	I Ran (So Far Away)	A Flock Of Seagulls	Pop	9	12
3	Try Again	Aaliyah	R&B	27	13
4	Don't Turn Around	Ace Of Base	Pop	17	12
5	Hand In My Pocket	Alanis Morissette	Rock	11	10
6	Save Me	American Hi-Fi	Alternative Rock	12	9
7	There's A Star	Ash	Punk	7	10
8	Super Trouper	A*Teens	Pop	13	11
9	Like Dylan In The Movies	Belle & Sebastian	Indie Rock	7	12
10	Summer House	Better Than Ezra	Alternative Rock	5	9
11	Crazy In Love	Beyoncé Feat. Jay-Z	R&B	24	9
12	White Wedding	Billy Idol	Pop	13	11
13	That's Cool	Blue County	Country & Folk	7	10
14	Parklife	Blur	Rock	10	7
15	Here Comes My Baby	Cat Stevens	Pop	7	6

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MULTIPLE-FEATURE FUSION BASED ONSET DETECTION FOR SOLO SINGING VOICE

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ABSTRACT

Onset detection is a challenging problem in automatic singing transcription. In this paper, we address singing onset detection with three main contributions. First, we outline the nature of a singing voice and present a new singing onset detection approach based on supervised machine learning. In this approach, two Gaussian Mixture Models (GMMs) are used to classify audio features of onset frames and non-onset frames. Second, existing audio features are thoroughly evaluated for this approach to singing onset detection. Third, feature-level and decision-level fusion are employed to fuse different features for a higher level of performance. Evaluated on a recorded singing database, the proposed approach outperforms state-of-the-art onset detection algorithms significantly.

1. INTRODUCTION

Accurate monophonic singing voice transcription depends heavily on singing onset detection. Current onset detection algorithms applied on monophonic singing voice produced poor results [5][8]. In MIREX2007, the best result for onset detection of solo singing voice only managed an F-measure of 51.0% [2]. The singing voice is a pitched non-percussive (PNP) instrument [17], which is still a challenging category of instruments for onset detection. The case is further complicated by the nature of the singing voice, which is inherently inconsistent and prone to pitching and timing dynamics.

Current onset detection methods for PNP instruments, such as methods by spectral difference [7][10], phase deviation [6], pitch shift [9], and sub-band energy change [11][12] are targeted at detecting spectral changes based on certain rules. It has been observed on a spectrogram that in general, a musical note onset occurs at locations where there is a visible change in the spectrum, and within the duration of the note, the spectrum is relatively stable. A violin signal is shown as an example in Fig.1a.

However, for the case of singing voice, this observation may not always hold (Fig.1b). Unlike most other instruments, where there is usually a high level of timbre consistency in the duration of a note, the singing voice is capable of producing much more variations of formant structures (for articulation); sometimes the formant structure may even change within the duration of a single note. Pitch irregularities, pitch modulations, and inadvertent noise also upset stability of the spectrum.

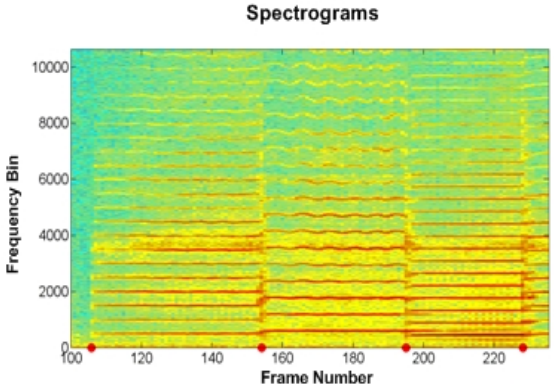

Figure 1: Comparison between spectrograms of violin and singing signal. Onsets are marked by red circles

Onset detection in the singing voice is much more complicated than in most other instruments. An onset detector that relies explicitly on a rule-based approach is bound to fail to capture the nuances of the singing onset. To address the difficulties in singing onset detection, we present a new onset detector based on supervised machine

learning, in order to capture the intricacies of singing note onsets, illustrated in Figure 2.

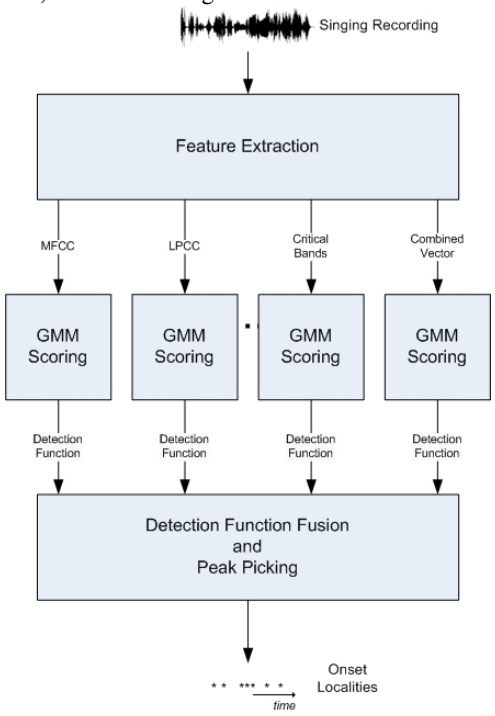

Figure 2: System diagram of the proposed system

Two GMMs are used to profile the distribution of audio features of onset frames and non-onset frames, respectively. We thoroughly evaluate existing audio features for their effectiveness in distinguishing onset and non-onset frames, including Mel-frequency Cepstral Coefficients (MFCCs), Linear Predictive Cepstrum Coefficients (LPCCs), Equal loudness phon values along critical bands. Feature concatenation fusion in feature-level and linear weighted sum fusion in decision-level are then employed to achieve a high level of onset detection accuracy. Evaluated using our singing database, our proposed approach to singing onset detection outperforms state-of-the-art methods, including methods based on phase change [6], pitch change [9], equal loudness [12], and inverse-correlation [7]. Re-implementations of these methods are used in the evaluation.

2. DEFINITION OF A SINGING ONSET

In a non-percussive continuous-pitch instrument, such as the melodic singing voice, there is no clear definition of an onset. A definition is especially difficult to establish when taking into account glissandos[1] and portamentos[2], which are vocal expressions involving pitch glides. Traditionally, an onset is the beginning of a new note event [5][8], characterized by a change in pitch, amplitude envelope, spectral shape, phase, or a combination of the mentioned factors [14].

However, in a singing voice, which is plagued by accidental glissandos, slurring (melisma[3]) and imperfect vocalization, the definition of an onset is a less clear-cut one. It is very often that a singer performs a glissando (more accurately, portamento in the singing voice) at the beginning or ending of a note, even when a music score does not stipulate so. When a musician is tasked to transcribe the singing performance, these portamentos are not transcribed into notes. However, slurred performances, which also involve pitch glides, are usually transcribed as actual notes, albeit with a legato denotation. There is no real physical difference between a portamento and a slur in singing performances. In both cases, the singing voice undergoes a continuous change in pitch. However, perceptually many glissandos are not considered as note onsets. This is because when listening to a singer's performance, a human subconsciously takes into account contextual cues like the rhythm, language and style of music.

The nature of singing onset is strongly related to the manner of articulation in speech. A singing voice has strong harmonic components producing numerous timbral differences, which are interpreted as different phonemes by the human ear. Singing onsets usually, though not always, occur during vowel-onsets.

Since there is no known existing definition for a singing onset, it is crucial to define one before even attempting to evaluate the effectiveness of a singing onset detector. In [5], an onset is defined as the start of the transient, but in a pitched singing voice (and many PNP instruments), there are numerous notes with no observable transient, particularly during slurs. A singing performance may also contain long stretches of unvoiced consonant segments, typically at the beginning or end of the note. These unvoiced segments are too varied and inconsistent to base the definition of a singing note onset on, and should simply be regarded as noise in the annotation process, not as part of a singing note. An onset should be marked at the end of a consonant, not before or during the consonant.

For practical purposes, it is intuitive and reasonable to define a singing onset as follows: The beginning of a new human-perceived note, taking into account contextual cues. This excludes erroneous portamentos during note beginnings, transitions and trail-offs, but includes pitch changes in slurring. The actual notes included in a slur

[1] Glissandos are glides from one pitch to another, by filling the slide with discrete intermediate pitches. In continuous-pitch instruments, glissando is often used interchangeably with the term portamento.

[2] Portamentos are continuous glides from one pitch to another, usually produced as an effect by continuous-pitch instruments.

[3] Melisma is the act of changing pitch on a single syllable of sung text.

may be subject to human interpretation, since it is possible that a singer reaches the correct pitch only very briefly during a slur. Slurred notes that occur too briefly for an average human auditory system to reliably detect should not be included. The precise location of an onset in a slur is also subjective, and allowances ought to be made for human bias.

3. FEATURE EXTRACTION

The effectiveness of any onset detection function ultimately depends on the audio features used in the system. It is necessary to employ features that provide useful information on whether or not a frame contains an onset.

Features capable of capturing timbral differences should provide a reliable measure for onset detection. Pitch and energy features could provide important information regarding onset detection as well, but are much less reliable. We have chosen to focus on features that provide information on the spectral shape, since a timbral change is the main characteristic of an onset.

3.1. Mel Frequency Cepstral Coefficients

MFCCs represent audio by a type of cepstral representation. In a mel-frequency cepstrum, the frequency bands are positioned on a mel-scale, which aims to approximate the human auditory system's response.

It is a common feature used in speech recognition, and can be derived by first taking the Fourier transform of a (windowed) signal, mapping the log-amplitudes of the resulting spectrum into the mel-scale, then performing DCT on the mel log-amplitudes. The amplitudes of the resultant cepstrum are the MFCCs. Since DCT holds most of the signal information in the lower bands, the higher coefficients can be truncated without excessive information loss. In our system, we extracted MFCC features with 81 mel-scale filter banks and 23 DCT coefficients.

The coefficients are concatenated with their first and second-order derivatives to form a feature vector of 69 dimensions. This improves the performance of the system, due to the correlation of the MFCCs and the derivatives at feature-level.

3.2. Linear Predictive Cepstrum Coefficients

LPCCs are LPC coefficients transformed into cepstra, and are widely used in speech recognition systems. An efficient method of obtaining the coefficients using Levinsin-Durbin recursion is covered in detail in [1]. Once we obtain the LPC coefficients, they can be transformed into cepstra by [4]:

$$c_m = a_m + \sum_{k=1}^{m-1} \frac{k}{m} c_k a_{m-k} \quad (1)$$

where $\{c_1 \ldots c_N\}$ is the set of cepstral coefficients of order N. In our system, we used 22 cepstral coefficients, appended with 44 coefficients of the first and second order derivatives, similar to the feature-level fusion performed in the extraction of MFCCs.

3.3. Critical bands

Grouping frequency bins into critical bands is a psychoacoustically motivated principle. In [11], 36 bands of triangular-response bandwidth were used along the critical bands, and power in each band was used to derive a detection function. Detection functions based on equal-loudness changes in Equivalent Rectangular Bandwidth (ERB) bands have also been used in the past [8]:

$$\mathrm{E} = \lfloor 21.4 \log_{10} (4.37\,\mathrm{F} + 1) \rfloor \quad (2)$$

where E is the ERB band number, F is the bin frequency in kHz. By grouping frequency bins into ERB bands, we greatly reduce the dimensionality of the spectra. In [8], the power in individual ERB bands were mapped into phon values using equal-loudness contours [3]. In our system, we replicate the method introduced in [8], but only 36 ERB bands are used (includes frequencies of up to 11kHz) for audio files sampled at 22.05kHz.

3.4. Other Features Combined

Pitch-stability, zero-crossing rate and signal periodicity are all simple features which may contain information about note onsets. These features are commonly used, and can be concatenated into a combined feature vector. This feature-level data fusion is sensible for features which are characteristic of onset frames, especially if the features are of low dimensionality. Features which are of no value to the onset detection problem should not be included, as they corrupt the feature space and hence degrade the system performance. In our system, we used pitch stability, zero-crossing rate and signal periodicity and their derivatives as the combined vector.

4. GMM TRAINING AND SCORING

GMM-based modeling of speech signals has been the principal approach for speech and speaker recognition systems in recent years. It has been successfully implemented in [16] for speaker identification with very good results. Like speaker identification, onset detection can be modeled as a classification problem. At each time step, we need to classify a frame of audio into the onset class or a non-onset class.

Onset detection based on probabilistic models has increasingly been the preferred approach in recent audio research. Of particular interest are systems that employ machine learning algorithms that produced promising results, like Lacoste and Eck's FNN system [13]. Inspired by the success in speech processing systems, and in view of the similarity between the two problems, we employed a supervised machine learning algorithm using GMM classifiers.

4.1. GMM-based supervised machine learning

For each feature type, we model the probability of onset and probability of non-onset as random variables drawn from a Gaussian probability distribution of the feature vectors. Using a GMM to model the probability of onset random variable, we have:

$$P(x_n \mid \lambda_{onset}) = \sum_{i=1}^{M} w_i p_i(x_n) \quad (3)$$

where $P(x_n \mid \lambda_{onset})$ is the probability that feature vector x_n belongs to the onset class; w_i gives the weight of each of the M mixtures, and for each mixture:

$$p_m(x_n) = \sqrt{\frac{1}{(2\pi)^D \mid \Sigma_i \mid}} e^{-\frac{1}{2}(x_n-\mu_i)'(\Sigma_i)^{-1}(x_n-\mu_i)} \quad (4)$$

The parameters μ_i and Σ_i are the mean vector of dimensionality D, and the DxD covariance matrix, respectively. The parameters of the GMM can then be denoted $\lambda_{onset} = \{w_i, \mu_i, \Sigma_i\}$ where $i=1...M$. In our system, diagonal covariance matrices are used to parameterize the GMM, and 64 mixtures are chosen based on experiments. To train the onset GMM, the onset class training features are selected as the feature vectors of frames containing a hand-marked onset plus 3 frames at either side of each onset frame, which includes a total of 560 onsets (3920 frames).

We train this GMM by using the Expectation-Maximization (EM) algorithm [15] to profile the distribution of the onset-class feature vectors. We then proceed to construct another GMM, $\lambda_{non-onset}$, from the remaining feature vectors (21060 frames) for the non-onset class using (3) and (4).

4.2. Derivation of a Detection Function

With our trained GMMs (onset and non-onset classes), the probabilities $P(x_n \mid \lambda_{onset})$ and $P(x_n \mid \lambda_{non-onset})$ of each new feature vector can then be obtained. That is, for each new feature vector in time, we measure the likelihood that they belong to the onset and non-onset class. A viable detection function for our system will then be:

$$df(t) = P(x_t \mid \lambda_{onset}) - P(x_t \mid \lambda_{non-onset}) + 1 \quad (5)$$

where x_t is a feature vector of the audio signal obtained at time t. The detection function is then normalized to a range of [0…1].

5. DETECTION FUNCTION FUSION AND PEAK-PICKING

As aforementioned, for each feature type, we train 2 GMMs (for onset-class and non-onset-class features). Since we could have several feature types, we train a pair of GMMs for each feature type and derive a detection function for them using (5). Thereafter, we linearly weigh each of the detection function, and compute a sum of the individual weighted detection functions to produce a combined detection function:

$$df(t) = \sum_{i=1}^{F} w_i df_i(t) \quad (6)$$

F is the number of feature types, and w_i gives the weight of each of the detection functions.

It is also possible to use only a pair of GMMs, by simply concatenating the dimensions of each of the individual feature vector into a single feature vector at the feature-level, but solely relying on such an approach will increase the dimensionality of the feature space considerably, and it will require exponentially more training data in order to fully train the GMMs, according to the curse of dimensionality[1]. It is therefore prudent to keep each of our feature space limited to a feature type and its derivatives, possibly except for simple features of very low dimensions.

Once we obtain the detection function described by (6), we apply a de facto standard median-filter peak-picking algorithm, used in both [5] and [8] to evaluate the detection functions. The output of the peak-picking process is a series of onset locations denoting the time points at which an onset has occurred.

6. EXPERIMENTAL RESULTS

6.1. Database Description

Our database consists of 18 singing recordings of pop songs, from 4 singers (2 male, 2 female) of varied singing styles. The pieces were annotated by hand for onsets, and contained a total of 1127 onsets. Approximately half

[1] Curse of dimensionality is a term coined by Richard Bellman (1920-1984) describing the exponential relationship between volume and dimensionality.

(560) of the onsets were used for training GMMs, and the other half (567) were used to evaluate the system.

Each recording was annotated and cross checked by two amateur musicians. Onsets were first identified by ear, and then marked at positions where the wave form was first observed to follow a periodic structure. Consonant noise and unintentional portamentos / glissandos do not constitute new onsets, as explained in Section 2. For legato, an onset was marked at the point where a pitch change was perceived to begin.

6.2. Evaluation of Individual Detection Functions

For evaluation of the individual onset detection functions, we extract all the features mentioned in Section 3 and score each feature type based on a pair of trained GMMs (onset and non-onset GMMs). From the detection function produced by (5) for each feature type, we apply the median-filter peak-picking process across different parameters. We evaluate the usefulness of the feature by the metrics of precision, recall and F-measure, based on an onset tolerance of 50ms. Based on an onset tolerance of 50ms. These evaluation conditions are identical to those in the annual MIREX Audio Onset Detection contest.

The onset detection results of individual features are shown in Figure 3. The best F-measure among the set of peak-picking parameters is selected for each feature. From our experiments, MFCC is revealed to be the best audio feature, producing the best results of 80.3% precision, 77.8% recall and 79.0% F-measure. LPCC is next best, with 72.9% precision, 77.7% recall, and F-measure 75.3%. ERB-bands produce 78.2% precision, 69.5% recall, and 73.6% F-measure. The combined vector produces 60.1% precision, 86.1% recall, and 70.8% F-measure.

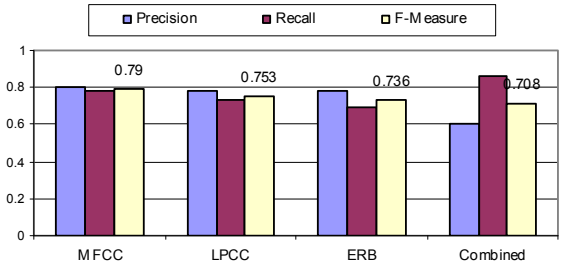

Figure 3: Evaluation results of individual detection function

6.3. Evaluation of Combined Detection Function

Based on the performance of all evaluated detection functions, we search the best linear weights based on extensive experiments and compute an overall detection function described by (6). The weight distribution is shown in Table 1.

Detection Function produced by:	Weight
MFCC	0.76
LPCC	0.12
ERB-bands	0.10
Combined vector	0.02

Table 1: Weight distribution for detection functions

Using the sum of linearly-weighted detection functions, we achieve the best performance of 86.5% precision, 83.9% recall, and 85.2% F-measure, which is superior to current state-of-the-art methods. As can be seen in Figure 4, most existing methods do not perform well on singing music. Equal loudness change based method produces an F-measure of 71.0% under the best set of parameters. Both the phase-based and pitch-based methods perform badly because the singing voice's pitch track is very unstable, and contains many noisy segments.

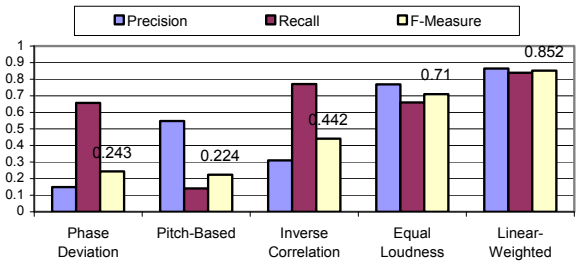

Figure 4: Results of linear-weighted sum detection function compared with state-of-the-art methods

The parameters for the system, especially those used in the GMM and individual features, were determined through extensive experimentation.

7. DISCUSSIONS AND FUTURE WORK

In our system, we have employed both feature-level and decision-level fusion. Feature-level fusion works well, but the higher dimensionality necessitates higher volume of training data. Our system was trained with only 560 onsets, due to the laborious process annotating pieces by hand. More training and testing data ought to be used to validate the system's performance.

We utilized only 4 pairs of GMMs: for MFCC, LPCC, ERB-bands, and a combined feature. In reality, any number of feature types can be used, and fusion can be

done at the feature-level (e.g. concatenation) and/or at decision-level (e.g. linear weighted sum). As a rough guide, more feature types and detection functions usually produce better onset detection results. This is provided the features extracted are representative of the onset classification problem, i.e. there should be a discernible difference in feature between onset and non-onset frames.

Weight-assignments for linear-weighted sum fusion are usually based on heuristics, and require lengthy experiments to optimize. Even though it generally works well, the simple linear weighting method can cause false peaks in detection functions to propagate into the combined detection function, making the overall detection function noisy. Other decision-level fusion techniques exist, and ought to be explored and tested.

We also look forward to expanding the system by incorporating note segmentation, as well as more functions in the post-processing section.

8. CONCLUSION

As shown by our experiments, the proposed supervised machine-learning approach based on GMM modeling produces higher accuracy for singing onset detection. Clearly, the system produces better results than state-of-the-art singing voice onset detection algorithms. Further improvements by decision-level fusion of the system boost the overall accuracy and completeness of the system. The singing onset detection problem cannot be considered solved by our system, but its potential is promising.

9. ACKNOWLEDGEMENT

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FIVE APPROACHES TO COLLECTING TAGS FOR MUSIC

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ABSTRACT

We compare five approaches to collecting tags for music: conducting a survey, harvesting social tags, deploying annotation games, mining web documents, and autotagging audio content. The comparison includes a discussion of both scalability (financial cost, human involvement, and computational resources) and quality (the cold start problem & popularity bias, strong vs. weak labeling, vocabulary structure & size, and annotation accuracy). We then describe one state-of-the-art system for each approach. The performance of each system is evaluated using a tag-based music information retrieval task. Using this task, we are able to quantify the effect of popularity bias on each approach by making use of a subset of more popular (short-head) songs and a set of less popular (long-tail) songs. Lastly, we propose a simple hybrid context-content system that combines our individual approaches and produces superior retrieval results.

1 INTRODUCTION

Tags are text-based tokens, such as "happy", "classic rock", and "distorted electric guitar", that can be used to annotate songs. They represent a rich source of semantic information that is useful for text-based music retrieval (e.g., [19]), as well as recommendation, discovery, and visualization [11]. Tags can be collected from humans using surveys [19, 5], social tagging websites [13], or music annotation games [20, 14, 12]. They can also be generated by text mining web-documents [10, 24] or by autotagging audio content [19, 7, 21]. In Section 2, we introduce key concepts associated with tag collection, and use them to highlight the strengths and weaknesses of each of these five approaches. In Section 3, we describe one implementation of a system for each approach and evaluate its performance on a tag-based music retrieval task. In the final section, we describe a simple hybrid system that combines the output from each of our individual systems.

2 COLLECTING TAGS

In this section, we describe five approaches to collecting music tags. Three approaches (surveys, social tags, games) rely on human participation, and as such, are expensive in terms of financial cost and human labor. Two approaches (text mining, autotagging) rely on automatic methods that are computationally intense, but require less direct human involvement.

There are a number of key concepts to consider when comparing these approaches. The *cold start problem* refers to the fact songs that are not annotated cannot be retrieved. This problem is related to *popularity bias* in that popular songs (in the *short-head*) tend to be annotated more thoroughly than unpopular songs (in the *long-tail*) [11]. This often leads to a situation in which a short-head song is ranked above a long-tail song despite the fact that the long-tail song may be more semantically relevant. We prefer an approach that avoids the cold start problem (e.g., autotagging). If this is not possible, we prefer approaches in which we can explicitly control which songs are annotated (e.g., survey, games), rather than an approach in which only the more popular songs are annotated (e.g., social tags, web documents).

A *strong labeling* [3] is when a song has been explicitly labeled or not labeled with a tag, depending on whether or not the tag is relevant. This is opposed to a *weak labeling* in which the absence of a tag from a song does not necessarily indicate that the tag is not relevant. For example, a song may feature drums but is not explicitly labeled with the tag "drum". Weak labeling is a problem if we want to design a MIR system with high recall, or if our goal is to collect a training data set for a supervised autotagging system that uses discriminative classifiers (e.g., [7, 24]).

It is also important to consider the size, structure, and extensibility of the tag vocabulary. In the context of text-based music retrieval, the ideal vocabulary is a large and diverse set of semantic tags, where each tag describes some meaningful attribute or characterization of music. In this paper, we limit our focus to tags that can be used consistently by a large number of individuals when annotating novel songs based on the audio content alone. This does not include tags that are personal (e.g., "seen live"), judgmental (e.g., "horrible"), or represent external knowledge about the song (e.g., geographic origins of an artist). It should be noted that these tags are also useful for retrieval (and recommendation) and merit additional attention from the MIR community.

A tag vocabulary can be *fixed* or *extensible*, as well as *structured* or *unstructured*. For example, the tag vocabulary associated with a survey can be considered fixed and structured since the set of tags and the grouping of tags into coherent semantic categories (e.g., genres, instruments, emotions, usages) is predetermined by experts using domain knowledge [19, 20]. By contrast, social tagging communities produce a vocabulary that is extensible since any user can

Approach	Strengths	Weaknesses
Survey	custom-tailored vocabulary high-quality annotations strong labeling	small, predetermined vocabulary human-labor intensive time consuming approach lacks scalability
Social Tags	collective wisdom of crowds unlimited vocabulary provides social context	create & maintain popular social website ad-hoc annotation behavior, weak labeling sparse/missing in long-tail
Game	collective wisdom of crowds entertaining incentives produce high-quality annotations fast paced for rapid data collection	"gaming" the system difficult to create viral gaming experience listening to short-clips, rather than entire songs
Web Documents	large, publicly-available corpus of relevant documents no direct human involvement provides social context	noisy annotations due to text-mining sparse/missing in long-tail weak labeling
Autotags	not affected by cold-start problem no direct human involvement strong labeling	computationally intensive limited by training data based solely on audio content

Table 1. Strengths and weaknesses of tag-based music annotation approaches

suggest any free-text token to describe music. This vocabulary is also unstructured since tags are not organized in any way. In general, we prefer an extensible vocabulary because a fixed vocabulary limits text-based retrieval to a small set of predetermined tags. In addition, a structured vocabulary is advantageous since the ontological relationships (e.g., genre hierarchies, families of instruments) between tags encode valuable semantic information that is useful for retrieval.

Finally, the accuracy with which tags are applied to songs is perhaps the most important point of comparison. Since there is no ideal ground truth and listeners do not always agree whether (or to what degree) a tag should be applied to a song (i.e., 'the subjectivity problem' [15]), evaluating accuracy can be tricky. Intuitively, it is preferable to have trained musicologists, rather than untrained non-experts, annotate a music corpus. It is also advantageous to have multiple individuals, rather then a single person, annotate each song. Lastly, individuals who are given incentives to provide good annotations (e.g., a high score in a game) may provide better annotations than unmotivated individuals.

2.1 Conducting a Survey

Perhaps the most well-known example of the music annotation survey is Pandora's [1] "Music Genome Project" [5, 23]. Pandora uses a team of approximately 50 expert music reviewers (each with a degree in music and 200 hours of training) to annotate songs using structured vocabularies of between 150-500 'musically objective' tags depending on the genre of the music [8]. Tags, such as "Afro-Latin Roots", "Electric Piano Riffs" and "Political Lyrics", can be considered objective since, according to Pandora, there is a high level of inter-reviewer agreement when annotating the same song. Between 2000 and 2007, Pandora annotated over 600,000 songs [23]. Currently, each song takes between 20 to 30 minutes to annotate and approximately 15,000 new songs are annotated each month. While this labor-intensive approach results in high-quality annotations, Pandora must be very selective of which songs they choose to annotate given that there are already millions of songs by millions of artists [2].

Pandora, as well as companies like Moodlogic [3] and All Media Guide (AMG) [4], have devoted considerable amounts of money, time and human resources to annotate their music databases with high-quality tags. As such, they are unlikely to share this data with the MIR research community. To remedy this problem, we have collected the CAL500 data set of annotated music [19]. This data set contains one song from 500 unique artists each of which have been manually annotated by a minimum of three non-expert reviewers using a structured vocabulary of 174 tags. While this is a small data set, it is strongly labeled, relies on multiple reviews per song, and as such, can be used as a standard data set for training and/or evaluating tag-based music retrieval systems.

2.2 Harvesting Social Tags

Last.fm [5] is a music discovery website that allows users to contribute *social* tags through a text box in their audio player interface. By the beginning of 2007, their large base of 20 million monthly users have built up an unstructured vocabulary of 960,000 free-text tags and used it to annotated millions of songs [16]. Unlike the Pandora and AMG, Last.fm makes much of this data available to the public through their Audiocrobbler [6] site. While this data is a useful resource for the MIR community, Lamere and Celma [11] point out a number of problems with social tags. First, there is often a sparsity of tags for new and obscure artists (cold start problem / popularity bias). Second, most tags are used to annotate artists rather than individual songs. This is problematic since we are interested in retrieving semantically relevant songs from eclectic artists. Third, individuals use

[1] www.pandora.com

[2] In February 2008, Last.fm reported that their rapidly growing database consisted of 150 million songs by 16 million artists.

[3] http://en.wikipedia.org/wiki/MoodLogic

[4] www.allmusic.com

[5] www.last.fm

[6] http://www.audioscrobbler.net/

ad-hoc techniques when annotating music. This is reflected by use of polysemous tags (e.g., "progressive"), tags that are misspelled or have multiple spellings (e.g., "hip hop", "hip-hop"), tags used for self-organization (e.g., "seen live"), and tags that are nonsensical. Finally, the public interface allows for malicious behavior. For example, any individual or group of individuals can annotate an artist with a misleading tag.

2.3 Playing Annotation Games

At the 2007 ISMIR conference, music annotation games were presented for the first time: ListenGame [20], Tag-a-Tune [12], and MajorMiner [14]. ListenGame is a real-time game where a large group of users is presented with a song and a list of tags. The players have to choose the best and worst tags for describing the song. When a large group of players agree on a tag, the song has a strong (positive or negative) association with the tag. This game, like a music survey, has the benefit of using a structured vocabulary of tags. It can be considered a strong labeling approach since it also collects information that reflects negative semantic associations between tags and songs. Like the ESPGame for image tagging [22], Tag-a-Tune is a two-player game where the players listen to a song and are asked to enter "free text" tags until they both enter the same tag. MajorMiner is similar in nature, except the tags entered by the player are compared against the database of previously collected tags in an offline manner. Like social tagging, the tags collected using both games result in a unstructured, extensible vocabulary.

A major problem with this game-based approach is that players will inevitably attempt to *game* the system. For example, the player may only contribute generic tags (e.g., "rock", "guitar") even if less common tags provide a better semantic description (e.g., "grunge", "distorted electric guitar"). Also, despite the recent academic interest in music annotation games, no game has achieved large scale success. This reflects the fact that it is difficult to design a viral game for this inherently laborious task.

2.4 Mining Web Documents

Artist biographies, album reviews, and song reviews are another rich source of semantic information about music. There are a number of research-based MIR systems that collect such documents from the Internet by querying search engines [9], monitoring MP3 blogs [4], or crawling a music site [24]. In all cases, Levy and Sandler point out that such web mined corpora can be *noisy* since some of the retrieved webpages will be irrelevant, and in addition, much of the text content on relevant webpages will be useless [13].

Most of the proposed web mining systems use a set of one or more documents associated with a song and convert them into a single document vector (e.g., tf-idf representation) [10, 25]. This *vector space* representation is then useful for a number of MIR tasks such as calculating music similarity [25] and indexing content for a text-based music retrieval system [10]. More recently, Knees et. al. [9] have proposed a promising new web mining technique called *relevance scoring* as an alternative to the vector space approaches. Both relevance scoring and vector space approaches are subject to popularity bias since short-head songs are generally represented by more documents than long-tail songs.

2.5 Autotagging Audio Content

All previously described approaches require that a song be annotated by humans, and as such, are subject to the cold start problem. Content-based audio analysis is an alternative approach that avoids this problem. Early work on this topic focused (and continues to focus) on music classification by genre, emotion, and instrumentation (e.g., [21]). These classification systems effectively 'tag' music with class labels (e.g., 'blues', 'sad', 'guitar'). More recently, *autotagging* systems have been developed to annotate music with a larger, more diverse vocabulary of (non-mutually exclusive) tags [19, 7, 17]. In [19], we describe a generative approach that learns a Gaussian mixture model (GMM) distribution over an audio feature space for each tag in the vocabulary. Eck et. al. use a discriminative approach by learning a boosted decision stump classifier for each tag [7]. Finally, Sordo et. al. present a non-parametric approach that uses a content-based measure of music similarity to propagate tags from annotated songs to similar songs that have not been annotated [17].

3 COMPARING SOURCES OF TAGS

In this section, we describe one system for each of the tag collection approaches. Each has been implemented based on systems that have been recently developed within the MIR research community [19, 9, 20]. Each produces a $|S|$x$|T|$ *annotation matrix* $\mathbf{X}$ where $|S|$ is the number of songs in our corpus and $|T|$ is the size of our tag vocabulary. Each cell $x_{s,t}$ of the matrix is proportional to the strength of semantic association between song s and tag t.

We set $x_{s,t} = \emptyset$ if the relationship between song s and tag t is missing (i.e., unknown). If the matrix $\mathbf{X}$ has many empty cells, then we refer to the matrix as *sparse*, otherwise we refer to it as *dense*. Missing data results from both weak labeling and the cold start problem. Sparsity is reflected by the *tag density* of a matrix which is defined as the percentage of non-empty elements of a matrix.

Our goal is to find a tagging system that is able to accurately retrieve (i.e., rank-order) songs for a diverse set of tags (e.g., emotions, genres, instruments, usages). We quantitatively evaluate music retrieval performance of system a by comparing the matrix $\mathbf{X}^a$ against the CAL500 matrix $\mathbf{X}^{\mathrm{CAL500}}$ (see Section 2.1). The $\mathbf{X}^{\mathrm{CAL500}}$ matrix is a binary matrix where $x_{s,t} = 1$ if 80% of the individuals annotate song s with tag t, and 0 otherwise (see Section V.a of [19] for details). For the experiments reported in this section, we use a subset of 109 of the original 174 tags. [7] We will assume that the subset of 87 songs from the Magnatunes [6]

[7] We have merged genre-best tags with genre tags, removed instrument-solo tags, removed some redundant emotion tags, and pruned other tags that are used to annotate less than 2% of the songs. For a complete list of tags, see http://cosmal.ucsd.edu/cal.

Approach	Songs	Tag Density	AROC	Avg. Prec	R-Prec	Top10 Prec
Survey (CAL500)	All Songs	1.00	1.00	1.00	1.00	0.97
Ground Truth	Long Tail	1.00	1.00	1.00	1.00	0.57
Baseline	All Songs	1.00	0.50	0.15	0.14	0.13
Random	Long Tail	1.00	0.50	0.18	0.15	0.12
Social Tags	All Songs	0.23	0.62	0.28	0.30	0.37
Last.fm	Long Tail	0.03	0.54	0.24	0.20	0.19
Game ListenGame†	All Songs	0.37	0.65	0.28	0.28	0.32
Web Documents	All Songs	0.67	0.66	0.29	0.29	0.37
SS-WRS	Long Tail	0.25	0.56	0.25	0.20	0.18
Autotags	All Songs	1.00	0.69	0.29	0.29	0.33
SML	Long Tail	1.00	0.70	0.34	0.30	0.27
Rank-based	All Songs	1.00	0.74	0.32	0.34	0.38
Interleaving (RBI)	Long Tail	1.00	0.71	0.33	0.27	0.28

Table 2. Tag-based music retrieval: Each approach is compared using all *CAL500* songs and a subset of 87 more obscure *long-tail* songs from the Magnatunes dataset. *Tag Density* represents the proportion of song-tag pairs that have a non-empty value. The four evaluation metrics (*AROC, Average Precision, R-Precision, Top-10 Precision*) are found by averaging over 109 tag queries. †Note that ListenGame is evaluated using half of the CAL500 songs and that the results do not reflect the realistic effect of the popularity bias (see Section 3.2).

collection that are included in the CAL500 data set are representative of long-tail music. As such, we can use this subset to gauge how the various tagging approaches are affected by popularity bias. [8]

Each system is compared to the CAL500 data set using a number of standard information retrieval (IR) evaluation metrics [9]: area under the receiver operation characteristic curve (AROC), average precision, R-precision, and Top-10 precision. An ROC curve is a plot of the true positive rate as a function of the false positive rate as we move down this ranked list of songs. The area under the ROC curve (AROC) is found by integrating the ROC curve and is upper-bounded by 1.0. A random ranking of songs will produce an expected AROC score of 0.5. Average precision is found by moving down our ranked list of test songs and averaging the precisions at every point where we correctly identify a relevant song. R-Precision is the precision of the top R-ranked songs where R is the total number of songs in the ground truth that have been annotated with a given tag. Top-10 precision is the precision after we have retrieved the top 10 songs for a given tag. This metric is designed to reflect the 10 items that would be displayed on the first results page of a standard Internet search engine.

Each value in Table 2 is the mean of a metric after averaging over all 109 tags in our vocabulary. That is, for each tag, we rank-order our 500 song data set and calculate the value of the metric using CAL500 data as our ground truth. We then compute the average of the metric using the 109 values from the 109 rankings.

3.1 Social Tags: Last.fm

For each of our 500 songs, we attempt to collect two lists of social tags from the Last.fm Audioscobbler website. One list is related specifically to the song and the other list is related to the artist. For the song list, each tag has a score ($x_{s,t}^{\text{Last.fm\_Song}}$) that ranges from 0 (low) to 100 (high) and is a secret function (i.e., trade secret of Last.fm) of both the number and diversity of users who have annotated song s with tag t. For the artist list, the tag score ($x_{s,t}^{\text{Last.fm\_Artist}}$) is again a secret function that ranges between 0 and 100, and reflects both tags that have been used to annotate the artist or songs by the artist. We found one or more tags for 393 and 472 of our songs and artists, respectively. This included at least one occurrence of 71 and 78 of the 109 tags in our vocabulary. While this suggests decent coverage, tag densities of 4.6% and 11.8%, respectively, indicate that the annotation matrices, $\mathrm{X}^{\text{Last.fm\_Song}}$ and $\mathrm{X}^{\text{Last.fm\_Artist}}$, are sparse even when we consider mostly short-head songs. These sparse matrices achieve AROC of 0.57 and 0.58.

To remedy this problem, we create a single Last.fm annotation matrix by leveraging the Last.fm data in three ways. First, we match tags to their synonyms. [9] For example, a song is considered to be annotated with 'down tempo' if it has instead been annotated with 'slow beat'. Second, we allow wildcard matches for each tag. That is, if a tag appears as a substring in another tag, we consider it to be a wildcard match. For example, "blues" matches with "delta electric blues", "blues blues blues", "rhythm & blues". Although synonyms and wildcard matches add noise, they increase the respective densities to 8.6% and 18.9% and AROC performance to 0.59 and 0.59. Third, we combine the song and artist annotation matrices in one annotation matrix:

$$\mathrm{X}^{\text{Last.fm}} = \mathrm{X}^{\text{Last.fm\_Song}} + \mathrm{X}^{\text{Last.fm\_Artist}}.$$

This results in a single annotation matrix that has a density of 23% and AROC of 0.62. 95 of the 109 tags are represented at least once in this matrix. However, the density for the Magnatunes (e.g., long-tail) songs is only 3% and produces retrieval results that are not much better than random.

[8] It should be noted that 87 songs is a small sample.

[9] Synonyms are determined by the author using a thesaurus and by exploring the Last.fm tag vocabulary.

3.2 Games: ListenGame

In [20], Turnbull et al. describe a music annotation game called ListenGame in which a community of players listen to a song and are presented with a set of tags. Each player is asked to vote for the single *best* tag and single *worst* tag to describe the music. From the game, we obtain the annotation matrix $\mathbf{X}^{\mathbf{Game}}$ by letting

$$[\mathbf{X}^{\mathrm{Game}}]_{s,t} = \#(\text{best votes}) - \#(\text{worst votes})$$

when song s and tag t are presented to the players.

During a two-week pilot study, 16,500 annotations (best and worst votes) were collected for a random subset of 250 CAL500 songs. Each of the 27,250 song-tag pairs were presented to users an average of 1.8 times. Although this represents a very small sample size, the mean AROC for the subset of 250 songs averaged over the 109-tag vocabulary is 0.65. Long-tail and short-head results do not accurately reflect the real-world effect of popularity bias since all songs were selected for annotation with equal probability. As such, these results have been omitted.

3.3 Web Documents: Weight-based Relevance Scoring

In order to extract tags from a corpus of web documents, we adapt the relevance scoring (RS) algorithm that has recently been proposed by Knees et. al. [9]. They have shown this method to be superior to algorithms based on vector space representations. To generate tags for a set of songs, the RS works as follows:

1. **Collect Document Corpus:** For each song, repeatedly query a search engine with each song title, artist name, or album title. Collect web documents in search results. Retain the (many-to-many) mapping between songs and documents.
2. **Tag Songs:** For each tag
   (a) Use the tag as a query string to find the relevant documents, each with an associated *relevance weight* (defined below) from the corpus.
   (b) For each song, sum the relevance scores for all the documents that are related to the song.

We modify this algorithm in two ways. First, the relevance score in [9] is inversely proportional to the rank of the relevant document. We use a weight-based approach to relevance scoring (WRS). The relevance weight of a document given a tag can be a function of the number of times the tag appears in the document (tag-frequency), the number of documents with the tag (document frequency), the number of total words in the document, the number of words or documents in the corpus, etc. For our system, the relevance weights are determined by the MySQL match function. [10]

We calculate an entry of the annotation matrix $\mathbf{X}^{\mathrm{WRS}}$ as,

$$\mathbf{X}^{\mathrm{WRS}}_{s,t} = \sum_{d \in D_t} w_{d,t} I_{d,s}$$

[10] http://dev.mysql.com/doc/refman/5.0/en/fulltext-natural-language.html

where D_t is the set of relevant documents for tag t, $w_{d,t}$ is the relevance weight for document d and tag t, and $I_{d,s}$ is an indicator variable that is 1 if document d was found when querying the search engine with song s (in Step 1) and 0 otherwise. We find that weight-based RS (WRS) produces a small increase in performance over rank-based RS (RRS) (AROC of 0.66 vs. 0.65). In addition, we believe that WRS will scale better since the relevance weights are independent of the number of documents in our corpus.

The second modification is that we use *site-specific* queries when creating our corpus of web documents (Step 1). That is, Knees et. al. collect the top 100 documents returned by Google when given queries of the form:

- "<artist name>" music
- "<artist name>" "<album name>" music review
- "<artist name> " "<song name>" music review

for each song in the data set. Based on an informal study of the top 100 webpages returned by non-site-specific queries, we find that many pages contain information that is only slightly relevant (e.g., music commerce site, ticket resellers, noisy discussion boards, generic biographical information). By searching music-specific sites, we are more likely to find detailed music reviews and in-depth artist biographies. In addition, the webpages at sites like Pandora and AMG All Music specifically contain useful tags in addition to natural language content.

We use site-specific queries by appending the substring 'site:<music site url>' to the three query templates, where <music site url> is the url for a music website that is known to have high quality information about songs, albums or artists. These sites include allmusic.com, amazon.com, bbc.co.uk, billboard.com, epinions.com, musicomh.com, pandora.com, pitchforkmedia.com, rollingstone.com, wikipedia.org. For these 10 music sites and one non-site-specific query, we collect and store the top 10 pages returned by the Google search engine. This results in a maximum of 33 queries and a maximum of 330 pages per song. On average, we are only able to collect 150 webpages per song since some of the long-tail songs are not well represented by these music sites.

Our *site-specific weight-based relevance scoring* (SS-WRS) approach produces a relatively dense annotation matrix (46%) compared with the approach involving Last.fm tags. However, like the Last.fm approach, the density of the annotation matrix is greatly reduced (25%) when we consider only long-tail songs.

3.4 Autotagging: Supervised Multiclass Labeling

In [19], we use a supervised multiclass labeling (SML) model to automatically annotate songs with a diverse set of tags based on audio content analysis. The SML model is parameterized by one Gaussian mixture model (GMM) distribution over an audio feature space for each tag in the vocabulary. The parameters for the set of GMMs are trained using annotated training data. Given a novel audio track, audio features

are extracted and their likelihood is evaluated using each of the GMMs. The result is a vector of probabilities that, when normalized, can be interpreted as the parameters of a multinomial distribution over the tag vocabulary. This *semantic multinomial* distribution represents a compact and interpretable index for a song where the large parameter values correspond to the most likely tags.

Using 10-fold cross validation, we can estimate a semantic multinomial for each of the CAL500 songs. By stacking the 50 test set multinomials from each of the 10 folds, we can construct a strongly-labeled annotation matrix $\mathbf{X}^{\mathrm{SML}}$ that is based purely on the audio content. As such, this annotation matrix is dense and not affected by the cold start problem.

3.5 Summary

Comparing systems using a two-tailed, paired t-test (N = 109, $\alpha = 0.05$) on the AROC metric, we find that all pairs of the four systems are significantly different, with the exception of Game and Web Documents. [11] If we compare the systems using the other three metrics (Average Precision, R-Precision, and Top 10 Precision), we no longer find statistically significant differences. It is interesting that Social Tags and Web Documents (0.37) have slightly better Top 10 precision than Autotags (0.33). This reflects the fact that for some of the more common individual tags, we find that Social Tags and Web Documents have exceptional precision at low recall levels. For both Web Documents and Social Tags, we find significant improvement in retrieval performance of short-head songs over long-tail songs. However, as expected, there is no difference for Autotags. This confirms the intuition that systems based on web documents and social tags are influenced by popularity bias, whereas content-based autotagging systems are not.

4 COMBINING SOURCES OF TAGS

While the purpose of this paper is to compare various approaches for collecting tags for music, our ultimate goal is to combine these approaches in order to create a more powerful tag-based music retrieval system. For example, if we interleave the top ranked songs from each of the four approaches (See Rank-based Interleaving (RBI) in Table 2), we observe a significant increase in performance (AROC 0.74) over the performance of the best single approach (Autotags with AROC = 0.69). Our improvement is consistent with the findings of Yoshii et. al. [26] and Aucountier et. al. [1], both of whom have recently proposed hybrid context-content systems for music recommendation and music classification, respectively. We explore alternative hybrid systems in some of our related work [2, 18].

5 ACKNOWLEDGEMENTS

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[11] Note that when we compare each system with the Game system, we compare both systems using the reduced set of 250 songs.

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UNCOVERING AFFINITY OF ARTISTS TO MULTIPLE GENRES FROM SOCIAL BEHAVIOUR DATA

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ABSTRACT

In organisation schemes, musical artists are commonly identified with a unique 'genre' label attached, even when they have affinity to multiple genres. To uncover this hidden cultural awareness about multi-genre affinity, we present a new model based on the analysis of the way in which a community of users organise artists and genres in playlists.

Our work is based on a novel dataset that we have elaborated identifying the co-occurrences of artists in the playlists shared by the members of a popular Web-based community, and that is made publicly available. The analysis defines an automatic social-based method to uncover relationships between artists and genres, and introduces a series of novel concepts that characterises artists and genres in a richer way than a unique 'genre' label would do.

1 INTRODUCTION

Musical genres have typically been used as a 'high level' classification of songs and artists. However, many musical artists cannot be described with a single genre label. Artists can change style throughout their career, and perform songs that do not entirely fall into single categories such as 'Rock', 'R&B' or 'Jazz'. In general terms, artists can be said to have a certain degree of *affinity* to each specific genre.

In this paper we propose to model relationships from artists to genres as *fuzzy sets*, where the *membership degree* indicates the affinity of an artist to a genre. This is in contrast to conventional labelling approaches, in which artists either belong or do not belong to a genre, and allows for a sentence like "Madonna is Pop and R&B, not Jazz" to be rephrased as "Madonna belongs with a membership degree of 0.8 to Pop genre, with 0.6 to R&B and with 0.1 to Jazz".

To uncover the affinity of different artists to different genres, we follow a *social* approach: we exploit information about how people aggregate and organise artists and genres in *playlists*. Playlists are the predominant form of casual music organisation. When playlists are collected in aggregate over a large sampling of the music listening populace, they reflect current cultural sensibilities for the association of popular music. Our assumption about *large repositories* of playlists is that when two artists or genres co-occur together and closely in many playlists, they have some form of shared cultural *affinity*. For example, if we observe that a large number of songs by an artist x co-occur often and closely with songs whose artists are classified in the 'Jazz' genre, then we argue that x has a certain affinity with Jazz, independently from the original 'genre' label attached to x.

On the basis of this assumption, we build a model of *multi-genre affinity* for artists, grounded on data about the *usage* of music objects by real people. We suggest this kind of analysis can provide results which are complementary to those provided by other content-based or social techniques, which we discuss in the following subsection.

In this paper, we first discuss different approaches towards musical genre analysis, and provide arguments for the consideration of a community-based approach. Next, we report on the analysis we performed on a novel dataset, which lists the co-occurrences of 4,000 artists in a large collection of playlists, to uncover associations between artists and genres. Finally, we depict possible applications, such as affinity-based generation of 'smooth sequences' of songs.

1.1 Related work about genre analysis

Genre analysis has been a constant theme in the MIR community, and researchers have had some success in classifying or predicting genre using either *content-analytic* approaches or *social behaviour* approaches.

Content-analytic methods are broken into two distinct approaches. The 'acoustic-analytic' approach was introduced by Tzanetakis and Cook [9], and investigates the acoustic signature of music according to a battery of relevant cognitive and musicological descriptors, such as beats per minute, timbre, and frequency spectrum. The 'score-analytic' approach concerns itself with symbolic representations of music as sequences of notes and events making up a musical score, generally as MIDI files, as in McKay and Fujinaga [6].

Social behaviour approaches classify music objects

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based upon the actions that a community of people perform with them. A large amount of effort has been directed towards harvesting textual correlations (of names of artists, songs, etc.) that co-occur on public Web sites. This has been the focus of Schedl et al. [8], who analysed artist-based term co-occurrences, Knees et al. [4], and Whitman and Smaragdis [10]. Text-based Web mining, however, can suffer from a lack of precision, mistaking band name terms with non-associated content as mentioned in [8].

Other attempts at providing richer representations for musical artists include describing an artist with a set of tags [2], with a set of moods [3] or identifying term profiles that characterise a specific cluster of similar artists [4]. All of these techniques maintain a 'Boolean' approach, identifying whether each artist matches specific tags, moods or clusters, while we seek to identify 'fuzzy' membership values relating each artist to each genre.

The social-based approach that we follow is the analysis of co-occurrences in playlists, which is currently not a common method of analysis. This is because playlists are not often publicly indexed or made available. Previously, a smaller public collection of about 29,000 playlists was used by Logan et. al [5] to analyse a set of 400 popular artists, and by Cano and Koppenberger [1] to provide a study of artists networks, while we work on a novel and much larger dataset of 1,030,068 human-compiled playlists.

Playlists have a special utility for social analysis of genres in that they are *specific* to music, with no extraneous "noise" information such as on arbitrary Web pages. They are also *intentional* in their creation, as opposed to a simple record of an individual's play history which may have been generated randomly.

2 A DATASET OF ARTISTS CO-OCCURRENCES

MusicStrands (http://music.strands.com) is a Web-based music recommendation community which allows members to publish playlists from their personal media players, such as Apple iTunes and Windows Media Player. Every published playlist is made of a sequence of IDs that univocally identify the songs and artists that the playlist contains.

On July 31st, 2007, we gathered the 1,030,068 user playlists published so far. We discarded playlists made by a single song or by a single artist. We also removed any information related to the creators (user name, creation date, playlist title, rating) to focus on the sequential nature of playlists in terms of co-occurrences of artists.

For each pair of artists (x, y), we extracted the number of times that a song by x occurs together with a song by y, and at what *distance* (i.e., how many other songs are in between). To reduce the dimension of the dataset, we considered only the co-occurrences of songs from the 4,000 most popular artists, where the popularity of an artist equals the number of playlists in which occurs. We also excluded non-specific popular artist labels such as "Various Artists" or "Original Soundtrack", and any co-occurrence with a distance larger than 2, to obtain the dataset sketched in Table 1, where each record contains: the ID of the first artist x, the ID of the second artist y, and the numbers $d_0(x, y)$, $d_1(x, y)$, and $d_2(x, y)$ of playlists where y co-occurs after x at distance 0, 1, and 2 respectively.

x	y	$d_0(x,y)$	$d_1(x,y)$	$d_2(x,y)$
11	11	137	96	77
11	27	0	0	1
11	91	0	1	2
...	...	...	...	...

Table 1. A portion of the provided dataset

An auxiliary table in the dataset lists the 4,000 artists IDs, together with their genre IDs and their Musicbrainz artist IDs. Both artist IDs and genre IDs can be easily translated to actual names using the OpenStrands developers API. [1]

Thanks to the support of MusicStrands, we now make public the complete dataset that we have elaborated, which is freely available to researchers wishing to perform social based analysis on music behaviour data from real users. [2]

3 GENRES AND AFFINITY

The artists in the previously described dataset already have a genre label attached. However, our motivation is to provide a richer representation, which overcomes the limitation of having only one genre per artist.

We denote with $\mathcal{G}$ the set of m genres, and with $\mathcal{A}$ the set of n artists in the dataset, where $m = 26$ and n = 4,000. Our goal is to describe each artist x as a vector $[M_x(g_1), M_x(g_2), \ldots, M_x(g_m)]$, where each value $M_x(g_i) \in [0, 1]$ indicates how much x has affinity to the genre g_i. We call $M_x(g_i)$ the **genre affinity degree** of artist x to genre g_i.

The process to calculate the genre affinity degrees is the following. First we measure the association from each artist $x \in \mathcal{A}$ to each other artist, based on their co-occurrences in the authored playlists. Then we aggregate these associations *by genre* to obtain the degree in which each artist x is associated with each genre $g \in \mathcal{G}$. Finally we combine artist-to-artist and artist-to-genre associations to measure the degree $M_x(g)$ of affinity between artist x and genre g.

The first step is to consider how much an artist x is associated with *any other* artist y. Let $\mathcal{X}$ be set of $n - 1$ artists other than x: $\mathcal{X} = \mathcal{A} - \{x\}$. We define the **artist-to-artist** association $A_x : \mathcal{X} \rightarrow \mathbb{R}$ as an aggregation of the number

[1] API available at http://my.strands.com/openstrands

[2] Dataset available at http://labs.strands.com/music/affinity

of co-occurrences of x with any artist $y \in \mathcal{X}$:

$$\begin{aligned} A_x(y) = \quad & \alpha \cdot [d_0(x,y) + d_0(y,x)] \\ + \quad & \beta \cdot [d_1(x,y) + d_1(y,x)] \\ + \quad & \delta \cdot [d_2(x,y) + d_2(y,x)] \,. \end{aligned}$$

This metric aggregates the number of playlists in which artists x and y co-occur, with different weights according to the *distance* at which x and y occur. Co-occurrences at distance 0 (d_0), distance 1 (d_1), and distance 2 (d_2) are weighted with α, β and δ respectively. These parameters, with values in $[0, 1]$, can be tuned to reflect the importance that the authors of the playlists assign to the distance between associated artists in their playlists. In the case of MusicStrands, we have observed that some authors compile playlists grouping together associated artists *one after the other* (for example, DJs for dance playlists), while most users create playlists as *unordered sets* of associated songs, where the distance between two artists is not related to their association. Based on this mixed behaviour, in our analysis, we have assigned values of $\alpha = 1$, $\beta = 0.8$, and $\delta = 0.64$.

According to the metric defined, the values of A_x are heavily influenced by the popularity of x. In fact, an artist x who is present in many playlists co-occurs with many artists, and has a generally higher artist-to-artist association than an uncommon artist. Therefore, we perform a normalisation and obtain an artist-to-artist association $\widehat{A}_x : \mathcal{A} \rightarrow [-1, 1]$ independent from the popularity of x, such that:

$$\widehat{A_x}(y) = \frac{A_x(y) - \overline{A_x}}{|max(A_x(y) - \overline{A_x})|}$$

where $\overline{A_x} = \frac{1}{n-1} \sum_{y \in \mathcal{X}} A_x(y)$, and $\widehat{A}_x(x) = 0$ by convention. A positive (resp. negative) value $\widehat{A_x}(y)$ indicates that the artist-to-artist association of x with y is above (resp. below) the average artist-to-artist association of x.

Now we turn to estimate the affinity of artists with respect to genres. First, let us consider how much an artist x is associated with any genre g. We define the association $P_x : \mathcal{G} \rightarrow \mathbb{R}$ by cumulating the artist-to-artist association A_x from x to *any artist of genre* g:

$$P_x(g) = \sum_{y \in \mathcal{X} : \gamma(y) = g} A_x(y)$$

where $\gamma(y)$ denotes the genre of y. Again, we normalise this association to counter-effect the uneven distribution of genres in the playlists (e.g., Rock/Pop artists occur in almost any playlist, while Folk artists are quite rare). The result is a normalised association $\widehat{P_x}(g) : \mathcal{G} \rightarrow [0, 1]$, independent from the popularity of genre g, defined as:

$$\widehat{P_x}(g) = \frac{\sum_{y \in \mathcal{X} : \gamma(y) = g} A_x(y)}{\sum_{y \in \mathcal{X}} A_x(y)} \,.$$

This function $\widehat{P}$ measures the degree in which artists associated with artist x belong to the genre g. Finally we obtain the **genre affinity degree** $M_x : \mathcal{G} \rightarrow [0, 1]$ by weighting the association $\widehat{P}$ with the artist-to-artist association $\widehat{A}$, and normalising the result to the range $[0, 1]$:

$$M_x(g) = \frac{1}{2} \left(\frac{\sum_{y \in \mathcal{A}} \widehat{A_x}(y) \widehat{P_y}(g)}{n} + 1 \right) \,.$$

Notice that $M_x(g)$ is high when artists that often co-occur with x belong to genre g and when artists that rarely co-occur with x do not belong to genre g. Each artist can be characterised by a **genre affinity vector** $[M_x(g_1), M_x(g_2), \ldots, M_x(g_m)]$ $\forall g_i \in \mathcal{G}$, which is a multi-dimensional representation of an artist in terms of genres.

For example, Table 2 shows , for three famous artists of the dataset labelled as 'Rock/Pop' (Madonna, Metallica and Bruce Springsteen), their artist-to-artist association $\widehat{A_x}(y)$, their association $\widehat{P_x}(g)$ and their genre affinity degrees $M_x(g)$ with respect to three popular genres (Rock/Pop, Country and R&B).

The 'Rock/Pop' genre is very common in our data (labelling 2,286 of the 4,000 artists in the dataset), so having these three artists generically labelled as 'Rock/Pop' is not very informative of their style. However, by observing the differences in their genre affinity degrees $M_x(g)$, we can spot their differences. For instance, Bruce Springsteen has a higher genre affinity to Country, while Madonna has a higher affinity to R&B. By plotting their $M_x(g)$ values with respect to these three genres on a graph, we can visually observe such difference in Fig. 1.

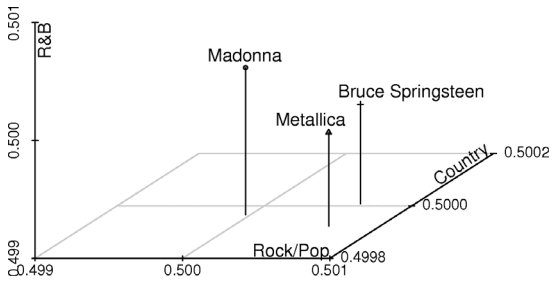

Figure 1. Genre affinity of 3 artists with 3 genres

4 ANALYSIS AND DISCUSSION

4.1 Genres and core artists

One advantage of describing artists in terms of affinity vectors, rather than with unique genre labels, is that we can identify how *central* an artist is to a genre. For each artist x, we define its **genre-centrality** $C_g : \mathcal{A} \rightarrow [0, 100]$ to a genre $g \in \mathcal{G}$ as the percentage of artists whose genre affinity to g

$\widehat{A}_x(y)$	Madonna	Metallica	Bruce Spr.
Mad.	0	0.0026	0.0516
Met.	0.0322	0	0.0106
B. Spr.	0.0602	0.0063	0

$\widehat{P}_x(g)$	Rock	Country	R&B
Mad.	0.628	0.028	0.065
Met.	0.892	0.013	0.034
B. Spr.	0.790	0.041	0.026

$M_x(g)$	Rock	Country	R&B
Mad.	0.49997	0.49998	**0.50007**
Met.	**0.50040**	0.49995	0.49997
B. Spr.	0.50038	**0.50001**	0.49991

Table 2. Associations and affinities between artists and genres, shown only for a subset of 3 artists and 3 genres

is lower or equal than the genre affinity of x to genre g:

$$C_g(x) = \frac{100}{n} \cdot card\{y \in \mathcal{A} : M_y(g) \leq M_x(g)\} \ .$$

For instance, the *Country*-centrality of Bruce Springsteen is 79.3%, because as many as 3,172 out of the 4,000 artists in the dataset have an affinity degree to Country lower than his, while his *R&B*-centrality is 39.5%, since only 1,580 artists have a genre affinity degree to R&B lower than his.

We will call those artists which are the most central to a genre g the **core artists** of g. Core artists are good representatives of a genre, since they occur very often in the set of playlists with artists from that genre and seldom with artists from other genres. For instance, we interestingly observe that the top core artists for the genre labelled 'Soundtrack' are James Horner, Alan Silvestri and Michael Giacchino, who are famous composers of original movie scores (e.g., James Horner's *"Titanic Original Soundtrack"*), and not Pop artists who have only sporadically performed famous songs which appeared in movies (e.g., Celine Dion's *"My Heart Will Go On"*).

The top core artists for other three genres are: (Folk) Luka Bloom, Greg Brown, The Chieftains; (Blues) Muddy Waters, Taj Mahal, Dr. John; (Jazz) Bob James, Donald Byrd, Peter White; (Electronic) Kaskade, Sasha, Junkie XL.

4.2 Centrality and cross-genre artists

Table 3 shows four artists originally labelled as Electronic and their genre-centrality with respect to four genres. From this table, we learn that St. Germain has a high *Jazz*-centrality (94%), Soul II Soul have a high *R&B*-centrality (94%) and M83 have a high *Rock/Pop*-centrality (95%). This is not surprising, since St. Germain is "a French electronica and *nu jazz* musician", [3] Soul II Soul is a "U.K. *R&B* collective", [4] and M83 offer "a luscious blend of shoegaze aesthetics, ambient *pop*, and progressive textures". [5] This is an example of how, using genre-centrality, we can obtain richer descriptions of the affinity of artists to multiple genres, without employing expert knowledge, but analysing the way in which each artists' works were *employed* by real users in playlists, in combination with other artists.

Table 3 also shows that Cameron Mizell, originally labelled as Electronic, has a higher genre-centrality to Jazz (99%) than to Electronic (90%). The reason is that Cameron Mizell is a cross-genre artist, whose creations mostly fall in the Electronic category, and mostly occur together with Jazz songs. Indeed, Cameron Mizell is labelled as 'Electronic' in MusicStrands, but as 'Jazz' according to other external musical sources, such as AMG (http://allmusic.com), iTunes Music Store (http://apple.com/itunes) and Amazon (http://amazon.com). This is not such a strange case, for several artists exist which behave as a sort of *musical bridge* from a genre to another. Formally, we call **bridge artist** any artist x whose original genre label $\gamma(x)$ differs from the genre $\phi(x)$ for which x has the highest genre-centrality, where $\phi(x) = g \in \mathcal{G} : C_g(x) \geq C_h(x)\ \forall h \in \mathcal{G}$.

$C_g(x)$	St. Germain	Soul II Soul	M83	C.Mizell
Electronic	98%	96%	98%	90%
Jazz	**94%**	10%	6%	**99%**
R&B	41%	**94%**	8%	62%
Rock/Pop	37%	1%	**95%**	17%

Table 3. Genre-centrality of 4 artists to 3 genres

Table 4 reports the number of bridge artists, aggregated by genre, for six of the 26 genres (with the main diagonal containing artists which are *not* bridge artists). According to our data, the distribution of bridge artists is uneven and depends on genres; for instance there are no bridge artists from Reggae to Country and vice versa, while there are 90 bridge artists from Rap to R&B (37% of all the Rap artists). This suggests that genres like Country and Reggae are "distinct", in the sense that artists from the two genres tend not to "play well together", while genres like R&B and Rap are "adjoining", meaning that artists from the two genres tend to "play well together". These abstract concepts are defined more precisely hereafter, with different typologies of associations between genres characterised.

$g \setminus h$	Country	Blues	Jazz	Reggae	R&B	Rap
Country	162	-	1	-	-	-
Blues	-	3	-	-	-	-
Jazz	-	2	101	-	-	-
Reggae	-	-	-	24	-	-
R&B	1	1	9	6	334	11
Rap	-	-	-	9	90	226

Table 4. Number of artists labelled as genre g, whose genre-centrality is maximum for genre h, for 6 different genres

[3] Excerpt from http://last.fm/music/St.+Germain
[4] Excerpt from http://music.strands.com/artist/15246
[5] Excerpt from http://music.aol.com/artist/m83/1655884

4.3 Correlation between two genres

After having analysed associations from artists to genres, we draw some observations about *how genres relate* with each other. The goal is to discern for each pair of genres (g, h) whether they are or not correlated. The idea is that two genres are correlated when artists belonging to one genre naturally tend to be played together with artists of the other genre. Using the genre affinity degree $M_x(g)$, we can more precisely state that two genres g and h are **correlated** (resp. **exclusive**) when artists with a high $M_x(g)$ (genre affinity degree to g) tend to have a high (resp. low) $M_x(h)$ (genre affinity degree to h); and two genres are **independent** when their affinity degrees do not show any sign of correlation.

As an example, consider Figure 2 (above) showing, for each of the 4,000 artists in the dataset, its affinity degree to Country (x axis), to Reggae (y axis), and its original genre label (point type). Artists with high affinity to Country (in the lower right section) tend to have a 'neutral' affinity to Reggae (with most values close to 0.5). Similarly, artists with high affinity to Reggae (in the upper left section) tend to have a 'neutral' affinity to Country. This is an example of independent genres, since the genre affinity degree to Country and to Reggae of an artist look unrelated. Figure 2 (below) shows instead that artists with high affinity to R&B often have a high affinity to Rap as well (in the upper right section), suggesting that the two genres are correlated.

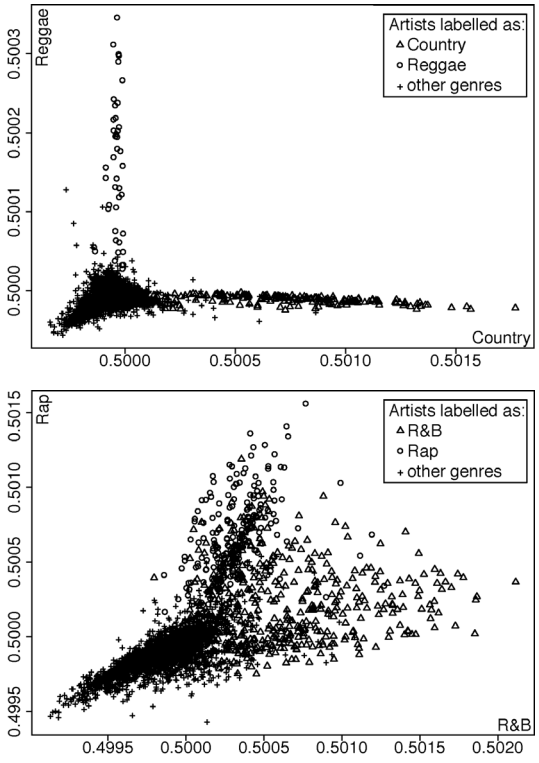

Figure 2. Comparing *Country* vs. *Reggae* (above), and *R&B* vs. *Rap* (below) genre-centralities

To measure the correlation between any two genres g and h, we employ Pearson's coefficient $\rho_{g,h} \in [-1, 1]$, which assesses the linear correlation between the affinity degree of artists from two genres, and has values close to 1 in the case of a positive relationship (i.e., artists with high affinity to g also have high affinity to genre h), and close to -1 in the case of a negative relationship (i.e., artists with high affinity to g have low affinity to genre h). Table 5 reports these coefficients for six genres, showing for instance that R&B and Rap (0.6) are positively correlated (as suggested by Fig. 2), as well as Jazz and Blues (0.4), while Blues and Rap (-0.2) are negatively correlated: artists with a high affinity to Blues tend to have a low affinity to Rap.

This social analysis of genres exposes cultural correlations between music that is often not present in an acoustic or score analysis. Most importantly, this analysis does not require expert human knowledge of genre relationships. Instead, relationships emerge from a larger cultural consensus.

$\rho_{g,h}$	Country	Blues	Jazz	Reggae	R&B	Rap
Country	1	0.2	0.1	0	0.1	-0.1
Blues	0.2	1	**0.4**	0.1	0	**-0.2**
Jazz	0.1	0.4	1	0.1	0.2	-0.1
Reggae	0	0.1	0.1	1	**0.4**	**0.4**
R&B	0.1	0	0.2	0.4	1	**0.6**
Rap	-0.1	-0.2	-0.1	0.4	0.6	1

Table 5. Pearson coefficient ρ for 6 genre-centrality vectors

5 APPLICATIONS

The analysis we have performed allows to describe artists in terms of affinity vectors and to uncover relations among genres. This knowledge is useful for many applications, some of which are presented hereafter.

First, consider the case of a Web radio, with a large repository of available music, and with the need to schedule different channels with different sequences of songs. The most common approach is to programme each channel with a randomly ordered selection of songs from the repository, matching the definition of each channel (e.g., a random sequence of jazz songs is played on a 'Jazz channel'). This method may work well for channels limited to a single genre, but may sound inadequate when multiple genres are permitted (e.g., in a channel generically defined as "Music from the '80s"), since unpleasant musical transitions could occur from one genre to another (e.g., a song by Abba, followed by a song by Metallica, and by a song by B.B. King). To procure *smooth* transitions from a song to the next one, the knowledge we have uncovered about affinity of each artist to each genre can be exploited, to generate better musical sequences that move, for example, from the core artists of 'Rock', to less central and cross-genre artists

('Rock'/'R&B'), to core artists of 'R&B', and so on, avoiding disturbing disruptions in the path. The same approach can work to generate playlists.

Another possible application of our analysis is to perform similarity assessments among artists. Having artists described as affinity vectors, we can obtain a measure of how close two artists are, independently from their 'genre' label. For instance, we can spot for each artist its nearest neighbour using Euclidean distance among these vectors. The results are interesting: in most cases we find that the nearest neighbour artists have the same genre label attached (e.g., Donna Summer goes to Diana Ross, both R&B), while sometimes they are differently labelled but still culturally associated, for example Nelly Furtado (Rock/Pop) goes to Missy Elliott (Rap), and Bob Sinclair (R&B) goes to Sonique (Rock/Pop). In this way, we exploit affinity fuzzy values to uncover associated artists, independently from their 'genre' label.

A third possible application is the creation of user profiles in music recommender systems. Rather than asking a user which musical genres she prefers, the system could ask her to name a few of her favourite artists, and then locate them on a multi-dimensional map of affinity degrees, to identify which styles of music the user seems to favour.

6 CONCLUSIONS

Music is a complex form of information that can be addressed both as an acoustic and a cultural phenomenon. Genre labels serve as simple and easily communicated classifications of musical style, but it can be shown that popular music often does not avail itself to such simple classifications. Musicians can share affinity with different genres, although in most organisational schemes (such as record stores) they have a unique genre label attached.

Our first contribution is providing a way to automatically uncover a richer relationship between artists and genres, based on social data in the form of playlists. Our proposal is moving from a Boolean concept of genres as exclusive categories to a view where artists have a degree of affinity with respect to each genre, and thus genres are conceptualised as fuzzy categories where affinity is represented as membership degrees in fuzzy set theory.

Our second contribution is developing a richer ontology for describing artist-to-genre and genre-to-genre associations, defining new concepts useful to better organise and understand large collections of music metadata. To enrich the characterisation of artists based on our first contribution (moving from Boolean categories to the notion of genre affinity), the ontology introduces the concepts of genre affinity degree, genre affinity vector, and genre centrality. To better characterise genres, the ontology introduces the concepts of core artists of a genre, bridge artists, and correlation or independence among genres.

To promote further social-based music analysis, we make publicly available the dataset of artists co-occurrences that we have produced from a very large repository of playlists. Since we identify each artist with its unique Musicbrainz ID, researchers have the chance to mash-up this dataset with a large number of other inter-linked music-related semantic Web datasets, following the methodology described in [7].

Future work includes expanding this approach to elucidate relationship between *songs*, *tags* and *genres*, viewing tags as user-provided fuzzy categories to determine the affinity of artists to tags, the core tags of genres and artists, the affinity and centrality of songs to genres, the relationship of songs with their artists, and the relationships among tags.

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FAST INDEX BASED FILTERS FOR MUSIC RETRIEVAL

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ABSTRACT

We consider two content-based music retrieval problems where the music is modeled as sets of points in the Euclidean plane, formed by the (on-set time, pitch) pairs. We introduce fast filtering methods based on indexing the underlying database. The filters run in a sublinear time in the length of the database, and they are lossless if a quadratic space may be used. By taking into account the application, the search space can be narrowed down, obtaining practically lossless filters using linear size index structures. For the checking phase, which dominates the overall running time, we exploit previously designed algorithms suitable for local checking. In our experiments on a music database, our best filter-based methods performed several orders of a magnitude faster than previous solutions.

1 INTRODUCTION

In this paper we are interested in content-based music retrieval (CBMR) of symbolically encoded music. Such setting enables searching for excerpts of music, or *query patterns*, that constitute only a subset of instruments appearing in the full orchestration of a musical work. Instances of the setting include the well-known query-by-humming application, but our framework can also be used for more complex applications where both the query pattern searched for and the music database to be searched may be polyphonic.

The design of a suitable CBMR algorithm is always a compromise between robustness and efficiency. Moreover, as robustness means high precision and recall, the similarity/distance measure used by the algorithm should not be too permissive to detect false matches (giving low precision) and not too restrictive to omit true matches (giving low recall). In this paper, we concentrate on a modeling of music that we believe is robust in this sense, and at the same time provides computationally feasible retrieval performance.

As symbolically encoded monophonic music can easily be represented as a linear string, in literature several solutions for monophonic CBMR problems are based on an appropriate method from the string matching framework (see e.g. [4, 6]). Polyphony, however, imposes a true challenge, especially when no voicing information is available or the occurrence is allowed to be distributed across the voices. In some cases it may suffice to use some heuristic, as for an example the SKYLINE algorithm [8], to achieve a monophonic reduction out of the polyphonic work. This, however, does not often provide musically meaningful results.

In order to be able to deal with polyphonic music, geometric-based modeling has been suggested [1, 7, 9, 10]. Most of these provide also another useful feature, i.e., extra intervening elements in the musical work, such as grace notes, that do not appear in the query pattern can be ignored in the matching process. The downside is that the geometric online algorithms [2, 5, 9, 10] are not computationally as efficient as their counterparts in the string matching framework. Moreover, the known offline (indexing) methods [1, 7] compromise on crucial matters.

These downsides are not surprising: the methods look at all the subsequences and the number of them is exponential in the length of the database. Thus, a total index would also require exponential space.

We deal with symbolically encoded, polyphonic music for which we use the pitch-against-time representation of note-on information, as suggested in [10] (see Figs. 1 and 2). The musical works in a database are concatenated in a single geometrically represented file, denoted by T. In a typical case the query pattern P to be searched for is often monophonic and much shorter than the database T to be searched. If P and T are readily not given in the lexicographic order, the sets can be sorted in $|P| \log |P|$ and $|T| \log |T|$ times, respectively. The problems of interest are the following:

(P1) Find translations of $P = p_1, p_2, \ldots, p_m$ such that each point in P match with a point in $T = t_1, t_2, \ldots, t_n$ ($p_i, t_j \in \mathbb{R}^2$ for $1 \leq i, j \leq m, n$).

(P2) Find translations of P that give a partial match of the points in P with the points in T.

Notice that the partial matches of interest in P2 need to be defined properly, e.g. one can use a threshold k to limit the minimum size of partial matches of interest.

Ukkonen et al. [9] presented algorithms PI and PII solving problems P1 and P2 in worst case times $O(mn)$ and $O(mn \log m)$, respectively. Their algorithms require $O(m)$ space. Noteworthy, the algorithm solving P1 has an $O(n)$ expected time complexity. Clifford et al. [2] showed that problem P2 is 3SUM-hard, i.e., it is unlikely that an exact solution could run faster than in quadratic time $O(mn)$, and give an approximation algorithm, called MSM, for P2, that runs in time $O(n \log n)$.

In this paper we introduce index-based filtering algorithms for the problems presented above. Our contribution

Figure 1. A musical excerpt.

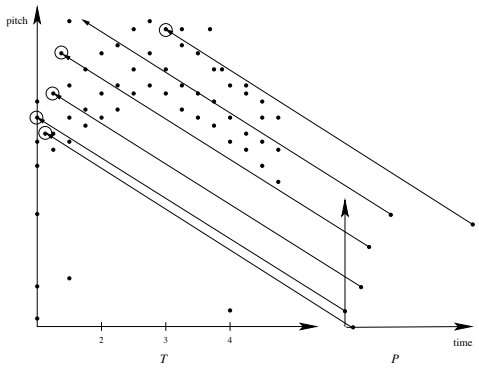

Figure 2. Pointset T represents Fig. 1 in the geometric representation. P corresponds to the first 2.5 bars of the melody line with a delayed 5th point. The depicted vectors form the translation f giving a partial match of P in T.

is twofold. Firstly, our methods outperform its competitors; in particular, the algorithms are *output sensitive*, i.e., the running time depends more on the output than on the input. This is achieved by exploiting a simple indexing structure that is not a total index. Secondly, we show how to keep the index of a practical, linear size. The index enables fast filtering; best results are obtained with filters running in $O(f(m)\log n + s)$ time. The found s ($s \leq n$) candidate positions are subsequently checked using Ukkonen et al's PI and PII algorithms. Thus, executing checking take time $O(sm)$ and $O(sm \log m)$, in the worst case, respectively.

2 INDEX BASED FILTERS

We will denote by $P + f$ a translation of P by vector f, i.e., vector f is added to each component of P separately: $P + f = p_1 + f, p_2 + f, \ldots, p_m + f$. Problem P1 can then be expressed as the search for a subset I of T such that $P + f = I$ for some f. Please note that a translation corresponds to two musically distinct phenomena: a vertical move corresponds to transposition while a horisontal move corresponds to aligning the pattern time-wise (see Fig. 2).

The idea used in [9, 10] is to work on trans-set vectors. Let $p \in P$ be a point in the query pattern. A translation vector f is a *trans-set vector*, if there is a point $t \in T$, such that $p + f = t$. Without loss of generality, let us assume all the points both in the pattern and database to be unique. So, the number of trans-set vectors is $O(n^2)$ in the worst case.

For the indexing purposes we consider translation vectors that appear within the pattern and the database. We call translation vector f *intra-pattern vector*, if there are two points p and p', $p, p' \in P$, such that $p + f = p'$. The *intra-database vector* is defined in the obvious way. The number of intra-pattern and intra-database vectors are $O(m^2)$ and $O(n^2)$, respectively. A nice property of Ukkonen et al's PI and PII algorithms is that they are capable of starting the matching process anywhere in the database. Should there be a total of s occurrences of the pattern within the database and an oracle telling where they are, we could check the occurrences in $O(sm)$ and $O(sm \log m)$ time, in the worst case, by executing locally PI and PII, respectively.

We will exploit this property by first running a filter whose output is subsequently checked by PI or PII. If a quadratic size for the index structure is allowed, we have a lossless filter: all intra-database vectors are stored in a balanced search tree in which translations can be retrieved in $O(\log n)$ time; Let $C(f)$ be the list of starting positions i of vector $f = t_j - t_i$, for some j, in the database, then the list is stored in the leaf of a binary search tree so that a search from root with key f leads this leaf. Let us denote by $|C(f)|$ the number of elements in the list. Since the points are readily in the lexicographic order, building such a structure takes a linear time in the number of elements to be stored.

However, for large databases, a quadratic space is infeasible. To avoid that, we store only a subset of the intra-database vectors. In CBMR, an occurrence is a compact subpart of the database typically not including too many intervening elements. Now we make a full use of this locality and that the points are readily sorted: for each point i in the database, $1 \leq i \leq n-1$, we store intra-database vectors to points $i+1, \ldots, i+c+1$ ($c = \min(c, n-i-1)$), where c is a constant, independent of n and m. Constant c sets the 'reach' for the vectors. Thus, the index becomes of linear, $O(n)$ size. Naturally such filters are no more totally lossless, but by choosing a large c and by careful thinking in the filtering algorithms, losses are truly minimal.

2.1 Solving P1

To solve the problem P1 we consider four straightforward filters. The first one works in $O(\log n + s)$ time, where s is the number of candidate position found, but we consider also two $O(m^2 \log n + s)$ time filters with a better filtration power. The simplicity of these filters are due to the fact that all the points need to find its counterpart within the database. Thus, to find candidate occurrences, we may consider oc-

currences of any of the intra-pattern vectors. The filters are represented in an increasing order in their complexities; the order also reflects their filtration power.

In FILTER0 we choose a random intra-pattern vector $f = p_j - p_i$. The candidate list $C(f)$ to be checked contains thus $s = |C(f)|$ candidates. For FILTER1 and FILTER2 we calculate frequencies of the distinct intra-database vectors. FILTER1 chooses the intra-pattern vector $f^* = p_j - p_i$ that occurs the least in the database, i.e., for which the $s = |C(f^*)|$ is smallest. In FILTER2, we consider the two intra-pattern vectors $f^* = p_j - p_i$ and $f^{**} = p_l - p_k$ that have the least occurrences within the database, i.e., for which $s' = |C(f^*)| + |C(f^{**})|$ is smallest. Then the set $S = \{i'' \mid t_{i'} - t_{i''} = p_i - p_1, i' \in C(f^*), t_{k'} - t_{i''} = p_k - p_1, k' \in C(f^{**})\}$ contains the candidates for starting positions of the pattern, such that both f^* and f^{**} are included in each such occurrence.

For the running time, FILTER0 uses $O(\log n)$ time to locate the candidate list. FILTER1 and FILTER2 execute at most $O(m^2)$ additional inquiries each taking $O(\log n)$ time. FILTER2 needs also $O(s')$ time for intersecting the two occurrence lists into the candidate list S; notice that values i'' can be scanned from left to right simultaneously to the scanning of lists $C(f^*)$ and $C(f^{**})$ from left to right, taking amortized constant time at each step of the intersection.

With all the filters we may consider only translations between consecutive points of the pattern. Thus, we would somewhat compromise on the potential filtration power, but the 'reach constant' c above would get an intuitive interpretation: it tells how many intervening points are allowed to be in the database between any two points that match with consecutive pattern points. For long patterns, the search for the intra-pattern vector that occurs the least in T may dominate the running time. Hence, we have FILTER3 that is FILTER2 with a random set of intra-pattern vectors as the input.

2.2 Solving P2

The same preprocessing as above is used for solving P2, but the search phase is modified in order to find partial matches. We will concentrate on the case where a threshold k is set for the minimum size of a partial match. Since any pattern point can be outside the partial match of interest, one should in principle check the existence of all the $O(m^2)$ vectors among the intra-database vectors, merge the candidate lists into multiset S' and accept any candidate position i'' into set S that occurs at least k times in S', and run the checking on each candidate position in S with algorithm PII. More formally, the multiset S' contains position i'' for each intra-pattern vector $f = p_j - p_i$ such that $i' \in C(f)$ and $p_i - p_1 = t_{i'} - t_{i''}$. We call this basic lossless filter FILTER4. We will also consider a lossy variant FILTER5, where for each pattern point p only one half of the intra-pattern vectors (the least frequent ones) having p as an endpoint is chosen.

The *pigeon hole principle* can be used to reduce the amount of intra-pattern vectors to check: If the pattern is split into $(m-k+1)$ distinct subsets, then at least one subset must be present in any partial occurrence of the complete pattern. Therefore, it is enough to run the filters derived for P1 on each subset independently and then check the complete set of candidates. The total amount of intra-subset vectors is bound by $O((m-k+1)(\frac{m}{m-k+1})^2) = O(\frac{m^2}{m-k+1})$. This is $O(m)$ whenever k is chosen as a fraction of m. FILTER0 and FILTER2 both select constant number of vectors among each subset, so the total number of candidate lists produced by each filter is $O(m-k+1)$. Hence, this way the filtration efficiency (number of candidates produced) can be expected to depend linearly on the number of errors $m-k$ allowed in the pattern. This is an improvement to the trivial approach of checking all $O(m^2)$ intra-pattern vectors.

Notice that these pigeon hole filters are lossless if all the intra-database vectors are stored. However, the approach works only for relatively small error-levels, as each subset needs to contain at least two points in order the filtering to be possible. Let us focus on how to optimally use FILTER1 for the subsets in the partition, as FILTER0 and FILTER2 are less relevant for this case. The splitting approach gives the freedom to partition the pattern into subsets in an arbitrary way. For optimal filtering efficiency, one should partition the pattern so that the sum of least frequent intra-subset vectors is minimized. This sub-problem can be solved by finding the *minimum weight maximum matching* in the graph whose nodes are the points of P, edges are the intra-pattern vectors, and edge weights are the frequencies of the intra-pattern vectors in the database. In addition, a set of dummy nodes are added each having an edge of weight zero to all pattern points. These edges are present in any minimum weight matching, so their amount can be chosen so that the rest of the matched edges define the $m-k$ non-intersecting intra-pattern vectors whose frequency sum is minimum.

We use an $O(m^3)$ time minimum weight maximum matching algorithm to select the $m-k+1$ intra-pattern vectors in our filter. Some experiments were also done with an $O(m^2)$ time greedy selection. We call this algorithm FILTER6 in the experiments.

3 EXPERIMENTS

We set the new algorithms against the original Ukkonen et. al.'s PI and PII [9]. We set the window length within the database (the reach constant c) to 50, the window length within the pattern in FILTERs 0–3 and 6 to 5, and in FILTERs 4–5 to 12. In FILTER 6, we experimented with different values of k in range $\lceil m/2 \rceil$ to $\lceil \frac{15}{16} m \rceil$. Overall, these settings are a compromise between search time, index memory usage and search accuracy for difficult queries. Larger window lengths may be required for good accuracy depending on the level of polyphony and the type of expected queries.

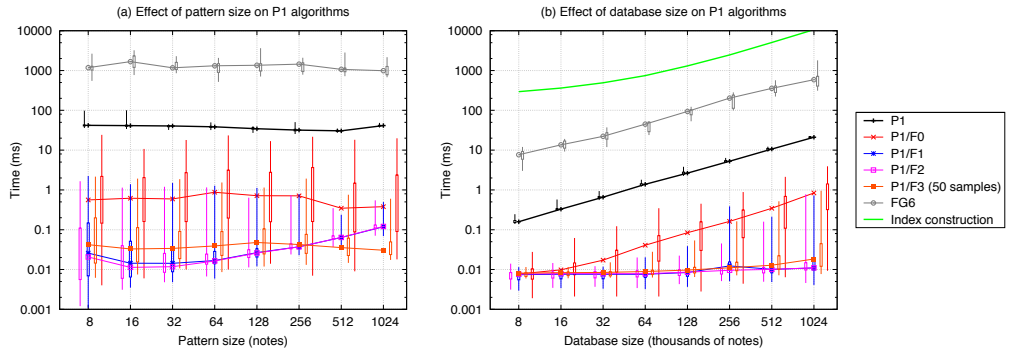

Figure 3. Solving P1. Search time as function of pattern and database sizes in the MUTOPIA collection of 1.9 million notes. Database size experiments were done with a pattern size of 64 notes. Note the log scales.

We also compare with Clifford et. al.'s MSM [2] and Fredriksson and Grabowski's FG6 algorithm [3, Alg. 6.]. The latter solves a slightly different problem; the time information is ignored, and the pitch values can differ by $\delta \geq 0$ after transposition. We set $\delta = 0$ to make the setting as close to ours as possible. We tested also other algorithms in [3] but Alg. 6 was constantly fastest among them on our setting.

To measure running times we experimented both on the length of the pattern and the database. Reported times are median values of more than 20 trials using randomly picked patterns. As substantial variations in the running times are characteristic to the filtering methods, we have depicted this phenomenon by using box-whiskers: The whisker below a box represents the first quartile, while the second begins at the lower edge and ends at the median; the borders of third and fourth quartiles are given by the median, the upper edge of the box and the upper whisker, respectively.

We also measured how robust the PII-based methods (FILTERs 4-6) are against noise, and calculated a precision-against-recall plot. Here we reported the mean value of a 100-trial experiment in which the set of occurrences found by PII was considered as the ground truth. All the experiments were carried out with the MUTOPIA database; at the time it consisted of 1.9 million notes in 2270 MIDI files.

3.1 Experimenting on P1

In experimenting on the pattern size, the variations in the running times of the PI-based filters are clearly depicted in Fig. 3a. Out of the four filters, FILTER2 performed most stably while FILTER0 had the greatest variation. The figure shows also that all our filtering methods constantly outperform both the original PI and the FG6 algorithm. With pattern sizes $|P| \lesssim 400$, FILTERs 1 and 2 are the fastest ones, but after that FILTER3 starts to dominate the results.

The evident variation in our filters is caused by difficult patterns that only contain relatively frequently appearing vectors. In our experiments, FILTERs 1–3 had search times of at most 2 ms. It is possible to generate patterns that have much longer search times, especially if the database is very monotonic or short windows are used. However, in practice these filters are at least 10 times faster than PI and 200 times faster than FG6 for every pattern of less than 1000 notes.

When experimenting on the size of the database as shown in Fig. 3b, execution times of online algorithms PI and FG6 increases linearly in the database size. Also FILTER0's search time increases at the same rate due to the poor filtering efficiency. FILTERS 1–3 have much lower slope because only few potential matches need to be verified after filtering. Fig. 3b also depicts the construction time of the index structure for the filters. Remember that this costly operation needs to be executed only once for any database.

3.2 Experimenting on P2

Fig. 4 shows the experimentally measured search times for P2 filters, PII and MSM. When varying the size of the pattern, our PII-based filters clearly outperformed MSM in all practical cases (Fig. 4a): MSM becomes faster than PII only when $|P| > 400$ and faster than FILTER4 when $|P| > 1000$. In this experiment FILTER6 dominates over FILTER5 until $|P| \gtrsim 250$ after which FILTER5 starts to dominate. However, a greedy version of FILTER6 outperforms them both by an order of magnitude.

Results of the experiments on the length of the database are rather similar (Fig. 4b), exceptions being that MSM is constantly outperformed by the others and that FILTER6 performs the best throughout the experiment. Again, the greedy FILTER6 is the fastest: it is nearly 100 times faster than FILTER6 and over million times faster than MSM.

3.2.1 Comparing whole musical works.

In [2], Clifford et al. carried out experiments for partial music matching and concluded that their algorithm is faster than PII when two whole documents are to be matched

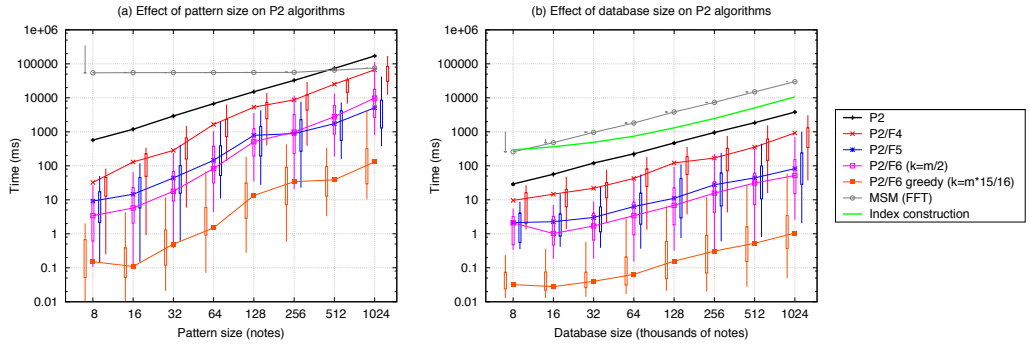

Figure 4. Solving P2. Search times as function of pattern and database sizes. Database and pattern sizes as in Fig 3.

against each other. Fig. 5 depicts results of our experiment using their setting but including also FILTERs 4–6. In our experiment, MSM becomes faster than PII when $|P| = |T| \gtrsim 600$ and dominates FILTERs 4 and 5 when the size of the matched documents exceeds 1000 and 5000 notes, respectively. Depending on the value of k, MSM becomes faster than FILTER6 at document sizes larger than 1500–20,000 notes. However, in this specific task algorithms would be expected to return matches that are relatively poor if measured as a ratio between the matched notes and pattern size. Solving the task by using FILTER6 with $k = m/16$ would not give good results, but with $k = m/2$ FILTERs 4–6 are comparable with MSM.

3.2.2 *Precision and recall.*

For experimenting on the robustness against noise we selected a random pattern of length 100 from the database and introduced mutations (substitutions) to it; parameter *error rate* gives their number. Then we retrieved all approximate occurrences of the pattern from the database by using PII to form the ground truth. To keep the ground truth at a manageable size, we used the parameter *ground truth similarity*, gts, as a similarity cutoff point for the retrieved matches. For example, ground truth similarity value 0.1 defines that each item in the ground truth must match with at least one tenth of the pattern notes. In the end, we measured the precision of the algorithms as the function of recall with different settings of ground truth similarity and error rate. This was repeated 100 times with randomly selected patterns and the resulting graphs were averaged into one result.

Basic evaluation of the measurements is doable by comparing areas: the larger the area below a curve the better the accuracy and robustness of the corresponding algorithm. Both increasing error rate and lower ground truth similarity values make the task harder: higher error rate makes the original patterns, with possibly several occurrences, be less pronounced in the retrieved result lists. When good matches have been toned down like this, algorithms need to find more difficult matches to score well. Also correct ordering of the

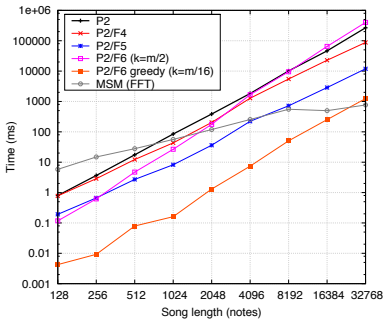

Figure 5. Matching whole music works against each other.

results by decreasing similarity with the pattern is necessary.

We bound ground truth similarity to 0.10 and let the error rate vary between 0, 0.5 and 0.70; this setting requires the algorithms to find potentially very sparse matches. Fig. 6 depicts that, at low error rates, there is not much difference between the index filters, and MSM performs similarly. However, at error rate 0.75, only FILTER4 achieves good scores. The results suggest that all the filters are well balanced when comparing their robustness to execution speed: FILTERs 5 and 6 are fast but somewhat inaccurate, while FILTER 4 has its place between them and PII.

4 CONCLUSIONS

We considered point pattern matching problems applicable to CBMR and showed how they can be solved using index-based filters. Given a point pattern P of m points, the problems are to find complete and partial matches of P within a database T of n points. The presented filters are lossless if $O(n^2)$ space is available for indexing. We also introduced a more practical, linear size structure and sketched how the filters based on this structure become virtually lossless.

After the preprocessing of the database, the proposed filters use $O(f(m) \log n + s)$ time to produce s candidate positions that are consequently checked for real occurrences with the existing checking algorithms. The filters vary on

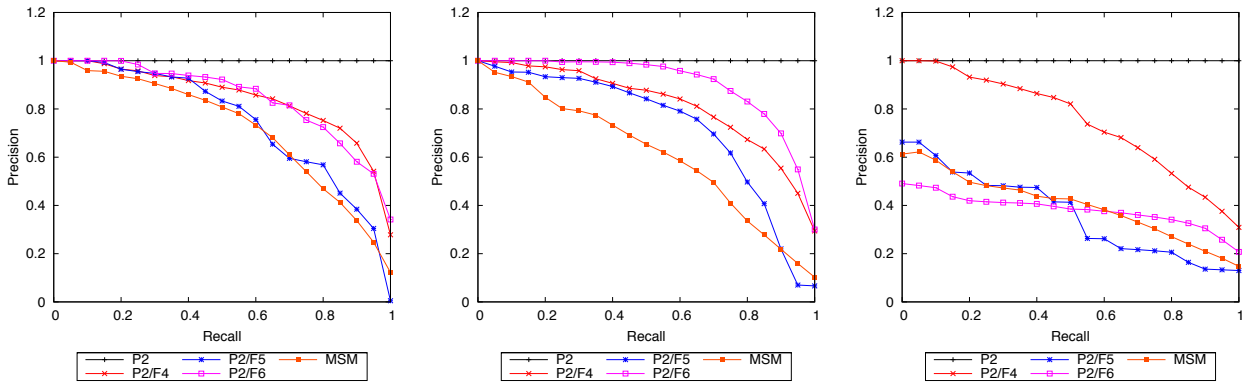

Figure 6. Precision-Recall at error rates 0, 0.5 and 0.75 (left to right), with $gts = 0.1$.

the complexity of $f(m)$ and on the output size s. Since the filtering power of the proposed filters is hard to characterize theoretically, we ran several experiments to study the practical performance on typical inputs.

The experiments showed that our filters perform much faster than the existing algorithms on typical application scenarios. Only when comparing large musical works in their entirety, MSM [2] outperforms our new filters.

Since the guarantee of losslessness in the filters is only valid on limited search settings (number of mismatches allowed, constants limiting maximal reach, etc.), it was important to study also the precision and recall. This comparison was especially fruitful against MSM, that is an approximation algorithm, and can hence be considered as a lossy filter as well. The experiments showed that our filters typically performs at the same level that MSM in this respect.

As a future work, we plan to study the extensions of the filters to approximate point pattern matching; in addition to allowing partial matching, we could allow matching a point to some ϵ-distance from its target. Such setting gives a more robust way to model the query-by-humming application. Although it is straightforward to extend the filters to consider the candidate lists of all intra-database vectors within the given ϵ-threshold from any intra-pattern vector, the overall amount of candidate positions to check grows fast as the threshold is loosen. Therefore, finding better strategies for filtering in this scenario is an important future challenge.

5 ACKNOWLEDGEMENTS

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DIRECT AND INVERSE INFERENCE IN MUSIC DATABASES: HOW TO MAKE A SONG FUNK?

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ABSTRACT

We propose an algorithm for exploiting statistical properties of large-scale metadata databases about music titles to answer musicological queries. We introduce two inference schemes called "direct" and "inverse" inference, based on an efficient implementation of a kernel regression approach. We describe an evaluation experiment conducted on a large-scale database of fine-grained musical metadata. We use this database to train the direct inference algorithm, test it, and also to identify the optimal parameters of the algorithm. The inverse inference algorithm is based on the direct inference algorithm. We illustrate it with some examples.

1. INTRODUCTION

Large databases of metadata are now available in many domains of digital content such as music (allmusic) or films (imdb). However, the scaling up of content databases makes it increasingly difficult and expensive to maintain manually these metadata databases. The apparition of collaborative tagging addresses this issue by exploiting the power of large distributed networks of users. However, the basic issue of maintaining these repositories of information and coping with their possible incompleteness remains open.

The aim of our study is to design algorithms to turn large-scale databases of musical information into knowledge bases. More precisely, we wish to exploit these databases to perform statistical inferences from partial descriptions of items in order to infer new information about the content.

The inferences we target must be both statistically coherent with the data and computationally tractable. The inferences we consider in this study are of two sorts: direct and inverse. Direct inference consists in inferring the most plausible value of an attribute given a set of "observed attributes". These inferences can be used typically to lower the maintenance cost of the database: when new items are encountered, only a subset of attributes would then be needed to be observed, and the other ones could be inferred with some degree of confidence. Inverse inference answers a different question: given a set of observed attributes (possibly complete, i.e. a fully described title), and a target attribute value, we wish to know the minimal modifications to apply to this initial observation set to reach this target attribute value. An application of direct inference is algorithms for automatic music categorization. Some attributes like 'Style Rap' or 'Distorted Guitar' can be relatively well estimated from the analysis of the acoustic signal [2]. However, acoustic classifiers fail to compute more "high-level" attributes like "Mood sad" or "Lyrics cynical" with a satisfactory performance. The idea of such a hybrid approach is to first compute the best acoustic attributes and, in a second stage, infer the remaining attributes using direct inference.

As an example of inverse inference, we can consider a computer-aided composition tool that suggests musicians how to modify their compositions: starting from an initial composition, the system would observe some "low-level" attributes. The musician could then ask the system what attributes to change, and how, in order to, say, make his title sound "Funk", or not "aggressive", or "harmonious". Inverse inference yields the minimal modifications of the initial title to increase optimally the corresponding probabilities.

In this paper, we propose an algorithm for direct and inverse inference, based on an efficient implementation of a kernel regression approach. We describe an evaluation experiment conducted on a large-scale database of fine-grained musical metadata. We use this database to train the direct inference algorithm, test it, and also to identify the optimal parameters of the algorithm. Inverse inference is based on the direct inference algorithm. We illustrate it with real world examples.

1.1. State-of-the-art

To perform our inferences, we would need ideally a statistical model of our attributes in the form of a Bayesian Network, yielding a compact representation of the underlying distribution of the attributes. If we had such a model inference would be carried out with approximate methods suitable for large networks.
We tried to learn such a model of our random variables with the SBNS algorithm proposed in [3]: since our data is sparse, we can use *frequent sets* (i.e. sets of attributes co-

occurring more than some threshold. For more details, see [1]) to build local models. Using frequent sets dramatically decreases the complexity of the structural learning of a Bayesian network model, as remarked in [5].

To carry out inferences in the large network we obtained, we used *Mean Field Theory* (MFT). MFT is a statistical physics approximation to solve the many-body problem. It consists in replacing all interactions to any one body with an average interaction, turning the N-body problem into $N(N-1)/2$ 2-body problems. When applied to Bayesian Networks for inference, MFT means estimating the real distribution of the inferred nodes given the observed nodes with a simpler distribution; this simpler distribution assumes independence between inferred nodes given observed nodes. An efficient implementation of MFT can be found in [8] and complete details about variational inference methods in [6].

Unfortunately, the network we obtained was too densely connected even for MFT to perform in reasonable time. As an example, the computation of the MFT equation for one title, given an observable set, would take approximately 5 minutes, so cross-validation on our database (described in the next section) would take about 5 months, which is not acceptable. In this context, inverse inference would be even less tractable. However, our database is probably not as sparse as those used in [3]. This paper is an attempt to address this challenge.

1.2. Description of the data

For this study we have used a music dataset comprising 37,000 popular songs of various styles. Each song is described by 948 binary ($0/1$ valued) attributes. Attributes describe low-level information (like the presence of "acoustic guitar", or the "tempo range" of the song) as well as high-level characteristics (like the "style" of a song, the mood emerging from it, etc.). The attributes are grouped in 17 categories: *Style*, *Genre*, *Musical setup*, *Main instruments*, *Variant*, *Dynamics*, *Tempo*, *Special*, *Era/Epoch*, *Metric*, *Country*, *Situation*, *Mood*, *Character*, *Language*, *Popularity*, and *Rhythm*. The complexity of the database makes it impossible to describe it in details here, but we are only concerned in this study by the number of attributes and the quality of the inferences obtained.

The database we considered for this study is *sparse*: The mean number of attributes set to true per song (occupation factor) is 4% (i.e. 40 on a total of 948). Sparseness suggests the dominant role of the true-valued attributes versus false-valued attributes for a given song. Therefore, in the analysis of the database, our inference algorithm should treat differently true values and false values. Another feature of the database is its *redundancy*. For instance, attributes like 'Country Greece' and 'Language Greek' are extremely correlated.

This justifies the presence of inter-attribute dependencies, that our inference algorithm will attempt to exploit.

1.3. Problem Statement

In the context of our Boolean multi-attribute database, we can formulate our problem as follows. Let D denote the number of attributes ($D = 948$), and $(A_i)_{1\le i\le D}$ the set of attributes modeled as $0/1$ valued random variables. We consider each song description as a realization of the random vector A and thus denote A_i^s the value of the i^{th} attribute ($1 \le i \le D$) for the s^{th} song ($1 \le s \le N$) where N is the number of songs in the database ($N = 37{,}000$). Let $\Pr(\cdot)$ denote the underlying probability distribution of the random vector A. We wish to accomplish two tasks:

Direct inference: Given an incomplete description of a music title, determine the most probable values of the remaining uninformed attributes. Formally, if for a subset $I \subset \{1, \dots, D\}$ the observed values of the attributes $\left(A_{I_1}, \dots, A_{|I|}\right)$ (denoted A_I) are $\left(a_{I_1}, \dots, a_{I_{|I|}}\right)$ (denoted a_I), we wish to estimate the most probable values of the non-observed attributes, i.e. compute:

$$e^* \in \operatorname{argmax}_{e\in\{0,1\}^{D-|I|}} \Pr(A_I = e \mid A_I = a_I)$$

Inverse inference: Given an incomplete description of a music title *and* target values for a subset of target attributes, how can one modify the initial description to achieve the target values with maximum confidence, with a minimal modification? Formally, for a subset $I \subset \{1, \dots, D\}$ and $J \subset \{1, \dots, D\}$ such that $I \cap J$ is empty, the observed values of A_I are a_I and the target fields A_T are meant to be a_T, but are not, so we suppose that $\Pr(A_T = a_T \mid A_I = a_I) < 0.5$. Let then δ_A be an indicator function with value 1 if A is true, and 0 otherwise. We say that σ is a *flip* of the observed attributes A_I if:

$$\sigma: \{0,1\}^{|I|} \longrightarrow \{0,1\}^{|I|}$$
$$(u_i)_{1\le i\le |I|} \mapsto \left(\delta_{i\notin D(\sigma)} u_i + \delta_{i\in D(\sigma)}(1-u_i)\right)_{1\le i\le |I|}$$

The subset $D(\sigma) \in I$ characterizes the flip σ: this definition means that the flip will inverse its parameters which index belong to $D(\sigma)$ and leave the remaining parameters unchanged. The *order* of the flip is $d(\sigma) = |D(\sigma)|$. It represents the number of modifications on the parameter of the flip. Let S denote the space of all flips (its cardinality is $2^{|I|}$). The inverse inference problem consists in solving the following combinatorial optimization problem:

$$\sigma^* \in \operatorname{argmin}_{\sigma\in S}[d(\sigma) - \lambda \Pr(A_T = a_T \mid A_I = a_I)]$$

Here $\lambda > 0$ is a free parameter representing a trade-off between minimal flip order and maximum probability of achieving the target state. Since I could be of cardinality up to 947, the search space S is huge so an important feature of our algorithm is the evaluation speed of this functional form.

2. DIRECT INFERENCE

In this section we describe our direct inference algorithm and evaluate its performance.

2.1. Description of the algorithm

Computing $\Pr(A_{\bar{I}} = e | A_I = a_I)$ for all $e \in \{0,1\}^{D-|I|}$ is combinatorial and computationally intractable for the dimensions we are dealing with ($D - |I| \approx 700$), so the basic assumption to break the intractability is that the variables $A_{\bar{I}_1}, \dots, A_{\bar{I}_{|\bar{I}|}}$ are independent given A_I. It then suffices to compute separately $\Pr(A_{\bar{I}_k} = 1 | A_I = a_I)$ for all $k \in \{1, \dots, |\bar{I}|\}$ to determine the most probable values of the inferred attributes $A_{\bar{I}}$. Although this assumption might seem contradictory with our primary aim to model dependencies, the intuition is that the inferred $A_{\bar{I}}$ attributes only depend on observed attributes A_I, meaning that subjective attributes emerge solely from objective (possibly acoustic) attributes that will form our observation I. This independence assumption is less strong as the size of the observation set grows. However, the choice of the attributes A_I will have to be dealt with carefully, and we discuss this issue later on.

To compute $\Pr(A_{\bar{I}_k} = 1 | A_I = a_I)$ we use a kernel regression estimator [4] and [7]:

$$\hat{p}_k(a_I) = \frac{\sum_{s=1}^{N} \omega_\theta\big(d(a_I, A_I^s)\big) A_{\bar{I}_k}^s}{\sum_{s=1}^{N} \omega_\theta\big(d(a_I, A_I^s)\big)}$$

where $d(a_I, A_I^s)$ is a distance function indicating the 'proximity' between our observed attributes a_I and the corresponding values of attributes A_I for the s-th song in the database. For instance, $d(\cdot,\cdot)$ could be the Euclidian distance in $\mathbb{R}^{|I|}$ counting the number of differences between two binary strings passed as arguments. However, the Euclidian distance treats equivalently both values 0 and 1. Yet the sparseness of the data indicates clearly that two binary strings are more similar if they have common 1s than if they have common 0s. To emphasize the important role of common ones, we rather use the Jaccard distance:

$$d(u,v) = d\big((u_i)_{1\le i\le p}, (v_i)_{1\le i\le p}\big) = \frac{\sum_{i=1}^{p} \delta_{u_i \neq v_i}}{\sum_{i=1}^{p} \delta_{u_i \neq v_i} + \sum_{i=1}^{p} \delta_{u_i=1}\delta_{v_i=1}}$$

The numerator is the number of differences between the two arguments, and the denominator is composed of two terms: the first one is again the number of differences and the second term is the number of "1" the strings have in common. Notice that for all $u, v \in \mathbb{R}^p$ $0 \le d(u,v) \le 1$.

$\omega_\theta(\cdot)$ is a weighting function with a real parameter $\theta \in \mathbb{R}^+$. An obvious property of this function is that it decreases towards 0 as its parameter increases. Intuitively, this means that the closer a song is from our observation a_I (using distance d) the greater its weight should be. The integral of the weighting function $\omega_\theta(\cdot)$ should also be finite. Notice that the weighting function needs not be normalized since the estimator $\hat{p}_k$ is already normalized. A simple choice for ω_θ is a Gaussian filter $\omega_\theta(x) = \exp\left(-\frac{x^2}{\theta^2}\right)$ but since θ is a free parameter that we have to tune, a nice property would be to have an intuitive understanding of the value θ. For example, with a simple linear decreasing function $\omega_\theta(x) = \left(1 - \frac{x}{\theta}\right)_+$, the value of θ is just a threshold distance above which songs have a weight set to 0, and are thus not taken into account in the estimation $\hat{p}_k$. Note that if $\theta \to \infty$ our estimator is an empirical mean on all the songs of the database, each song having identical importance, no matter the distance. If $\theta \to 0^+$ then $\omega_\theta(x) = \begin{cases} 1 \text{ if } x = 0 \\ 0 \text{ if } x \neq 0 \end{cases}$ and we are only estimating the empirical distribution on the data.

2.2. Evaluation & Experiments

This section introduces experiments to assess the performance of our algorithm. We also study the influence of several parameters of the algorithm.

2.2.1. Performance criterion

Our database contains about 4% of attributes set to 1. This unbalancedness forbids the use of precision or recall to assess our classifiers (a dumb classifier that always predicts an attribute $A_{\bar{I}_k}$ to 0 would have an accuracy of 96%). Instead, we use the more suited F-measure. For a given attribute, *F-measure* (for the true class) is defined as the mean of precision and recall, $F = \frac{2\pi\rho}{\pi + \rho}$. A classifier always responding false has a F-measure of 0.

2.2.2. Observed Attributes

In this paragraph we discuss the different choices for the observation set I. The choice can either be driven by applications, or by performance: if the goal is to complete an acoustic automatic categorization system, then there is a canonical order for all the attributes defined by the decreasing performance of some acoustic classifier. We call this order the *acoustic order*. Another possible choice for I is the set of *modifiable attributes*. This set of observations will be used in the inverse inference problem and has been determined by hand: modifiable attributes are low-level attributes that a musician can easily modify (like instruments, tempo, etc.). On the other hand, if what we are looking for is to increase the performance of our inference algorithm, the choice for I is not obvious. The simplest way to formulate the problem is to choose I as one of the subsets of $\{1, \dots, D\}$ of fixed cardinality m maximizing some global performance criterion $\mathcal{L}$ for inference:

$$I_m^* \in \underset{\substack{I \subset \{1,\dots,D\} \\ |I| = m}}{\operatorname{argmax}} \mathcal{L}\left(\left[\hat{p}_k(a_I) \right]_{\substack{1 \le k \le |\bar{I}| \\ a_I \in \{0,1\}^m}} \right)$$

The arguments of $\mathcal{L}$ are all possible responses to an inference query. The global performance criterion should 'integrate' in a certain sense all possible inference responses to all $a_I \in \{0,1\}^m$. Of course, this optimization problem is completely intractable and we shall restrict our empirical study to the case of the *acoustic* and *modifiable* attributes.

2.2.3. Experiments

In this paragraph, we assess the influence of the free parameters of the algorithm, namely θ and ω_θ and the influence of the choice of the observation I. The performances were computed using leave-one-out cross-validation on the whole dataset.

2.2.3.1 Influence of the smoothing parameter

First, we evaluate the influence of the free parameter $\theta > 0$ on the overall performance. In this experiment, we thus fix the observation I to be the 100 first attributes in the acoustic order.

On Figure 1, we show the performance of our algorithm with θ varying in the interval $[0,1[$. The reason we restrict the search to this interval is because intuitively we do not want to take into account all the songs in the database: songs with large distance (close to 1) should be ignored and, as we mentioned earlier, θ can be interpreted as a threshold distance. We plotted the F-measure for several individual attributes versus the smoothing parameter θ on the range $[0,1[$.

All of the attributes exhibit a bell-shaped F-measure vs. θ curve. This shows that there exists a maximum of the performance for some optimal value of the smoothing parameter. The figure also shows that this optimal value is different from one attribute to another. If the smoothing parameter was unique for all attributes and chosen to maximize the performance of, say, attribute 'Country France', the performance of the attribute 'Variant Wah-wah' would be zero whereas it could be near 80% with an individual fine smoothing tuning. This suggests that we should use one smoothing parameter per attribute to achieve maximum performance as opposed to the classic approach of maximizing a global performance criterion over all attributes. On figure 1 we plotted the cumulative distribution function (cdf) for the set of F-measures (over all inferred attributes) obtained for several choices of the smoothing parameter. A perfect classifier would have a cdf starting at the origin straight to point $(1,0)$ and from then straight to point $(1,1)$.

The worst classifier would have a cdf curve starting at the origin, straight to point $(0,1)$ and then straight to point $(1,1)$. Visually, the best classifier among our choices for the smoothing parameter is thus the closest curve to that of the perfect classifier. Figure 2 confirms that choosing individually optimized smoothing parameters for each attribute largely outperforms a global optimized parameter.

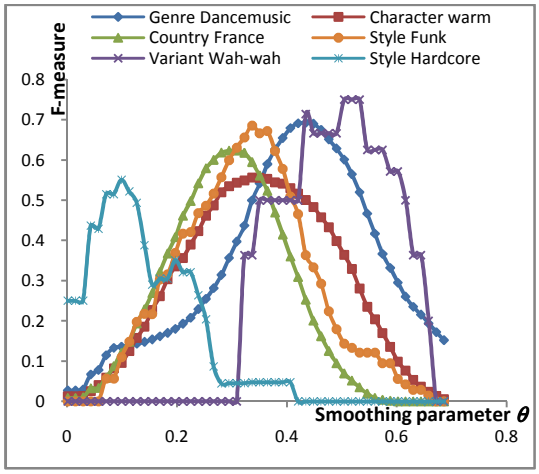

Figure 1: Influence of the smoothing parameter on the performance of several classifiers.

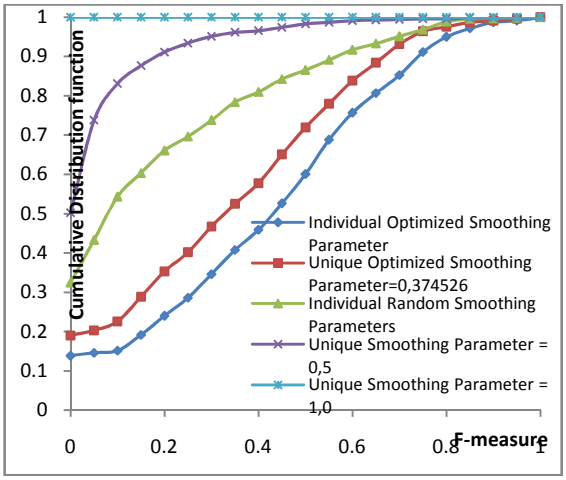

Figure 2: CDF of the F-measures over all inferred attributes for several choices of smoothing parameters. We select a unique parameter (for all attributes) that maximizes the sum of F-measures.

2.2.3.2 Influence of the weighting function

A well-known fact of kernel regression is that the choice of the smoothing parameter θ is more important than the choice of the weighting function. To confirm this in our case, we computed the performances for all inferred attributes for several functional forms for ω_θ and where θ was optimally chosen for each attribute as discussed above. The observation I is still the first 100 attributes in the acoustic order. On Figure 3, we plotted the cdf for each set of F-measures obtained for several choices of the weighting function.

Figure 3 shows that although there is a slight performance increase for the Gaussian function over other functional forms (with individually optimized smoothing parameters),

the improvement is negligible compared to the performance of a classifier with Gaussian weighting function and random choice for the smoothing parameters. Clearly, the choice of the smoothing parameters is critical and no particular effort should be dedicated to the choice of the weighting function. We thus choose to use the linear weighting function, as discussed in the algorithm description, since exponentials are expensive to compute. Linear weighting functions are a rather good tradeoff between performance and time.

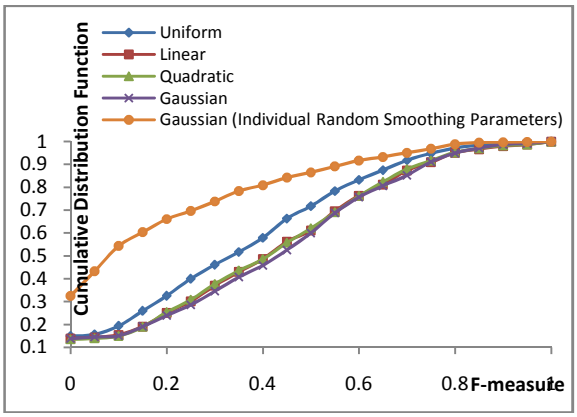

Figure 3: CDF of F-measures for several functional forms of the weighting function and for individual optimized smoothing parameters. We also plotted the cdf of the F-measures for random choice of the smoothing parameters.

2.2.3.3 *Choice of the distance*

To assess the choice of our distance, we compare the performance of our algorithm using the Jaccard distance and the Euclidian distance. Although computing an Euclidian distance is slightly faster than computing a Jaccard distance, the gain in performance is worth it: on Figure 4 we plotted the cdf for the Hamming and Jaccard algorithms with individual optimal smoothing parameters and a Gaussian weighting function.

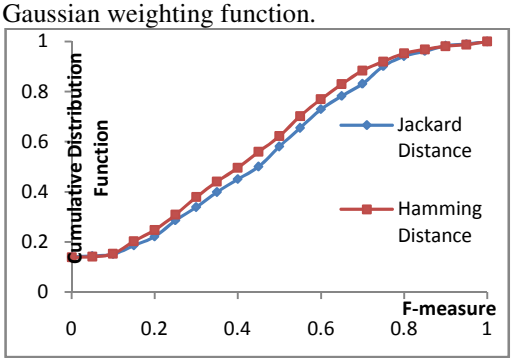

Figure 4: CDF of the F-measures of inferred attributes for the Hamming or Euclidian distance and the Jaccard-based distance.

2.2.3.4 *Choice of the observation*

The last parameter that may impact the quality of the inference is the observation set I. For the acoustic order, we would like to confirm the intuition that as $|I|$ increases, the overall performance converges towards an upper limit. On Figure 5 we plotted the cdf of the F-measures over all inferred attributes for several values of $|I|$. It can be seen that although there is an improvement of the performance for small values of $|I|$, there is a 'glass ceiling' for the classifier's performance.

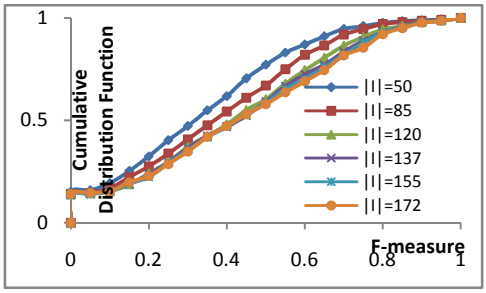

Figure 5: CDF of the F-measures of all inferred attributes for several values of the size of the observation I.

3. INVERSE INFERENCE

3.1. Description

As mentioned earlier, the observation is fixed when performing inverse inference so I is the set of modifiable attributes, meaning all low-level attributes and of which we can easily think of as attributes that a musician can easily change in a song: instruments, tempo, language, etc. Some other attributes are clearly not that easily modifiable like the "mood" of the song, or its "style". These more subjective attributes are supposed to "emerge" from the observation. The modifiable attributes set is of cardinality 232. So there are 2^{232} possible flips of the observation. To reduce this search space, we introduce the following heuristic: recall that there are categories in which all attributes can be grouped. For instance, all sorts of moods (calm, aggressive, dreamy, etc.) can be grouped in one category. In each category we define a limit to the number of true attributes. This limit is determined using the statistical distribution of the number of true attributes per category on the songs of the database. The limit we fix is the 90%-quantile of this distribution for each category. If the initial state a_I does not respect these limits once flipped, the flip will not be evaluated. We call a_I-*authorized flips* such flips and S_{a_I} the subset of authorized flips. The problem we will solve as an approximation of the general inverse inference problem we mentioned earlier is the following:

$$\sigma^* \in \operatorname{argmin}_{\sigma \in S_{a_I}} \left[d(\sigma) - \lambda \prod_{k \in T} \hat{p}_k(a_I)^{a_k} \left(1 - \hat{p}_k(a_I)\right)^{1-a_k} \right]$$

The product is just the estimated value of $\Pr(A_T = a_T | A_I = a_I)$.

```
Do
  flip = get next authorized flip
  If ( fitness(flip) is one of the M best
fitness's of order 1 )
  Then add flip to M best flip list
Until no more authorized flips of order 1
For k=2..max order Do
    flip = get next authorized flip
    If flip contains same attributes as at least
an element of M best flip list
    And fitness(flip) is one of the M best
    Then add flip to temporary M best flip list
  Until no more authorized flips of order k
  M best flip list = temporary M best flip list
End For
```

Table 1. The inverse inference algorithm.

We finally explore the search space with the following strategy: we scan through the a_I-authorized order 1 flips and evaluate their fitness; we select the M best authorized order 1 flips ($M$is a parameter of the approximation method). We then loop on the order of the flip, and only consider a_I-authorized flips that contain the same attributes as at least one of the M best flips of previous order, and update the M best flip list (Table 1).

3.2. Examples

We illustrate here inverse inference with an example. We tried to make "Black or White" by Michael Jackson sound 'Funk' (its style was initially tagged as 'Pop' and 'Rock/Pop'). We set tradeoff parameter $\lambda = 1000$ so that in the optimization process, an additional modification is only accepted if it increases the probability of being 'Funk' by at least 0.01. This 'Probability of Target' is the probability of being 'Funk' given the 232 modifiable attributes of the song "Black or White" (or its flipped version). Initially, this probability is about 0.05 and even though we are not sure the flip we found is optimal, it yields a probability of being 'Funk' of about 0.87 which is much better. The set of attributes to flip is the following:

Main Instruments Vocals (Choir)=false
Variant forming/shaping=true
Main Instruments Voice (Effects/Sample)=true
Main Instruments SFX (Sound Effects)=true
Rhythm funky=true

To confirm the funkiness of the song has been increased, we can verify that within the nearest neighbors (NN) of the modified song there are more 'Funk' songs than initially. Among the 20 NN of the initial "Black or White", 4 songs are tagged as "Funk". Among the 20 NN of its flipped version, 11 songs tagged as "Funk". So the algorithm is indeed able to translate the high-level query "more funk" into a minimal set of low-level modifications.

4. CONCLUSION

We have presented an attribute inference algorithm for large-scale Boolean databases. The algorithm produces optimally plausible inferences in reasonable time. The algorithm can also be used to perform inverse inference, i.e. answering questions about how to modify a description to make it fit optimally a given attribute. The algorithm is particularly well suited to the exploitation of metadata databases, both for traditional search applications, and for applications that exploit these databases for creation purposes. As such, it can be seen as a first step in turning these databases into knowledge bases.

5. ACKNOWLEDGEMENT

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PERFORMER IDENTIFICATION IN CELTIC VIOLIN RECORDINGS

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ABSTRACT

We present an approach to the task of identifying performers from their playing styles. We investigate how violinists express and communicate their view of the musical content of Celtic popular pieces and how to use this information in order to automatically identify performers. We study note-level deviations of parameters such as timing and amplitude. Our approach to performer identification consists of inducing an expressive performance model for each of the interpreters (essentially establishing a performer dependent mapping of inter-note features to a timing and amplitude expressive transformations). We present a successful performer identification case study.

1 INTRODUCTION

Music performance plays a central role in our musical culture today. Concert attendance and recording sales often reflect people's preferences for particular performers. The manipulation of sound properties such as pitch, timing, amplitude and timbre by different performers is clearly distinguishable by the listeners. Expressive music performance studies the manipulation of these sound properties in an attempt to understand expression in performances. There has been much speculation as to why performances contain expression. Hypothesis include that musical expression communicates emotions and that it clarifies musical structure, i.e. the performer shapes the music according to her own intentions

In this paper we focus on the task of identifying violin performers from their playing style using high-level descriptors extracted from single-instrument audio recordings. The identification of performers by using the expressive content in their performances raises particularly interesting questions but has nevertheless received relatively little attention in the past.

The data used in our investigations are violin audio recordings of Irish popular music performances. We use sound analysis techniques based on spectral models [15] for extracting high-level symbolic features from the recordings. In particular, for characterizing the performances used in this work, we are interested in inter-note features representing information about the music context in which expressive events occur. Once the relevant high-level information is extracted we apply machine learning techniques [9] to automatically discover regularities and expressive patterns for each performer. We use these regularities and patterns in order to identify a particular performer in a given audio recording.

The rest of the paper is organized as follows: Section 2 sets the background for the research reported here. Section 3 describes how we process the audio recordings in order to extract inter-note information. Section 4 describes our approach to performance-driven performer identification. Section 5 describes a case study on identifying performers based on their playing style and discusses the results, and finally, Section 6 presents some conclusions and indicates some areas of future research.

2 BACKGROUND

Understanding and formalizing expressive music performance is an extremely challenging problem which in the past has been studied from different perspectives, e.g. [14], [4], [2]. The main approaches to empirically studying expressive performance have been based on statistical analysis (e.g. [12]), mathematical modeling (e.g. [17]), and analysis-by-synthesis (e.g. [3]). In all these approaches, it is a person who is responsible for devising a theory or mathematical model which captures different aspects of musical expressive performance. The theory or model is later tested on real performance data in order to determine its accuracy. The majority of the research on expressive music performance has focused on the performance of musical material for which notation (i.e. a score) is available, thus providing unambiguous performance goals. Expressive performance studies have also been very much focused on (classical) piano performance in which pitch and timing measurements are simplified.

Previous research addressing expressive music performance using machine learning techniques has included a number of

approaches. Lopez de Mantaras et al. [6] report on SaxEx, a performance system capable of generating expressive solo saxophone performances in Jazz. One limitation of their system is that it is incapable of explaining the predictions it makes and it is unable to handle melody alterations, e.g. ornamentations.

Ramirez et al. [11] have explored and compared diverse machine learning methods for obtaining expressive music performance models for Jazz saxophone that are capable of both generating expressive performances and explaining the expressive transformations they produce. They propose an expressive performance system based on inductive logic programming which induces a set of first order logic rules that capture expressive transformation both at an inter-note level (e.g. note duration, loudness) and at an intra-note level (e.g. note attack, sustain). Based on the theory generated by the set of rules, they implemented a melody synthesis component which generates expressive monophonic output (MIDI or audio) from inexpressive melody MIDI descriptions.

With the exception of the work by Lopez de Mantaras et al and Ramirez et al, most of the research in expressive performance using machine learning techniques has focused on classical piano music where often the tempo of the performed pieces is not constant. The works focused on classical piano have focused on global tempo and loudness transformations while we are interested in note-level tempo and loudness transformations.

Nevertheless, the use of expressive performance models, either automatically induced or manually generated, for identifying musicians has received little attention in the past. This is mainly due to two factors: (a) the high complexity of the feature extraction process that is required to characterize expressive performance, and (b) the question of how to use the information provided by an expressive performance model for the task of performance-based performer identification. To the best of our knowledge, the only group working on performance-based automatic performer identification is the group led by Gerhard Widmer. Saunders et al [13] apply string kernels to the problem of recognizing famous pianists from their playing style. The characteristics of performers playing the same piece are obtained from changes in beat-level tempo and beat-level loudness. From such characteristics, general performance alphabets can be derived, and pianists' performances can then be represented as strings. They apply both kernel partial least squares and Support Vector Machines to this data.

Stamatatos and Widmer [16] address the problem of identifying the most likely music performer, given a set of performances of the same piece by a number of skilled candidate pianists. They propose a set of very simple features for representing stylistic characteristics of a music performer that relate to a kind of 'average' performance. A database of piano performances of 22 pianists playing two pieces by Frdric Chopin is used. They propose an ensemble of simple classifiers derived by both subsampling the training set and subsampling the input features. Experiments show that the proposed features are able to quantify the differences between music performers.

3 MELODIC DESCRIPTION

First of all, we perform a spectral analysis of a portion of sound, called analysis frame, whose size is a parameter of the algorithm. This spectral analysis consists of multiplying the audio frame with an appropriate analysis window and performing a Discrete Fourier Transform (DFT) to obtain its spectrum. In this case, we use a frame width of 46 ms, an overlap factor of 50%, and a Keiser-Bessel 25dB window. Then, we compute a set of low-level descriptors for each spectrum: energy and an estimation of the fundamental frequency. From these low-level descriptors we perform a note segmentation procedure. Once the note boundaries are known, the note descriptors are computed from the low-level values. the main low-level descriptors used to characterize note-level expressive performance are instantaneous energy and fundamental frequency.

Energy computation. The energy descriptor is computed on the spectral domain, using the values of the amplitude spectrum at each analysis frame. In addition, energy is computed in different frequency bands as defined in [5], and these values are used by the algorithm for note segmentation.

Fundamental frequency estimation. For the estimation of the instantaneous fundamental frequency we use a harmonic matching model derived from the Two-Way Mismatch procedure (TWM) [7]. For each fundamental frequency candidate, mismatches between the harmonics generated and the measured partials frequencies are averaged over a fixed subset of the available partials. A weighting scheme is used to make the procedure robust to the presence of noise or absence of certain partials in the spectral data. The solution presented in [7] employs two mismatch error calculations. The first one is based on the frequency difference between each partial in the measured sequence and its nearest neighbor in the predicted sequence. The second is based on the mismatch between each harmonic in the predicted sequence and its nearest partial neighbor in the measured sequence. This two-way mismatch helps to avoid octave errors by applying a penalty for partials that are present in the measured data but are not predicted, and also for partials whose presence is predicted but which do not actually appear in the measured sequence. The TWM mismatch procedure has also the benefit that the effect of any spurious components or partial missing from the measurement can be counteracted by the presence of uncorrupted partials in the same frame.

Note segmentation is performed using a set of frame descriptors, which are energy computation in different frequency bands and fundamental frequency. Energy onsets are first detected following a band-wise algorithm that uses some psycho-acoustical knowledge [5]. In a second step, fundamental frequency transitions are also detected. Finally, both results are merged to find the note boundaries (onset and offset information).

Note descriptors. We compute note descriptors using the note boundaries and the low-level descriptors values. The low-level descriptors associated to a note segment are computed by averaging the frame values within this note segment. Pitch histograms have been used to compute the pitch note and the fundamental frequency that represents each note segment, as found in [8]. This is done to avoid taking into account mistaken frames in the fundamental frequency mean computation. First, frequency values are converted into cents, by the following formula:

$$c = 1200 \cdot \frac{\log\left(\frac{f}{f_{ref}}\right)}{log2} \quad (1)$$

where $f_{ref} = 8.176$ (fref is a the reference frequency of the C0). Then, we define histograms with bins of 100 cents and hop size of 5 cents and we compute the maximum of the histogram to identify the note pitch. Finally, we compute the frequency mean for all the points that belong to the histogram. The MIDI pitch is computed by quantization of this fundamental frequency mean over the frames within the note limits.

Musical Analysis. It is widely recognized that humans perform music considering a number of abstract musical structures. In order to provide an abstract structure for the recordings under study, we decided to use Narmour's theory of perception and cognition of melodies [10] to analyze the performances.

The Implication/Realization model proposed by Narmour is a theory of perception and cognition of melodies. The theory states that a melodic musical line continuously causes listeners to generate expectations of how the melody should continue. The nature of these expectations in an individual are motivated by two types of sources: innate and learned. According to Narmour, on the one hand we are all born with innate information which suggests to us how a particular melody should continue. On the other hand, learned factors are due to exposure to music throughout our lives and familiarity with musical styles and particular melodies. According to Narmour, any two consecutively perceived notes constitute a melodic interval, and if this interval is not conceived as complete, it is an implicative interval, i.e. an interval that implies a subsequent interval with certain characteristics. That is to say, some notes are more likely than others to follow the implicative interval. Two main principles recognized by Narmour concern registral direction and intervallic difference. The principle of registral direction states that small intervals imply an interval in the same registral direction (a small upward interval implies another upward interval and analogously for downward intervals), and large intervals imply a change in registral direction (a large upward interval implies a downward interval and analogously for downward intervals). The principle of intervallic difference states that a small (five semitones or less) interval implies a similarly-sized interval (plus or minus 2 semitones), and a large interval (seven semitones or more) implies a smaller interval. Based on these two principles, melodic patterns or groups can be identified that either satisfy or violate the implication as predicted by the principles. Such patterns are called structures and are labeled to denote characteristics in terms of registral direction and intervallic difference. Figure 1 shows prototypical Narmour structures. A note in a melody often belongs to more than one structure. Thus, a description of a melody as a sequence of Narmour structures consists of a list of overlapping structures. We parse each melody in the training data in order to automatically generate an implication/realization analysis of the pieces. Figure 2 shows the analysis for a fragment of a melody.

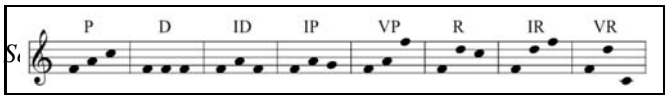

Figure 1. Prototypical Narmour Structures

Figure 2. Narmour analysis of a melody fragment

4 PERFORMANCE-DRIVEN PERFORMER IDENTIFICATION

4.1 Note features

The note features represent both properties of the note itself and aspects of the musical context in which the note appears. Information about the note includes note pitch and note duration, while information about its melodic context includes the relative pitch and duration of the neighboring notes (i.e. previous and following notes) as well as the Narmour structures to which the note belongs. The note's Narmour structures are computed by performing the musical analysis described before. Thus, each performed note is characterized

by the tuple

(Pitch, Dur, PrevPitch, PrevDur, NextPitch, NextDur, Nar1, Nar2, Nar3)

4.2 Algorithm

We are ultimately interested in obtaining a classification function F of the following form:

$$F(MelodyFragment(n1, \ldots, nk)) \longrightarrow Performers$$

where $MelodyFragment(n1, \ldots, nk)$ is the set of melody fragments composed of notes $n1, \ldots, nk$ and $Performers$ is the set of possible performers to be identified. For each performer i to be identified we induce an expressive performance model M_i predicting his/her timing and energy expressive transformations:

$$M_i(Notes) \rightarrow (PDur, PEner)$$

where $Notes$ is the set of score notes played by performer i represented by their inter-note features, i.e. each note in $Notes$ is represented by the tuple (Pitch, Dur, PrevPitch, PrevDur, NextPitch, NextDur, Nar1, Nar2, Nar3) as described before, and the vector $(PDur, PEner)$ contains the model's predictions for note duration ($PDur$) and energy ($PEner$). Once a performance model is induced for each performer P_i we apply the following algorithm:

```
F([N1,...,Nm], [P1,...,Pn])
   for each performer Pi
      Scorei = 0
   for each note Nk
      FNk = inter-note_features(Nk)
      Mi(FNk) = (PDk,PEk)
      for each performer Pi
         ScoreNKi = sqrt(((Dur(NK)-PDk)^2)
                  + ((Ener(Nk)-PEk)^2))
         Scorei = Scorei  + ScoreNKi
return Pj (j in {1,...,m}) with minimum score
```

This is, for each note in the melody fragment the classifier F computes the set of its inter-note features. Once this is done, for each note N_k and for each performer P_i, performance model M_i predicts the expected duration and energy for N_k. This prediction is based on the note's inter-note features. The score $Score_i$ for each performer i is updated by taking into account the Euclidean distance between the note's actual duration and energy and the predicted values. Finally, the performer with the lower score is returned.

Clearly, the expressive models M_i play a central role in the output of classifier F. For each performer, M_i is induced by applying Tildes top-down decision tree induction algorithm ([1]). Tilde can be considered as a first order logic extension of the C4.5 decision tree algorithm: instead of testing attribute values at the nodes of the tree, Tilde tests logical predicates. This provides the advantages of both propositional decision trees (i.e. efficiency and pruning techniques) and the use of first order logic (i.e. increased expressiveness). The musical context of each note is defined by predicates $context$ and $narmour$. $context$ specifies the note features described above and $narmour$ specifies the Narmour groups to which a particular note belongs, along with its position within a particular group. Expressive deviations in the performances are encoded using predicates $stretch$ and $dynamics$. $stretch$ specifies the stretch factor of a given note with regard to its duration in the score and $dynamics$ specifies the mean energy of a given note. The temporal aspect of music is encoded via the predicates $pred$ and $succ$. For instance, $succ(A, B, C, D)$ indicates that note in position D in the excerpt indexed by the $tuple(A, B)$ follows note C.

5 CASE STUDY

5.1 Training data

In this work we are focused on Celtic jigs, fast tunes but slower that reels, that usually consist of eighth notes in a ternary time signature, with strong accents at each beat. The training data used in our experimental investigations are monophonic recordings of nine Celtic jigs performed by two professional violinists. Apart from the tempo (they played following a metronome), the musicians were not given any particular instructions on how to perform the pieces.

5.2 Results

Initially, we evaluated the expressive performance model for each of the musicians we considered. Thus, we obtained two expressive performance models M_1 and M_2. For M_1 we obtained correlation coefficients of 0.88 and 0.83 for the duration transformation and note dynamics prediction tasks, respectively, while we obtained 0.91 and 0.85 for M_2. These numbers were obtained by performing 10-fold cross-validation on the training data. The induced models seem to capture accurately the expressive transformations the musicians introduce in the performances. Figure 3 contrasts the note duration deviations predicted by model M_1 and the deviations performed by the violinist. Similar results were obtained for M_2.

We then proceed to evaluate the classification function F by spliting our data into a training set and a test set. We held out approximately 30% of the data as test data while the remaining 70% was used as training data (we held out 3 pieces for each violinist). When selecting the test data, we left out the same number of melody fragments per class. In order to avoid optimistic estimates of the classifier performance, we

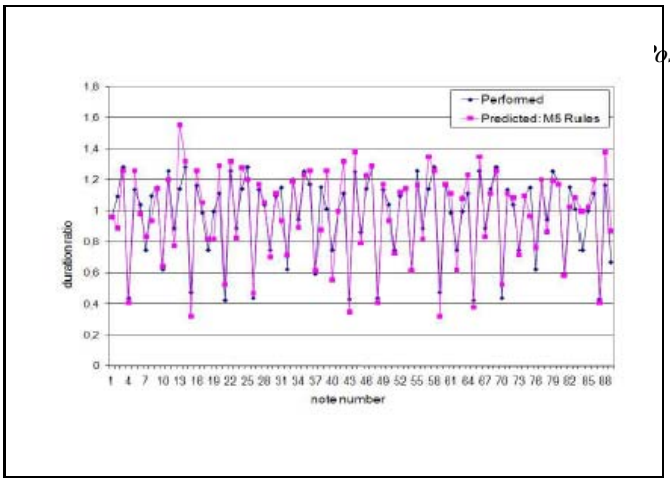

Figure 3. Note deviation ratio for a tune with 89 notes. Comparison between performed and predicted by M_1

explicitly removed from the training set all melody fragment repetitions of the hold out fragments. This is motivated by the fact that musicians are likely to perform a melody fragment and its repetition in a similar way. We tested our algorithm in each of the six test pieces (three pieces of each class) and obtained 100% accuracy (correctly classified instances percentage). This is, the six pieces in the test set were classified correctly.

6 CONCLUSION

In this paper we focused on the task of identifying performers from their playing style using note descriptors extracted from audio recordings. In particular, we concentrated in identifying violinists playing Irish popular pieces (Irish jigs). We characterized performances by representing each note in the performance by a set of inter-note features representing the context in which the note appears. We then induced an expressive performance model for each of the performers and presented a successful performer identification case study. The results obtained seem to indicate that the inter-note features presented contain sufficient information to identify the studied set of performers, and that the machine learning method explored is capable of learning performance patterns that distinguish these performers. This paper present preliminary work so there is further work in several directions. Our immediate plans are to extend our database and to test different distance measures for updating the performer scores in our algorithm. We also plan to evaluate our algorithm considering melody fragments of different size and to evaluate the predictive power of timing expressive variations relative to energy expressive variations.

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MACHINE ANNOTATION OF SETS OF TRADITIONAL IRISH DANCE TUNES

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ABSTRACT

A *set* in traditional Irish music is a sequence of two or more dance tunes in the same time signature, where each tune is repeated an arbitrary number of times. A *turn* in a set represents the point at which either a tune repeats or a new tune is introduced. Tunes in sets are played in a *segue* (without a pause) and so detecting the turn is a significant challenge. This paper presents the MATS algorithm, a novel algorithm for identifying turns in sets of traditional Irish music. MATS works on digitised audio files of monophonic flute and tin-whistle music. Previous work on machine annotation of traditional music is summarised and experimental results validating the MATS algorithm are presented.

1. INTRODUCTION

Several papers address the necessity of developing MIR (Music Information Retrieval) systems that are adapted to the specific requirements of ethnic music and also to the needs of musicologists studying ethnic music [1-3]. While there are MIR systems that allow users to search for traditional Irish dance tunes using text based musical queries [4,5] and there are MIR systems that allow users to search for melodies using sung queries [6,7], there are no MIR systems that we are aware of that allow musicians to search for traditional Irish dance tunes using queries played on traditional instruments. Some examples of the above include the website thesession.org [4] which contains an extensive collection of over seven thousand traditional dance tunes in the ABC language; the system supports text queries by any of the metadata associated with a tune or melodic queries in the ABC language. Similarly, Melodyhound [6] a publicly accessible MIR system that supports sung queries and contains a large collection of traditional Irish dance tunes does not generate positive results when queries are presented in the form of melodies played on the tin-whistle or wooden flute.

Such a system would have many applications in the field of music archiving and retrieval, particularly given the many thousands of hours of archive music collected by organisations involved in the cataloguing of traditional music such as Na Píobairí Uilleann, Comhaltas Ceoltóirí Éireann and the Irish Traditional Music Archive. Similarly it is common at traditional music sessions, recitals and even on commercial recordings for tunes to be named *gan ainm* (without name) when the tune in question does in fact have a name, composer and history. For a typical example see the CD recording [8].

Previous work proposes MATT2 (Machine Annotation of Traditional Tunes) as a system that can identify tunes played on either the flute or the tin whistle [9]. MATT2 takes advantage of a number of novel subsystems that significantly increase matching accuracy for traditional tunes played in a variety of regional styles by different musicians. These include an onset detection function developed for windblown instruments, an ornamentation compensation algorithm based on fuzzy histograms, a two thousand tune corpus of tunes in the ABC language (a natural fit for traditional music) and a melody normalisation algorithm that adapts tunes in the corpus to the way they might be played by a human musician. MATT2 is described in detail in [9] and we present an overview in section 3. The main purpose of this paper is to present our enhancements to the MATT2 system and specifically to present a new algorithm for annotating sets of traditional Irish dance tunes. Previous versions of MATT2 could only annotate single tunes, however in traditional music tunes are rarely played singly. More commonly tunes are played in groups of at least two tunes known as a *set* of tunes. A set typically consists of two three or four tunes played in succession without an interval [10,9]. Typically each tune in the set is played twice or three times before musicians advance to the subsequent tune in the set. A repetition or a change from one tune to the next in a set is known as a *turn*. As tunes in sets are always in the same time signature and often in the same key, the challenge therefore is in segmenting sets into tunes and repetitions. The approach presented in this paper tackles this problem by making use of melodic similarity calculated using a variant of the *edit distance* string matching algorithm described in section 3. The MATS algorithm described in this paper can identify the start and end of each repetition of a tune, can count the repetitions and can identify the title and associated metadata associated with each tune in a set.

Section 2 of this paper briefly explains the domain of traditional Irish dance music. In Section 3 existing work on the MATT2 system is presented. Section 4 presents MATS (Machine Annotation of Traditional Sets), a novel

annotation algorithm which annotates sets of traditional tunes. Section 5 presents experimental results which establish the effectiveness of this new algorithm and section 6 presents conclusions and future work.

2. TRADITIONAL IRISH DANCE MUSIC

The most common forms of dance music are *reels, double jigs* and *hornpipes*. Other tune types include *marches*, *set dances*, *polkas, mazurkas, slip jigs, single jigs and reels, flings, highlands, scottisches, barn dances, strathspeys* and *waltzes* [11]. These forms differ in time signature, tempo and structure. For example a reel is generally played at a lively tempo and is in 4/4 time (four crochets in a bar, though usually transcribed as eight quavers in a bar), while a waltz is generally played at slower pace and is in 3/4 time. Most tunes consist of a common structure of two parts called the *A* part and *B* part. Tunes are typically played as *sets*. Certain common sets were originally put together to accompany set dances [10], while other sets have become popular as a result of being recorded by emigrant Irish musicians in America in the early part of the twentieth century.

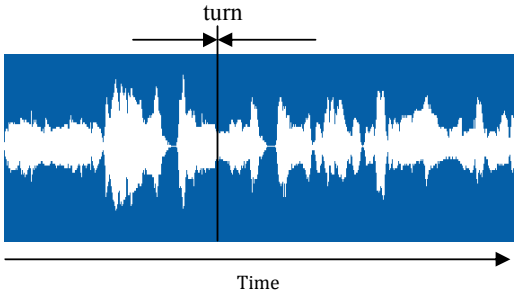

Figure 1: Waveform of the last phrase from the tune "Jim Coleman's" and the first phrase from the tune "George Whites Favourite" played in a set

The origin of many sets of tunes is unknown and musicians often compile new sets "on the fly" in traditional music sessions. Figure 1 shows a waveform plot from two tunes played in a set. The tunes were played on a wooden flute and as can be seen in the plot, there is no interval between the end of the first tune and the start of the second tune. Maddage *et al.* and other segmentation approaches generally look for repetitive patters in a music recording [12]. This is not the case in our approach, where each tune in the set can be played once or many times.

When a traditional musician plays a tune, it is rarely played exactly as transcribed. In fact an experienced musician never plays the same tune twice identically, employing the subtleties of *ornamentation* and *variation* to interpret the tune [11]. For a discussion on the use of ornamentation in traditional music we refer to [11,13,14].

Ornamentation plays a key role in the individual interpretation of traditional Irish music [10]. The usage of ornamentation is highly personal and large variations exist in the employment of ornamentation from region to region, instrument to instrument and from musician to musician. Tansey colourfully describes ornamentation in the following way:

> "*I put it to you therefore that it had to come from the throats of birds, the wild animals, the ancient chants of our forefathers, the hum of the bees and the mighty rhythms of the galloping hooves of wild horses all moulded together…*" [15]

Ornamentation is difficult to detect correctly and state of the art ornamentation detection algorithms report a success rate of just 40% for multi-note ornaments [16,17]. Similarly, related work in classical music suggests that the playing of ornamentation (grace notes) requires adaptation of melodic similarity measures [18].

It is clear from this brief introduction that an MIR system for traditional dance music must therefore deal with many special problems, such as stylistic variation even within the same instance of a tune, the use of ornamentation which can skew melodic similarity measures and the collection of tunes into sets creating segmentation problems. Transposition invariance is not a requirement for MIR in traditional music as it is uncommon for tunes to be transposed into different keys [16].

3. MACHINE ANNOTATION OF TRADITIONAL TUNES (MATT2)

MATT2 works on mono, digital audio files in the WAV format recorded at 44KHz. A high level diagram of the subsystems that make up MATT2 are presented in Figure 2. MATT2 is described in detail in [9] and so a brief description is presented here.

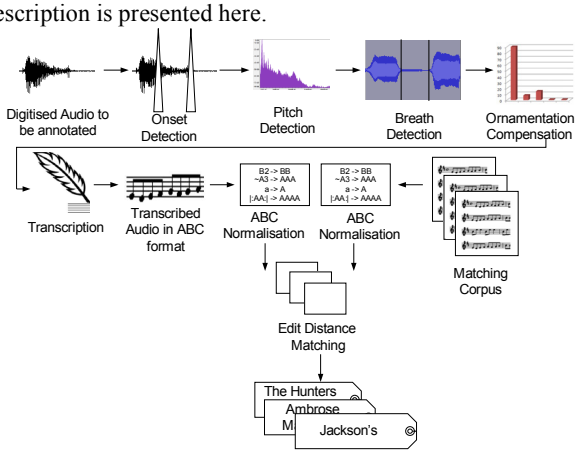

Figure 2: High level diagram of the MATT2 tune annotator

The audio file to be annotated is first segmented into candidate note onsets using an onset detection function

adapted from Gainza [16,17]. The onset detection function ODCF is based on time domain FIR (Finite Impulse Response) comb filters. ODCF discovers harmonic characteristics of the input signal and is therefore useful for detecting onsets in *legato* playing typical of windblown traditional instruments such as the flute and the tin whistle.

In order to detect the perceived pitch of a frame, the pitch detection sub-system performs a STFT (Short Term Fast Fourier Transform) on segments bounded by onsets detected by the onset detection system. The algorithm then calculates the pitch as being the interval between the two most prominent peaks in the FFT graph. This simple approach works well for the harmonics of the wooden flute and the tin whistle.

MATT2 incorporates a breath detector subsystem to transcribe a breath in the signal. A breath is marked if either the pitch detected by the pitch detector is less than 100Hz or the average amplitude of a candidate note c*n* is less than a 10% threshold *th* of the average amplitude of the entire signal *s*. Breaths detected before the transcription of the first pitched note and at the end of the transcription are ignored by the system.

MATT2 uses a heuristic to determine if the input signal was generated by a tin whistle or a wooden flute. A tin whistle in the key of D is pitched exactly one octave above a flute in the key of D, so if the algorithm counts more notes with a pitch above G5 (783.99hz) than below G5, then the algorithm concludes that the input signal contains a tin whistle and the pitches in the pitch spelling algorithm are shifted up accordingly.

Both the wooden flute and the tin whistle have a range of two octaves, though this can be extended by cross fingering techniques [11,19,13]. To tag each candidate note *cn* with a pitch spelling *pS(cn)*, each calculated note frequency is compared with the frequencies of the notes in the key of D4 Major and D5 Major $k_1 \ldots k_{16}$ the two octaves playable on a wooden flute.

The system eliminates notes whose durations are close to zero by merging their durations with subsequent notes. This has the effect of eliminating consecutive onsets (false positives in the ODF caused by noisy onsets) and also eliminating ornamentation notes such as those found in *rolls*, *cuts taps* and *crans* typical of traditional Irish music [11,20,19,13,15]. To achieve this, the quantisation subsystem first generates a histogram of all the note durations. The histogram bin with the highest value is considered to be the length of a quaver note. The histogram counts notes within +/-30% of the bin width. The algorithm also updates the bin width each time a candidate is counted, so that the bin widths represent the cumulative average lengths of notes counted. A transcription *t* is then generated in the ABC language of the input signal from the features extracted by the subsystems in MATT2.

MATT2 has a corpus *Z* of two thousand known tunes (and variations) in the ABC language drawn from the collections of Norbeck [21]. To identify a tune, MATT2 firstly normalises both the transcription *t* and each string *c* $\in Z$. This process is described in detail in [9]. Normalisation minimises the effect of transcription errors and stylistic variation on the calculation of melodic similarity. The *edit distance* is then calculated for *t* in every $c \in Z$ and the tune with the lowest edit distance is returned as a match.

Edit distance, also known as *Levenshtein distance* or *evolutionary distance* [22,23], is a concept from information retrieval and it describes the number of edits (insertions, deletions and substitutions) that have to be made in order to change one string to another. It is the most common measure to expose the similarity between 2 strings.

The edit distance *ed(x, y)* between strings $x=x_1 \ldots x_m$ and $y=y_1 \ldots y_n$, where $x, y \in \Sigma*$ is the minimum cost of a sequence of editing steps required to convert *x* into *y*. Σ is the alphabet of possible characters and $\Sigma*$ is the set of all possible sequences of $ch \in \Sigma$. Edit distance can be calculated using dynamic programming [23]. To compute the edit distance *ed(x,y)* a matrix $M_{1\ldots m+1,1\ldots n+1}$ is constructed where $M_{i,j}$ is the minimum number of edit operations needed to match $x_{1\ldots i}$ to $y_{1\ldots j}$. Each matrix element $M_{i,j}$ is calculated as per (1). The minimum edit distance between *x* and *y* is given by the matrix entry at position $M_{m+1,n+1}$.

$$M_{i,1} \leftarrow i-1, M_{1,j} \leftarrow j-1 \qquad \text{if } x_i=y_i$$
$$M_{i,j} \leftarrow \begin{cases} M_{i-1,j-1} & \text{else} \\ 1+\min\left(M_{i-1,j-1}, M_{i-1,j}, M_{i,j-1}\right) \end{cases} \qquad (1)$$

The algorithm can be adapted to find the lowest edit distances for *x* in substrings of *y*. This is achieved by setting $M_{1,j} = 0$ for all $j \in 1 \ldots n+1$. In contrast to the edit distance algorithm described above, the last row $M_{m+1,j}$ is then used to give a *sliding window* edit distance for *x* in substrings of *y* [23].

		D	G	G	G	D	G	B	D	E	F	G	A	B
	0	0	0	0	0	0	0	0	0	0	0	0	0	0
B	1	1	1	1	1	1	1	0	1	1	1	1	1	0
D	2	1	2	2	2	1	2	1	0	1	2	2	2	1
E	3	2	2	3	3	2	2	2	1	0	1	2	3	2
E	4	3	3	3	4	3	3	3	2	1	1	2	3	3

Table 1: Edit distance for the string BDEE in DGGGDGBDEFGAB. This string represents the first 13 notes from the tune "Jim Coleman's" in normalised ABC format

An example of this variation on the edit distance applied to search for the pattern "BDEE" in "DGGGDGBDEFGAB" is given in Table 1. The

minimum edit distance positions are highlighted. Variations on the edit distance algorithm have been applied in domains such as DNA analysis and automated spell checking and are commonly used in MIR systems [7,24].

With test input drawn from the playing of ten different musicians playing flute, whistle and fiddle, the system was able to correctly identify the tune in 86% of cases. In 96% of cases, the correct tune was identified within the top five closest matches [9].

4. MACHINE ANNOTATION OF TRADITIONAL SETS ALGORITHM (MATS)

In this section MATS is described. MATS is an enhancement to MATT2 described in the previous section. The purpose of MATS is to annotate tunes played in sets.

The shortest tune in the corpus *Z* used by MATT2 is a single jig. A single jig *sj* is a tune in 6/8 time with an A and B part played singly (48 quaver notes in duration). The shortest possible set therefore would contain two single jigs (96 notes) played with no repetitions. To annotate a set of tunes, MATS first uses a heuristic to determine if the string of transcribed notes *t* is longer than the length of the shortest set *length(sj)×2*.

When this is the case, the MATS algorithm is used instead of the minimum edit distance algorithm described in section 3. Pseudocode for the MATS algorithm is presented in Figure 3.

MATS first extracts a substring *ss* from *t* the transcription such that *length(ss) = length(sj)* at position *p=1* in *t*. MATS then searches the corpus *Z* using the edit distance algorithm described in section 3 to find a the closest match for *ss*. When a match is found MATS knows the name of the first tune and has *c′*, a transcription of the tune played with no repetitions from the corpus Z. MATS then generates an edit distance profile *edp* for *c′,* the matching tune, in *t* the transcription. *edp* is given as the last row of the edit distance matrix and can be understood as the positions where substrings in *t* match *c′* with the minimum edit distance .

Figure 4 shows the edit distance profiles for the set of tunes "Jim Coleman's", "George Whites Favourite" and "the Virginia" played in a set. The algorithm has identified the first tune as "Jim Coleman's" and has subsequently generated an edit distance profile (the top plot in Figure 4) for the first tune in the transcription. The two troughs in this graph indicate the end of the two repetitions of the tune in the transcription. These can be considered as turns in the set.

The MATS algorithm then normalises the edit distance profile *edp* and passes the graph through a low pass filter that filters frequencies less than 10Hz. This has the effect of smoothing the graph. An example of a smoothed edit distance profile is given in Figure 5. This graph illustrates the top graph in Figure 4 after filtering has been applied.

The algorithm then detects troughs in the graph less than a threshold initially set to *t=0.3*. The algorithm varies this threshold dynamically by trying different values until the number of troughs in the graph is between one and five. It is rare in traditional music for a tune to be repeated more than five times in a set.

```
p ← 0
rem ← length(t) - p
while (rem >= sj)
begin
   ss ← substring(t, p, p + sj)
   foreach (c in Z)
   begin
        ed_c ← min(ed(ss, c))
        if (ed_c < min_ed)
        begin
           min_ed ← ed_c
           c' ← c
        end
   end
   edp ← ed(c', t)
   edp ← normalise(edp)
   edp ← filter(edp, 10)
   th ← 0.3
   v ← troughs(edp, th)
   foreach (tr in v)
   begin
      convertToTime(tr)
   end
   r ← length(v)
   p ← v[r]
   print c', r
   rem ← length(t) - p
end
```

Figure 3: Pseudocode for the MATS set annotation algorithm

The trough detection algorithm in MATS returns a vector of troughs $\vec{v}$, such that *length($\vec{v}$)* is the number of troughs and the elements in $\vec{v}$ are the positions of the bottom of the troughs. A trough in MATS need only have a descending wall as a trough can occur at the end of a tune and hence may not contain an ascending wall. An example of this is the third plot in Figure 4.

The algorithm repeats this process with a new *p* given as the last entry in the troughs vector to extract the second and subsequent tunes in the set until it is no longer possible to extract a substring *ss* of length *length(sj)* starting at *p* because we have reached the end of *t*. The second tune in the set, "George Whites Favourite" was played once and there is a corresponding single trough in the graph of the edit distance function (the middle plot in Figure 4) for the tune from the corpus *c′* in the transcription *t*. The third tune "the Virginia" was repeated twice and so there are two troughs in the bottom plot in Figure 4.

5. RESULTS

In order to test the robustness of MATS we had a traditional musician record ten audio files of flute tunes played in sets. The recorded files are available at http://www.comp.dit.ie/bduggan/mats. The sets played in the input audio were taken from the Foinn Seisiún series of books published by Comhaltas Ceoltóirí Éireann [25].

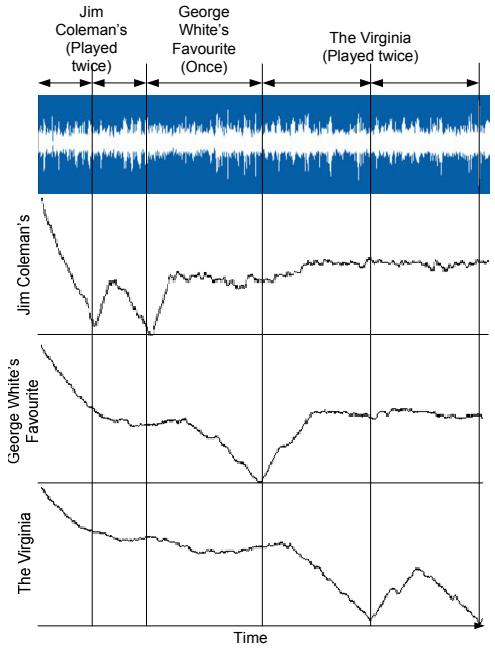

Figure 4: Edit distance profiles for three tunes played in a set

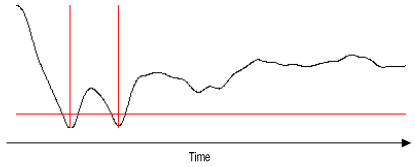

Figure 5: Filtered version of first graph in Figure 4. The dynamic threshold and detected troughs are marked

The sets consisted of single and double jigs and reels played multiple times in sets. In total, the sets contained 23 separate tunes with 48 turns we were interested in annotating. In carrying out this experiment, we were interested in establishing if MATT2 could correctly figure out the timings of turns and could identify the names of the tunes.

Correctly identified	96%
Incorrectly identified	4%

Table 2: Correctly and incorrectly identified tunes

MATT2 successfully identified 22 out of the 23 tunes, and recognised each input audio file as a set and so used the MATS set annotation algorithm (Table 2).

Table 3 shows a sample of the data collected in this experiment for the audio file used to generate Figure 4 and Figure 5. To establish a ground truth for the experiment, a human domain expert manually annotated the turns in the sets of tunes. In the human and machine columns are listed the onset time for turns in the set. Onset times for changes from one tune to the next are highlighted. From this table it can be seen that on average MATS was within .85 seconds of the human annotations.

Tune	Human	Machine	Difference
1	20.68	21.10	0.43
1	41.42	41.9	0.48
2	82.72	83.15	0.43
3	123.88	124.44	0.56
3	164.49	166.85	2.36
Average			0.85

Table 3: Human & machine annotated turns

The overall annotation accuracy is obtained by calculating two different measures *precision* and *recall*. The value of *precision* is calculated as per (2) where *TP* and *FP* are the true positives (correctly identified turns) and false positives (incorrectly identified turns). *recall* is calculated as per equation (3) where *FN* is the number of false negatives (turns in the input signal not detected by the algorithm).

$$precision = \frac{TP}{TP + FP} \quad (2)$$

$$recall = \frac{TP}{TP + FN} \quad (3)$$

TP	FN	FP	precision(%)	recall(%)
39	9	6	87%	81%

Table 4: Annotation accuracy

Table 4 shows the annotation accuracy. In can be seen from *precision* and *recall* that the algorithm provides a high degree of accuracy at detecting turns. Because the algorithm can successfully identify turns, it can also correctly extract a suitable prefix from the subsequent tune in the set and so can identify the tune. *FN*'s were caused by the algorithm failing to correctly identify the transitions between tunes in a set. When this happens the algorithm cannot extract a representative prefix from the next tune and so all subsequent turns are usually misidentified. In

some cases, *FP*'s were within a few seconds of the two second threshold we had set.

6. CONCLUSIONS & FUTURE WORK

This paper presented a novel algorithm that addresses a problem in the domain of Irish traditional dance music, that of annotating sets of tunes. As a set can contain an arbitrary number of tunes played segue without an interval and as tunes in sets are repeated an arbitrary number of times, are always in the same time signature and often in the same key, the significant challenge in this problem is in recognising where one tune ends and the next tune starts. The results presented prove that MATS is effective at segmenting sets, counting repetitions and at annotating individual tunes played in a set. To our knowledge this is the first time this problem has been addressed in an MIR system and we suggest that the proposed approach can be adapted to segmenting repeated tunes from other genres played in a segue.

The corpus used currently contains reels and jigs and in future work it will be augmented with the full complement of traditional tunes in different time signatures. One interesting feature not yet exploited is the metadata typically present in an ABC transcription. Effectively the time and key signature of an input audio file can be determined by *melodic similarity* with a known tune. This can be exploited in several interesting ways. Firstly, if the first tune in a set were to be identified as a reel, the search for subsequent tunes can be limited to reels, thus speeding up annotation. Conversely, if a number of reels were to be identified in a set and a single tune in a different time signature was to be identified this could be recognised as a potential error.

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RESOLVING OVERLAPPING HARMONICS FOR MONAURAL MUSICAL SOUND SEPARATION USING PITCH AND COMMON AMPLITUDE MODULATION

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ABSTRACT

In mixtures of pitched sounds, the problem of overlapping harmonics poses a significant challenge to monaural musical sound separation systems. In this paper we present a new algorithm for sinusoidal parameter estimation of overlapping harmonics for pitched instruments. Our algorithm is based on the assumptions that harmonics of the same source have correlated amplitude envelopes and the phase change of harmonics can be accurately predicted from an instrument's pitch. We exploit these two assumptions in a least-squares estimation framework to resolve overlapping harmonics. This new algorithm is incorporated into a separation system and quantitative evaluation shows that the resulting system performs significantly better than an existing monaural music separation system for mixtures of harmonic instruments.

1 INTRODUCTION

Musical sound separation attempts to isolate the sound of individual instruments in a polyphonic mixture. In recent years this problem has attracted significant attention as the demand for automatic analysis, organization, and retrieval of a vast amount of online music data has exploded. A solution to this problem allows more efficient audio coding, more accurate content-based analysis, and more sophisticated manipulation of musical signals [1]. In this paper, we address the problem of monaural musical sound separation, where multiple harmonic instruments are recorded by a single microphone or mixed to a single channel.

A well known difficulty in music separation arises when the harmonics of two or more pitched instruments have frequencies that are the same or similar. Since Western music favors the twelve-tone equal temperament scale [2], common musical intervals have pitch relationships very close to simple integer ratios (≈3/2, 4/3, 5/3, 5/4, etc.). As a consequence, a large number of harmonics of a given source may be overlapped with another source in a mixture.

When harmonics overlap, the amplitude and phase of individual harmonics become unobservable. To recover an overlapped harmonic, it has been assumed that the amplitudes of instrument harmonics decay smoothly as a function of frequency [3]. Based on this assumption, the amplitude of an overlapped harmonic can be estimated from the amplitudes of neighboring non-overlapped harmonics of the same source. For example, Virtanen and Klapuri [4] estimated an overlapped harmonic through non-linear interpolation of neighboring harmonics. Every and Szymanski [5] used linear interpolation instead. Recently, Virtanen [1] proposed a system which directly imposes spectral smoothness by modeling the amplitudes of harmonics as a weighted sum of fixed basis functions having smooth spectral envelopes. However, for real instrument sounds, the spectral smoothness assumption is often violated (see Figure 1). Another method of dealing with overlapping harmonics is to use instrument models that contain the relative amplitudes of harmonics [6]. However, models of this nature have limited success due to the spectral diversity in recordings of different notes, different playing styles, and even different builds of the same instrument type.

Although in general, the absolute value of a harmonic's amplitude with respect to its neighboring harmonics is difficult to model, the amplitude envelopes of different harmonics of the same source exhibit similar temporal dynamics. This is known as common amplitude modulation (CAM) and it is an important organizational cue in human auditory perception [7] and has been used in computational auditory scene analysis [8]. Although CAM has been utilized for stereo music separation [9, 10], to our knowledge, this cue has not been applied in existing monaural systems. In this paper we demonstrate how CAM can be used to resolve overlapping harmonics in monaural music separation.

Many existing monaural music separation systems operate only in the amplitude/magnitude domain [5, 6, 11, 12]. However, the relative phase of overlapping harmonics plays a critical role in the observed amplitude of the mixture and must be considered in order to accurately recover the amplitudes of individual harmonics. We will show that the phase change of each harmonic can be accurately predicted from the signal's pitch. When this and the CAM observation are combined within a sinusoidal signal model, both the amplitude and phase parameters of overlapping harmonics can be accurately estimated.

This paper is organized as follows. Section 2 presents the sinusoidal model for mixtures of harmonic instruments. In Section 3 we justify the CAM and phase change prediction assumptions and propose an algorithm where these assumptions are used in a least-squares estimation framework for resolving overlapping harmonics. In Section 4 we present a monaural music separation system which incorporates the proposed algorithm. Section 5 shows quantitative evaluation results of our separation system and Section 6 provides a final discussion.

2 SINUSOIDAL MODELING

Modeling a harmonic sound source as the summation of individual sinusoidal components is a well established technique in musical instrument synthesis and audio signal processing [13, 14]. Within an analysis frame m where sinusoids are assumed constant, the sinusoidal model of a mixture consisting of harmonic sounds can be written as

$$x_m(t) = \sum_n \sum_{h_n=1}^{H_n} a_n^{h_n}(m)\cos(2\pi f_n^{h_n}(m)t + \phi_n^{h_n}(m)), \quad (1)$$

where $a_n^{h_n}(m)$, $f_n^{h_n}(m)$, and $\phi_n^{h_n}(m)$ are the amplitude, frequency, and phase of sinusoidal component h_n, respectively, of source n at time frame m. H_n denotes the number of harmonics in source n. The sinusoidal model of $x_m(t)$ can be transformed to the spectral domain by using the discrete Fourier transform (DFT). With an appropriately chosen time-domain analysis window (in terms of frequency resolution and sidelobe suppression), and assuming perfect harmonicity, the spectral value of $x_m(t)$ at frequency bin k can be written as

$$X(m,k) = \sum_n S_n^{h_n}(m)W(kf_b - h_nF_n(m)). \quad (2)$$

Here, W is the complex-valued DFT of the analysis window, f_b is the frequency resolution of the DFT, and $F_n(m)$ denotes the pitch of source n at time frame m. We call $S_n^{h_n}(m)$ the sinusoidal parameter of harmonic h_n of source n, where $S_n^{h_n}(m) = \frac{a_n^{h_n}(m)}{2}e^{i\phi_n^{h_n}(m)}$. As a proof of concept, we further assume that pitches of individual sources are known.

3 RESOLVING OVERLAPPING HARMONICS

Given ground truth pitches of each source, one can identify non-overlapped and overlapped harmonics. When a harmonic of a source is not overlapped, the estimation of the sinusoidal parameter, $S_n^{h_n}(m)$, from observed spectral values, $X(m,k)$, in corresponding frequency bins is straightforward (see Section 4). However, when harmonics from different sources overlap, finding $S_n^{h_n}(m)$ for each active harmonic is an ill-defined problem. To address this, we make use of non-overlapped harmonics of the same source as well as the phase change estimated from the pitch information.

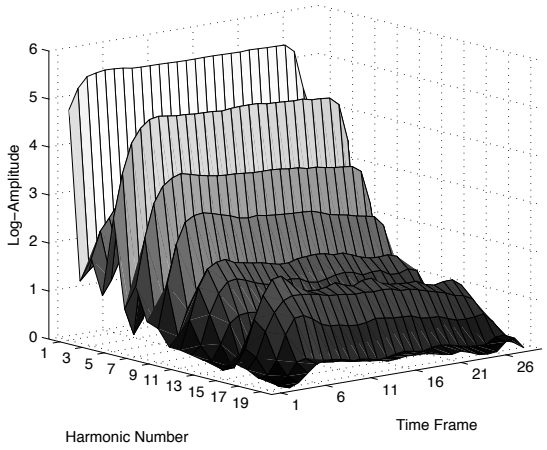

Figure 1. Logarithm of the amplitude envelopes for the first 20 harmonics of a clarinet playing a G#.

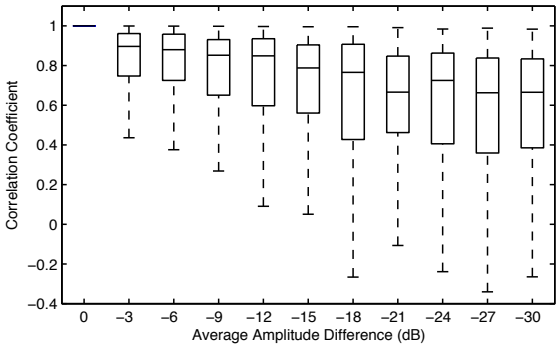

Figure 2. Box plots of correlation coefficient between different harmonics of the same source. Results are calculated using 40 five second instrument performances, with correlation calculated for each note of each performance.

3.1 Common Amplitude Modulation (CAM)

CAM assumes that the amplitude envelopes of sinusoidal components from the same source are correlated. Figure 1 shows the envelopes of the first 20 harmonics of a clarinet tone. We can see that in this case the CAM assumption holds while the spectral smoothness assumption does not. As further support for the CAM assumption, we calculated the correlation coefficient between the strongest harmonic of an individual instrument tone with other harmonics in the same tone as a function of difference in amplitude. The amplitude envelope of each harmonic was calculated by predicting each harmonic's frequency from the ground truth pitch

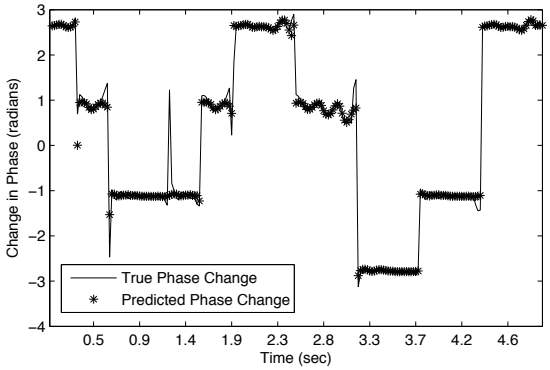

Figure 3. True and predicted phase change for harmonic 1 from a five second excerpt of a flute recording.

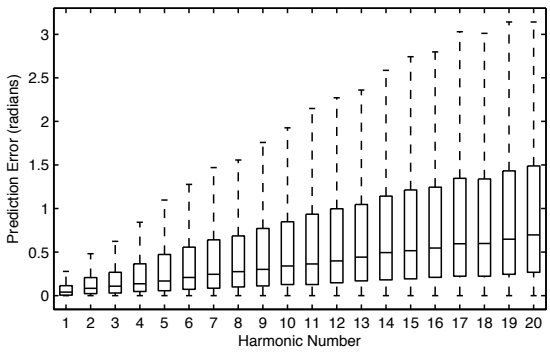

Figure 4. Box plots of phase change prediction error as a function of harmonic number. Results are calculated using 40 five second instrument performances, with error (in radians) calculated as absolute difference between true change in phase from frame m to $m+1$ and predicted change in phase for frame m.

and using the approach described in Section 4 for parameter estimation of non-overlapped harmonics (since every harmonic in a performance by a single instrument is non-overlapped).

Figure 2 shows box plots of the results obtained using 40 five second instrument performances with the correlation calculated for each note of each performance. The upper and lower edges of each box represent the upper and lower quartile ranges, the middle line shows the median value and the whiskers extend to the extent of the sample. For clarity, outliers are excluded from the plot. We can see that the correlation is high for harmonics with energy close to that of the strongest harmonic and tapers off as the energy in the harmonic decreases. This suggests that the amplitude envelope of an overlapped harmonic could be approximated from the amplitude envelopes of non-overlapped harmonics of the same source. Since the low-energy harmonics do not have a strong influence on the perception of a signal, the decreased correlation between the strongest harmonic and lower energy harmonics does not significantly degrade performance.

3.2 Predicting Phase Change using Pitch

According to the sinusoidal model in the time domain (see Equation (1)), the phase of a sinusoid at frame $m+1$ is related to the phase at frame m by

$$\phi_n^{h_n}(m+1) = 2\pi h_n F_n(m) T + \phi_n^{h_n}(m), \qquad (3)$$

where T denotes the frame shift in seconds. Equivalently we can write

$$\Delta\phi_n^{h_n}(m) = 2\pi h_n F_n(m) T. \qquad (4)$$

Therefore the phase change can be predicted from the pitch of a harmonic source. Figure 3 shows the phase change between successive time frames as measured from the first harmonic of a flute recording, and the predicted phase change using the true pitch of the signal. The predicted phase from Equation (4) is wrapped to $[-\pi, \pi]$. This example clearly shows that the phase change of a harmonic component can be accurately predicted from the pitch.

In Figure 4 we show box plots of the error between the true phase change of a harmonic component and the predicted phase change of a harmonic component as a function of harmonic number. The results are taken over the same performances as in Figure 2. As can be seen, for lower-numbered harmonics, the predicted phase change matches well to the true changes.

3.3 Estimating Amplitudes and Phases of Overlapped Harmonics

Using the amplitude envelope and phase change information, we can express the sinusoidal parameter of harmonic h_n, $S_n^{h_n}(m)$, in terms of a reference time frame m_0 as follows:

$$S_n^{h_n}(m) = S_n^{h_n}(m_0) r_{m_0 \to m}^{h_n} e^{i \sum_{l=m_0}^{m} \Delta\phi_n^{h_n}(l)}. \qquad (5)$$

Here, $r_{m_0 \to m}^{h_n} = \frac{a_n^{h_n}(m)}{a_n^{h_n}(m_0)}$ is the amplitude scaling factor between frames m_0 and m for harmonic h_n. The discussion in Section 3.1 suggests that the scaling factor of harmonic h_n can be approximated from the scaling factor of another harmonic of source n, i.e., $r_{m_0 \to m}^{h_n} \approx r_{m_0 \to m}^{h_n^*} = \frac{a_n^{h_n^*}(m)}{a_n^{h_n^*}(m_0)}$, where h_n^* is a non-overlapped harmonic with strong energy. As discussed in Section 3.2, the phase change of harmonic h_n can be predicted using the pitch $F_n(m)$. If we write

$$R_n(m,k) = r_{m_0 \to m}^{h_n^*} e^{i \sum_{l=m_0}^{m} \Delta\phi_n^{h_n}(l)} W(k f_b - h_n F_n(m)), \qquad (6)$$

then Equation (2) becomes

$$X(m,k) = \sum_n R_n(m,k) S_n^{h_n}(m_0). \qquad (7)$$

If harmonics from different sources overlap in a time-frequency (T-F) region with time frames from m_0 to m_1 and frequency bins from k_0 and k_1, we can write Equation (7) for each T-F unit in the region and the set of equations can be represented as

$$\mathbf{X} = \mathbf{RS}, \tag{8}$$

where,

$$\mathbf{X} = \begin{pmatrix} X(m_0, k_0) \\ \vdots \\ X(m_0, k_1) \\ \vdots \\ X(m_1, k_1) \end{pmatrix}, \tag{9}$$

$$\mathbf{R} = \begin{pmatrix} R_1(m_0, k_0) & \dots & R_N(m_0, k_0) \\ \vdots & & \vdots \\ R_1(m_0, k_1) & \dots & R_N(m_0, k_1) \\ \vdots & & \vdots \\ R_1(m_1, k_1) & \dots & R_N(m_1, k_1) \end{pmatrix}, \text{and} \tag{10}$$

$$\mathbf{S} = \begin{pmatrix} S_1^{h_1}(m_0) \\ \vdots \\ S_N^{h_N}(m_0) \end{pmatrix}. \tag{11}$$

The coefficient matrix $\mathbf{R}$ is constructed according to Equation (6) for each T-F unit. $\mathbf{X}$ is a vector of the observed spectral values of the mixture in the overlapping region. We seek a solution for $\mathbf{S}$ to minimize the sum of squared error

$$J = (\mathbf{X} - \mathbf{RS})^{\mathrm{H}}(\mathbf{X} - \mathbf{RS}). \tag{12}$$

The least-squares solution is given by

$$\mathbf{S} = (\mathbf{R}^{\mathrm{H}}\mathbf{R})^{-1}\mathbf{R}^{\mathrm{H}}\mathbf{X}, \tag{13}$$

where H denotes conjugate transpose. After $S_n^{h_n}(m_0)$ is estimated for each of the sources active in the overlapping region, we use Equation (5) to calculate $S_n^{h_n}(m)$ for all $m \in [m_0, m_1]$.

Figure 5 shows the effectiveness of the proposed algorithm in recovering two overlapping harmonics for two instruments. In this case, the third harmonic of the first source overlaps with the fourth harmonic of the second source. Figure 5(c) shows the magnitude spectrum of the mixture in the overlapping region. Note that the amplitude modulation results from the relative phase of the two harmonics. The estimated magnitude spectra of the two harmonics are shown in Figure 5(d) and (e). For comparison, the magnitude spectra of the two sources obtained from pre-mixed signals are shown in Figure 5(a) and (b). It is clear that the estimated magnitude spectra are very close to the true magnitude spectra.

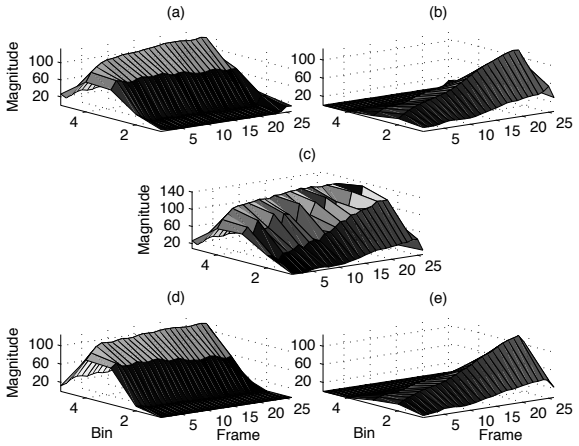

Figure 5. LS estimation of overlapped harmonics. (a) The magnitude spectrum of a harmonic of the first source in the overlapping T-F region. (b) The magnitude spectrum of a harmonic of the second source in the same T-F region. (c) The magnitude spectrum of the mixture at the same T-F region. (d) The estimated magnitude spectrum of the harmonic from the first source. (e) The estimated magnitude spectrum of the harmonic from the second source.

4 A MONAURAL MUSIC SEPARATION SYSTEM

We incorporate the proposed algorithm into a monaural music separation system to evaluate its effectiveness. The diagram of the system is shown in Figure 6. The input to the system is a polyphonic mixture and pitch contours of individual sources. As mentioned previously, we use ground truth pitch estimated from the clean signals for each source. In the harmonic labeling stage, the pitches are used to identify overlapping and non-overlapping harmonics.

To formalize the notion of overlapping harmonics, we say that harmonics h_{n_1} and h_{n_2} for sources n_1 and n_2, respectively, overlap when their frequencies are sufficiently close, $|f_{n_1}^{h_{n_1}}(m) - f_{n_2}^{h_{n_2}}(m)| < \theta_f$. If one assumes the signals strictly adhere to the sinusoidal model, the bandwidth of W determines how many frequency bins will contain energy from a sinusoidal component and one can set an amplitude threshold to determine θ_f.

For non-overlapped harmonics, sinusoidal parameters are estimated by minimizing the sum of squared error between the mixture and the predicted source energy,

$$J = \sum_{k \in K_n^{h_n}(m)} |X(m,k) - W(kf_b - h_n F_n(m))S_n^{h_n}(m)|^2, \tag{14}$$

where $K_n^{h_n}(m)$ is the set of frequency bins associated with harmonic h_n in frame m. The solution is given by:

$$S_n^{h_n}(m) = \frac{\sum_{k \in K_n^{h_n}(m)} X(m,k)W(hF_n(m) - kf_b)}{\sum_{k \in K_n^{h_n}(m)} |W(hF_n(m) - kf_b)|^2}. \tag{15}$$

As described in Section 3.3, we utilize the amplitude envelope of non-overlapped harmonics to resolve overlapping harmonics. Since the envelope information is sequential,

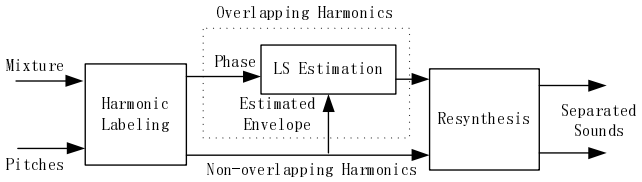

Figure 6. System diagram

we resolve overlapped h_n for time frames $[m_0, m_1]$ using a non-overlapped harmonic h_n^*. To determine appropriate time frames for this processing, we first identify sequences of time frames for which a harmonic h_n is overlapped with one or more other harmonics. If the pitch of any of the sources contributing to the overlapping region changes, we break the sequence of frames into subsequences. Given a sequence of frames, we choose the strongest harmonic for each source that is unobstructed in the entire sequence as h_n^*. We use θ_f to determine the bin indices, $[k_0, k_1]$, of the overlapping region.

For each overlapping region, we perform least-squares estimation to recover the sinusoidal parameters for each instrument's harmonics. To utilize the mixture signal as much as possible, we estimate the source spectra differently for the overlapped and non-overlapped harmonics. For all non-overlapped harmonics, we directly distribute the mixture energy to the source estimate,

$$\hat{Y}_n^{no}(m,k) = X(m,k) \quad \forall k \in K_n^{h_n}(m). \tag{16}$$

For the overlapped harmonics, we utilize the sinusoidal model and calculate the spectrogram using

$$\hat{Y}_n^{o}(m,k) = S_n^{h_n}(m) W(kf_b - f_n^{h_n}(m)). \tag{17}$$

Finally, the overall source spectrogram is $\hat{Y}_n = \hat{Y}_n^{no} + \hat{Y}_n^{o}$ and we use the overlap-add technique to obtain the time-domain estimate, $\hat{y}_n(t)$, for each source.

5 EVALUATION

5.1 Database

To evaluate the proposed system, we constructed a database of 20 quartet pieces by J. S. Bach. Since it is difficult to obtain multi-track recordings, we synthesize audio signals from MIDI files using samples of individual notes from the RWC music instrument database [15]. For each line selected from the MIDI file, we randomly assign one of four instruments: clarinet, flute, violin or trumpet. For each note in the line, a sample with the closest average pitch is selected from the database for the chosen instrument. We create two source mixtures (using the alto and tenor lines from the MIDI file) and three source mixtures (soprano, alto and tenor), and select the first 5-seconds of each piece for evaluation. All lines are mixed to have equal level, thus lines in the two instrument mixtures have 0 dB SNR and those in the three instrument mixtures have roughly -3 dB SNR. Details about the synthesis procedure can be found in [16]. Admittedly, audio signals generated in this way are a rough approximation of real recordings, but they show realistic spectral and temporal variations.

	SNR improvement
Virtanen (2 sources, 2006)	11.1 dB
Proposed System (2 sources)	14.5 dB
Proposed System (3 sources)	14.7 dB

Table 1. SNR improvement

5.2 Results

For evaluation we use the signal-to-noise ratio (SNR),

$$\text{SNR} = 10 \log_{10} \frac{\sum_t y^2(t)}{\sum_t (\hat{y}(t) - y(t))^2}, \tag{18}$$

where $y(t)$ and $\hat{y}(t)$ are the clean and the estimated instrument signals, respectively. We calculate the SNR gain after separation to show the effectiveness of the proposed algorithm. In our implementation, we use a frame length of 4096 samples with sampling frequency 44.1 kHz. No zero-padding is used in the DFT. The frame shift is 1024 samples. We choose $\theta_f = 1.5 f_b$, one and half times the frequency resolution of the DFT. The number of harmonics for each source, H_n, is chosen such that $f_n^{H_n}(m) < \frac{f_s}{2}$ for all time frames m, where f_s denotes the sampling frequency.

Performance results are shown in Table 1. The first row of the table is the SNR gain for the two source mixtures achieved by the Virtanen system [1], which is also based on sinusoidal modeling. At each frame, this approach uses pitch information and the least-squares objective to simultaneously estimate the amplitudes and phases of the harmonics of all instruments. A so-called adaptive frequency-band model is used to estimate the parameters of overlapped harmonics. To avoid inaccurate implementation of this system, we asked the author to provide separated signals for our set of test mixtures. The second row in Table 1 shows the SNR gain achieved by our system. On average, our approach achieved a 14.5 dB SNR improvement, 3.4 dB higher than the Virtanen system. The third row shows the SNR gain of our system on the three source mixtures. Note that all results were obtained using ground truth pitches. Sound demos of the our separation system can be found at: www.cse.ohio-state.edu/∼woodrufj/mmss.html

6 DISCUSSION AND CONCLUSION

In this paper we have proposed an algorithm for resolving overlapping harmonics based on CAM and phase change estimation from pitches. We incorporate the algorithm in a separation system and quantitative results show significant improvement in terms of SNR gain relative to an existing

monaural music separation system. In addition to large increases in SNR, the perceptual quality of the separated signals is quite good in most cases. Because reconstruction of overlapped harmonics is accurate and we utilize the mixture for non-overlapped harmonics, the proposed system does not alter instrument timbre in the way that synthesis with a bank of sinusoids can. A weakness of the proposed approach is the introduction of so-called *musical noise* as performance degrades. One aspect of future work will be to address this issue and create higher quality output signals.

In this study we assume that the pitches of sources are known. However, for practical applications, the true pitches of sources in a mixture are not available and must be estimated. Since our model uses pitch to identify overlapped and non-overlapped harmonics and pitch inaccuracy affects both the least-squares estimation and phase change prediction, good performance is reliant on accurate pitch estimation. We are currently investigating methods that relax the need for accurate prior knowledge of pitch information. Preliminary results suggest that performance similar to the Virtanen system using ground truth pitch can still be achieved by our approach even with prior knowledge of only the number of sources (when combining our system with multi-pitch detection) or rough pitch information (as provided by MIDI data).

Acknowledgment

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FAST MIR IN A SPARSE TRANSFORM DOMAIN

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ABSTRACT

We consider in this paper sparse audio coding as an alternative to transform audio coding for efficient MIR in the transform domain. We use an existing audio coder based on a sparse representation in a union of MDCT bases, and propose a fast algorithm to compute mid-level representations for beat tracking and chord recognition, respectively an onset detection function and a chromagram. The resulting transform domain system is significantly faster than a comparable state-of-the-art system while obtaining close performance above 8 kbps.

1 INTRODUCTION

Music recordings are now widely available in coded format. The reason is that state-of-the-art audio coders such as MP3 [6] or AAC [7] are able to reduce the size of a PCM audio signal more than 10 times, while guaranteeing a near-transparent quality. Consequently, such technology allows users to easily exchange and store music on mobile devices and networks.

On the other hand, state-of-the-art audio indexing algorithms such as beat tracking [8, 2] and chord recognition [1, 10] are designed to process PCM audio signals. Consequently, to use them with coded audio, one has to decode to PCM first and then apply the audio indexing algorithm on the PCM signal (*Processing in the time domain*, see Fig. 1). To save computational cost, which is often required e.g. when using such algorithms with mobile devices or on very large databases, it would be more efficient to design audio indexing algorithms that work directly with the coded data.

There are two ways to process coded data depending on which stage of the decoding process we are working on (see Fig. 1). The first way is to use directly the bitstream, this approach is called *processing in the compressed domain*. The second way is to use the transform representation, this approach is called *processing in the transform domain*. The first approach is faster as we avoid the cost of decoding the transform representation, however, for certain cases the information available in the bitstream is not sufficiently explicit, and it it thus necessary to use the transform domain representation.

We consider in this paper two audio indexing applications, beat tracking [8, 2] and chord recognition [1, 10].

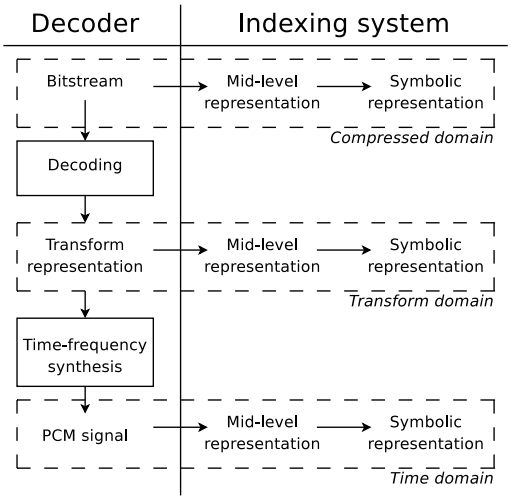

Figure 1. *Block diagram of a common audio decoder and three possible audio indexing systems.*

Beat tracking has already been investigated using MP3 audio files in the transform domain [13] and in the compressed domain [14]. However, no work related to chord recognition using coded data has been found in the literature. This may be due to the limited frequency resolution of the time-frequency analysis used in state-of-the-art transform audio coders such as MP3 [6] and AAC [7].

Due to this limitation of the transform based coders, it is interesting to consider other kinds of audio coders, such as parametric coding (e.g. [3]) or sparse representation based coding (e.g. [12]). We have chosen here to use a new prototype audio coder based on a union of MDCT bases [12], which has the advantage to provide a sparse representation which has both precise time and frequency resolution, contrary to transform based coders. This coder, available for testing in open source [1], has also the interesting property of fine-grain scalability.

We show in this paper that this new signal representation approach is not only useful for audio coding, but also for audio indexing. We propose a fast method to calculate, in the transform domain, mid-level representations similar to those used in state-of-the-art systems, namely an onset

[1] `http://www.emmanuel-ravelli.com/downloads.html`

detection function and a chromagram. The onset detection function and the chromagram are then used to perform respectively beat tracking and chord recognition using same machine learning systems as used in state-of-the-art systems [2, 1].

The remainder of this paper is as follows. In section 2, we describe the signal representation used in [12]. In section 3, the details of the calculation of the mid-level representations are given. In section 4, the machine learning systems we use to perform beat tracking and chord recognition are briefly described. Finally, section 5 gives the performance evaluation, section 6 discuss about computation times, and we conclude in section 7.

2 SIGNAL REPRESENTATION IN A UNION OF MDCT BASES

In state-of-the-art audio coders such as AAC [7], the Modified Discrete Cosine Transform (MDCT) is used. While such coders allow transparent quality at high bitrates, they give limited performance at low bitrates. In [12], Ravelli et al proposed a generalization of the transform coding approach where the signal is decomposed in a union of MDCT bases with different scales. The results showed that this new approach allows improved performance at low bitrates. We briefly present in the following the signal model and the decomposition algorithm.

2.1 Signal model

The signal is modeled using a union of 8 MDCT bases, where the window length ranges from 128 to 16384 samples (i.e. from 2.9 to 370 ms) in powers of 2. The smallest windows are needed to model very sharp attacks while larger windows are useful for modeling long stationary components. The signal $f \in \mathbb{R}^N$ is then decomposed as a weighted sum of functions $g_\gamma \in \mathbb{R}^N$ plus a residual of negligible energy r

$$f = \sum_{\gamma \in \Gamma} \alpha_\gamma g_\gamma + r \tag{1}$$

where α_γ are the weighting coefficients. The set of functions $\mathcal{D} = \{g_\gamma, \gamma \in \Gamma\}$ is called the dictionary and is a union of M MDCT bases (called blocks). The functions g, called atoms are defined as:

$$g_{m,p,k}(n) = w_m(u). \sqrt{\frac{2}{L_m}} \cos\left[\frac{\pi}{L_m}\left(u + \frac{1+L_m}{2}\right)\left(k + \frac{1}{2}\right)\right] \tag{2}$$

where $u = n - pL_m - T_m$ and m is the block index, p is the frame index, k is the frequency index, L_m is the half of the analysis window length of block m (defined as power of two $L_m = L_0 2^m$), P_m is the number of frames of block m, T_m is a time offset introduced to "align" the windows of different lengths ($T_m = \frac{L_m}{2}$) and $w_m(u)$ is the sine window defined on $u = 0, .., 2L_m - 1$.

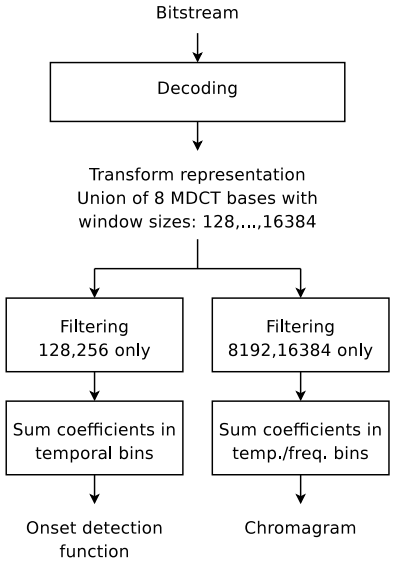

Figure 2. *Block diagram of the proposed system for the calculation of the mid-level representations.*

2.2 Decomposition algorithm

The signal is decomposed using Matching Pursuit (MP). MP [11] is an iterative algorithm which select at each iteration the atom in the dictionary that is the most correlated with the residual; subtracts the selected atom from the residual; and iterates until a stopping condition (e.g. target SNR) is met. The decomposition algorithm has been implemented in the Matching Pursuit ToolKit (MPTK) framework [9], which is to date the fastest available generic implementation of Matching Pursuit for audio signals.

3 CALCULATION OF MID-LEVEL REPRESENTATIONS

The signal representation used in the audio coder of [12] is based on a union of 8 MDCT bases with analysis window sizes from 128 to 16384 samples. We have remarked that high amplitude atoms with small window sizes (128 and 256) are often located around attacks; consequently, we can build a very cheap onset detection function by filtering the decomposition such that we keep only small window sizes atoms, and then sum the absolute value of the coefficients in temporal bins to construct a downsampled signal with peaks located at attacks. We have also remarked that strong tonal components are modeled by the large window sizes atoms (8192 and 16384) with a precise frequency resolution. Consequently, it is possible to build a very cheap chromagram by

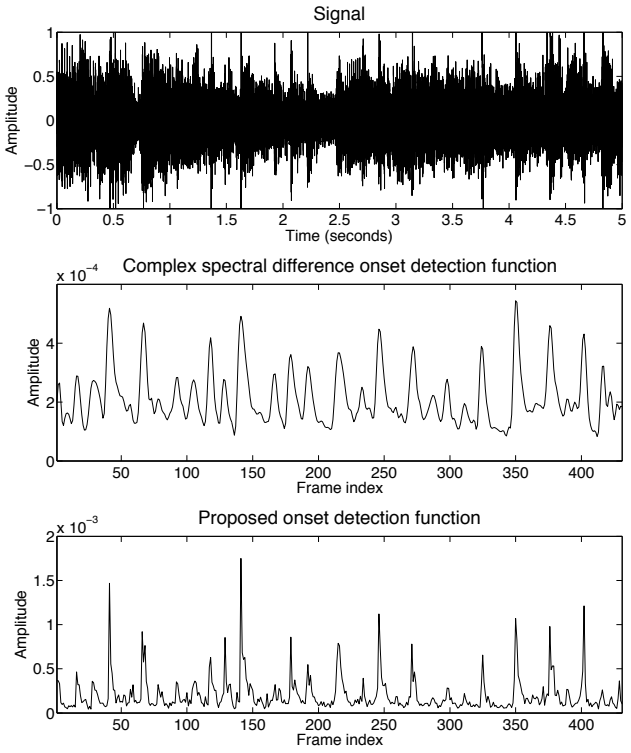

Figure 3. *A 5 seconds signal of rock music; the complex spectral difference onset detection function of [2]; the proposed onset detection function.*

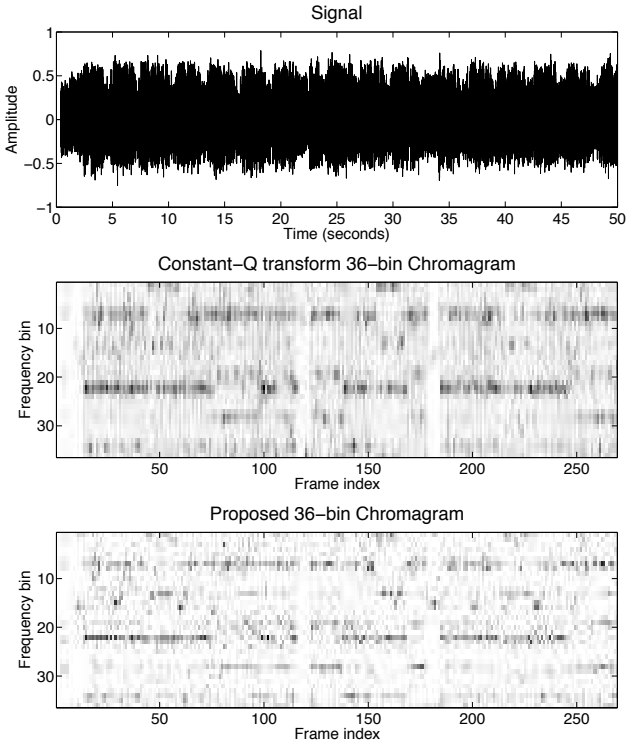

Figure 4. *A 50 seconds signal of rock music; the constant-Q transform 36-bin chromagram of [1]; the proposed 36-bin chromagram*

summing the absolute value of the coefficients of the largest atoms in time/frequency bins. The complete system is in Fig. 2 and we describe it in more details in the following.

3.1 Onset detection function

The onset detection function Γ is computed on a frame-by-frame basis. The length of one frame is defined such that the corresponding time resolution is the same as in [2] and [8] which is 11.6 ms and it is equivalent to $t_{DF} = 512$ samples at 44.1 kHz. The function $\Gamma(q)$ at frame q is thus defined as

$$\Gamma(q) = \sum_{m,p,k} |\alpha_{m,p,k}| \tag{3}$$

where we sum only the atoms satisfying the following two conditions:

- the window size is 128 or 256 samples

$$m < 2 \tag{4}$$

- the center is in the temporal support of the frame q

$$floor\left(\frac{(p+1)L_m + T_m}{t_{DF}}\right) = q. \tag{5}$$

Fig. 3 shows the onset detection function obtained with a 5 seconds signal of rock music, and as a reference the onset detection function of [2].

3.2 Chromagram

The chromagram is computed on a frame-by-frame basis too. In [1], the time resolution is 92.9 ms, while in [10] the time resolution is the double 185.8 ms. We decide to use here a time resolution of 185.8 ms as this is the hop size of the MDCT with largest window size. This is equivalent to a frame size of $t_{CH} = 8192$ samples at 44.1 kHz. The Chromagram $CH(q, b)$ at frame q and frequency bin b is thus defined as

$$CH(q, b) = \sum_{m,p,k} |\alpha_{m,p,k}| \tag{6}$$

where we sum only the atoms satisfying the following three conditions:

- the window size is 8192 or 16384 samples

$$m \geq 6 \tag{7}$$

- the center is in the temporal support of the frame q

$$floor\left(\frac{(p+1)L_m + T_m}{t_{CH}}\right) = q. \quad (8)$$

- the frequency value k maps to the frequency bin b of the chromagram

$$\text{mod}\left(\text{round}\left(B\,\text{log2}\left(\frac{22050\text{ k}/L_m}{f_{min}}\right)\right), B\right) = b \quad (9)$$

with $B = 36$ the number of bins per octave and f_{min} is the minimum frequency.

Fig. 4 shows the chromagram obtained with a 50 seconds signal of rock music, and as a reference the chromagram of [1].

4 MACHINE LEARNING SYSTEMS

After calculation of the mid-level representation, it is passed into a machine learning system to produce a symbolic representation. The onset detection function produces a sequence of beats (frame index of the detected beats) and the chromagram produces a sequence of chords (one detected chord per frame). We use the same machine learning as in other works in order to compare only the mid-level representations in the evaluation of the final system. We describe briefly in the following the machine learning systems.

4.1 Beat tracking

The same system as in [2] is used. The onset detection function is first post-processed using an adaptive moving average threshold. Then the onset detection function is partitioned into overlapping frames to allow variable tempo. In each frame, the unbiased autocorrelation function of the onset detection function is calculated. The autocorrelation function is then passed into a shift-invariant context-dependant comb filterbank in order to estimate the tempo of the current frame. Finally, a beat train at the estimated tempo is built and aligned with the current frame by passing the detection function into a tuned context-dependant comb filterbank.

4.2 Chord recognition

The same system as in [1] is used: the 36-bin chromagram is first circularly shifted according to the estimated tuning of the piece, low-pass filtered, and mapped to a 12-bin chromagram by simply summing within semitones. Then, the Expectation Maximization (EM) algorithm is used to train the initial states probabilities and the transition matrix of an Hidden Markov Model (HMM). Finally, the sequence of chords is estimated using the Viterbi algorithm with the chromagram and the trained HMM.

5 EVALUATION

We evaluate in the following the performance of our proposed system, in comparison with the reference systems [2, 1]. As the coding/decoding process is lossy, we also study the influence of the codec bitrate on the system performance. The audio files are coded with the coder described in [12], using the standard matching pursuit with a target SNR of 60 dB, and the bitplane encoder with a bitrate of 128 kbps. As we use an embedded coding method, the decoder simply truncates the bitstream to obtain lower bitrates.

5.1 Beat tracking

We compare the performance of our system with the system of Davies et al [2] on the same database provided by S. Hainsworth [4]. There are 222 files of several music genres. As we use the same beat tracking system, this evaluation compares only the detection function used in the system. Davies *et al.* use a complex spectral difference onset detection function, which is sensitive not only for percussive sounds but also for soft tonal onsets (contrary to our detection function). Results are in Fig. 5. We give four measures of beat accuracy, as proposed in [8] and used also in [2]: correct metrical level with continuity required (CML cont); correct metrical level with continuity not required (CML total); allowed metrical levels with continuity required (AML cont); allowed metrical levels with continuity not required (AML total). The performance of our system is close to the performance of the reference system at high bitrates (5-12% less) and even at 8 kbps, with a very bad quality of the synthesized audio, the performance is still high.

5.2 Chord recognition

We compare the performance of our system with the system of Bello et al [2]. We use the same evaluation database as in [1]. It consists of 2 albums of the Beatles: Please Please Me (14 songs) and Beatles for Sale (14 songs). The database has been annotated by C. Harte et al [5]. Only the chromagram calculation is different, the machine learning system is exactly the same. To calculate the 36-bin chromagram, Bello et al use a constant-Q transform on a downsampled PCM audio signal at 11.25 kHz, with a analysis window length of 8192 samples (32768 at 44.1 kHz, twice as ours). To obtain a relevant comparison, we have modified the time resolution of the Bello system such that it is the same as our system: 185.8 ms. Recognition rates (percentage of well detected frames) for both Beatles CD are in Fig. 6. The performance of our system is close to the performance of the reference system at high bitrates (5-7% less) and even at 4 kbps, with a very bad quality of the synthesized audio, the performance is still high.

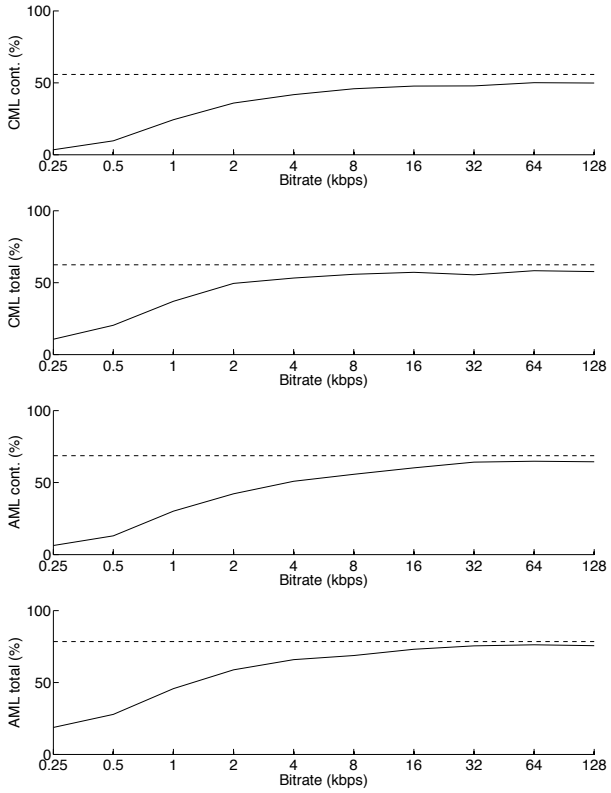

Figure 5. *Performance of the proposed beat tracking system in function of the decoding bitrate. The dotted line corresponds to the performance of the reference system (with the complex spectral difference onset detection function) on the original PCM audio.*

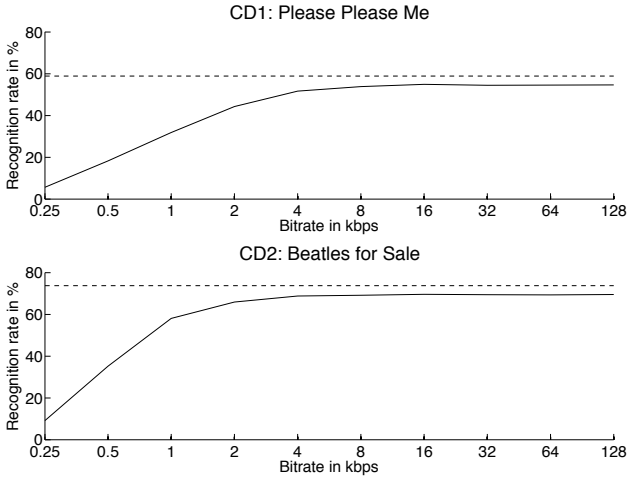

Figure 6. *Performance of the proposed chord recognition system in function of the decoding bitrate. The dotted line corresponds to the performance of the reference system (with the constant-Q transform chromagram) on the original PCM audio.*

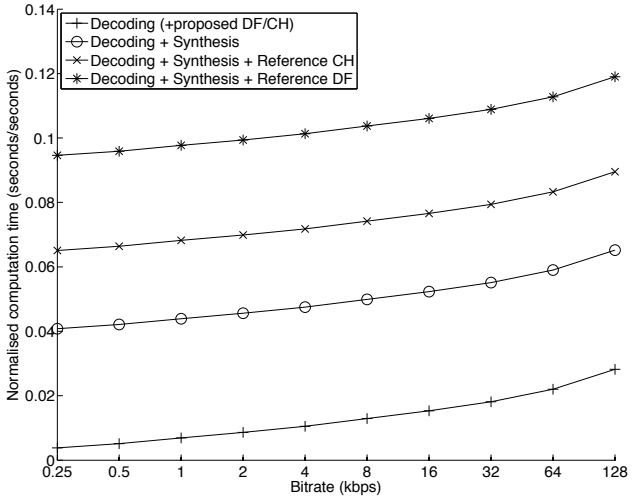

Figure 7. *Computation times of the different stages: bitstream decoding; signal synthesis; reference mid-level representations. The computation times for the proposed mid-level representations are not given as they are negligible in comparison with the bitstream decoding*

6 COMPUTATION TIME

As stated in the previous section, the performance of our transform domain systems is slightly lower than the one of the reference systems. However, working in the transform domain allows a huge gain in computation times.

As we are working with coded audio, there are two possibilities for a user to calculate mid-level representations from the coded audio. The first possibility is to decode the bitstream, synthesize the decoded transform representation, and calculate state-of-the-art mid-level representations on the synthesized PCM audio. This is the best approach in terms of performance. However, it requires several operations with a non negligible computational cost: the bitstream decoding; the transform representation synthesis, which requires 8 IMDCT; the mid-level representation, which requires a time-frequency analysis (FFT) plus additional computations. The second possibility is to compute mid-level representations from the transform representation as explained in the previous sections. This approach is a bit less efficient in terms of performance but it requires less operations that the first approach and thus is faster. Only two operations are required: the bitstream decoding; and the mid-level representation calculation, which involves only histogram-like calculations and has thus small computation cost as compared to the bitstream decoding cost.

Fig. 7 shows the computation times of the different operations: the bitstream decoding; the synthesis; the reference chromagram from PCM audio; and the reference onset detection function from PCM audio. The proposed mid-level

representations cost is not given as it is very small in comparison with the bitstream decoding. Results show that the computation of the proposed transform-domain mid-level representations is from 3 times (CH at high bitrates) to 24 times (DF at low bitrate) faster than the computation of the reference mid-level representations.

7 CONCLUSION

We have considered in this paper an alternative approach for audio indexing in the transform domain. While common approach use standard transform based coders, we show that using sparse representation allows to go beyond the limitations of the time-frequency resolution of the transform approach. We have chosen in this paper the audio coder of [12], which is based on a union of MDCT bases and we show that MIR in the sparse transform domain allows user to trade performance and computational complexity.

We have considered in this paper two applications, beat tracking and chord recognition. And we have proposed the calculation of two mid-level representations for these applications; an onset detection function and a chromagram. We have showed that the performance of the resulting systems are close to the performance of the reference systems above 8kbps. We have also shown that the computation time of the proposed systems depend mainly on the bitstream decoding module, as the cost of the proposed mid-level representations is small in comparison. Moreover, the saving of the synthesis and the time-frequency analysis allows a gain in the computation time from 3 to 24 times.

Given the results of this work, it would be interesting to consider in the future other audio indexing applications, such as e.g. musical genre classification, by using equivalent low level features such as e.g. MFCC. It would be also interesting to try other signal representations, such as MDCT with greater window sizes, or union of MCLT, and study the influence of such representations on the audio coding and audio indexing systems.

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PLAYLIST GENERATION USING START AND END SONGS

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ABSTRACT

A new algorithm for automatic generation of playlists with an inherent sequential order is presented. Based on a start and end song it creates a smooth transition allowing users to discover new songs in a music collection. The approach is based on audio similarity and does not require any kind of meta data. It is evaluated using both objective genre labels and subjective listening tests. Our approach allows users of the website of a public radio station to create their own digital "mixtapes" online.

1 INTRODUCTION

This work is concerned with the creation of playlists with an inherent sequential order. Such a playlist consists of a start and an end song, both chosen by a user. The songs in between should form a smooth transition, with songs at the beginning sounding similar to the start song, songs at the end similar to the end song and songs in the middle similar to both start and end songs. Our approach is based solely on audio analysis and does not require any kind of metadata. It could therefore easily replace or at least support manual creation of playlists. It allows to explore audio collections by simply choosing two songs and a desired length for the playlist. It also enables efficient discovery of new music if applied to collections of yet unknown songs by automatically creating a smooth transition between only two supporting songs.

Most existing approaches to playlist generation rely on the usage of one seed song or a group of seed songs. The playlist then consists of songs which are somehow similar to this seed. Some authors use different kinds of audio similarity to create the playlists [7, 10]. Others work with some kind of metadata [11, 14, 16]. Seed based creation of playlists has the problem of producing too uniform lists of songs if applied to large data bases with lots of similar music. If a data base does not contain enough similar music to a seed song there is the danger of "playlist drift" towards music that sounds very different.

Few authors report about generating playlists with an inherent sequential order. Most approaches are solely based on metadata and not audio similarity. Case Based Reasoning has been applied to create new playlists with inherent temporal structure based on patterns of subsequences of a collection of existing playlists [4]. Creation of playlists satisfying user constraints based on rich metadata has also been reported [3]. These constraints may also concern the temporal order of the playlists (e.g. rising tempo, change of genre). This constraint based approach has been extended [2] to include notions of audio similarity as yet another constraint (e.g. timbre continuity through a playlist). Related approaches have been formulated based on simulated annealing [12] and linear programming [1]. Travelling Salesman algorithms applied to audio similarity have been used to generate a sequential order of all songs in a data base [15]. Since all songs have to be part of the playlist, this is a quite different kind of organising principle for the playlist. Direct interaction with a two dimensional mapping of music spaces based on audio similarity also allows creation of playlists with inherent sequential structure [9]. Computations are based on the lower dimensional representations and not directly on the audio models of the songs themselves. A related approach also using a two dimensional display which is enriched with various kinds of meta data has also been presented [5].

Contrary to the work reviewed above, our approach is (i) based on audio similarity, (ii) requires very little user interaction and (iii) results in playlists with smooth temporal transitions potentially including songs previously unknown to the users. Our playlist generation algorithm has been developed for the internet portal of an Austrian radio station to allow creation of digital "mixtapes" online.

2 DATA

This work is part of a project aiming at providing novel ways of accessing the music of an Austrian music portal. The FM4 Soundpark is an internet platform [1] of the Aus-

[1] http://fm4.orf.at/soundpark

	HiHo	Regg	Funk	Elec	Pop	Rock
No.	226	60	56	918	158	1148
%	9	2	2	36	6	45

Table 1. Number of songs and percentages across genres in our data base. Genres are Hip Hop, Reggae, Funk, Electronic, Pop and Rock.

trian public radio station FM4. This internet platform allows artists to present their music free of any cost in the WWW. All interested parties can download this music free of any charge. At the moment this music collection contains about 10000 songs but it is only organised alphabetically and in a coarse genre taxonomy. The artists themselves choose which of the six genre labels "Hip Hop, Reggae, Funk, Electronic, Pop and Rock" best describe their music. We use a development data base of 2566 songs for our experiments. Number of songs and percentages across genres are given in Tab. 1. The distribution of genres is quite unbalanced with "Electronic" and "Rock" together taking up 81%. This is representative of the full data base.

From the 22050Hz mono audio signals two minutes from the center of each song are used for further analysis. We divide the raw audio data into non-overlapping frames of short duration and use Mel Frequency Cepstrum Coefficients (MFCC) to represent the spectrum of each frame. MFCCs are a perceptually meaningful and spectrally smoothed representation of audio signals. MFCCs are now a standard technique for computation of spectral similarity in music analysis (see e.g. [6]). The frame size for computation of MFCCs for our experiments was $46.4ms$ (1024 samples). We used the first 20 MFCCs for all our experiments.

3 METHODS

Our playlist generation algorithm consists of two basic parts: (i) computation of similarities between songs, (ii) computation of the actual playlists based on these similarities. Please note that the actual generation of playlists does not rely on a specific similarity function and could therefore also be done using different approaches towards computation of similarity.

3.1 Computing spectral similarity of songs

We use the following approach to music similarity based on spectral similarity. For a given music collection of songs, it consists of the following steps:

1. for each song, compute MFCCs for short overlapping frames as described in Sec. 2
2. train a single Gaussian (G1) to model each of the songs
3. compute a similarity matrix between all songs using the Kullback-Leibler divergence between respective G1 models

We use one single Gaussian (G1) with full covariance to represent the MFCCs of each song [8]. For single Gaussians, $p(x) = \mathcal{N}(x; \mu_p, \sigma_p)$ and $q(x) = \mathcal{N}(x; \mu_q, \sigma_q)$, there is a closed form of the Kullback-Leibler divergence [13]:

$$KL_N(p\|q) = 0.5 \log\left(\frac{\det(\Sigma_p)}{\det(\Sigma_q)}\right) + 0.5Tr\left(\Sigma_p^{-1}\Sigma_q\right) + 0.5\left(\mu_p - \mu_q\right)'\Sigma_p^{-1}\left(\mu_q - \mu_p\right) - \frac{d}{2} \quad (1)$$

where $Tr(M)$ denotes the trace of the matrix M, $Tr(M) = \Sigma_{i=1..n} m_{i,i}$. Dropping constants and symmetrizing the divergence yields the following approximation [17]:

$$D_{KL}(p,q) = Tr\left(\Sigma_p^{-1}\Sigma_q\right) + Tr\left(\Sigma_q^{-1}\Sigma_p\right) + Tr\left(\left(\Sigma_p^{-1} + \Sigma_q^{-1}\right)\left(\mu_p - \mu_q\right)\left(\mu_q - \mu_p\right)'\right) \quad (2)$$

Please note that this approximation is symmetric, i.e. $D_{KL}(p,q) = D_{KL}(q,p)$, and that the self-similarity is non-zero, i.e. $D_{KL}(p,p) \neq 0$. Actually, $D_{KL}(p,p) = 2d$ with d being the dimensionality of the data vectors (20 MFCCs in our case).

3.2 Computing playlists

Our algorithm for computation of a playlist of length p (excluding start and end song) for a database of n songs S_i, starting at song S_s and ending at song S_e consists of the following steps:

1. for all $i = 1, ..., n$ songs compute the divergences to the start song $D_{KL}(i, s)$ and the end song $D_{KL}(i, e)$
2. find the $d\%$ songs with greatest divergence $D_{KL}(i, s)$ to the start song S_s; find the $d\%$ songs with greatest divergence $D_{KL}(i, e)$ to the end song S_e; discard all songs which are in both of these groups; keep remaining m songs for further processing
3. for all $i = 1, ..., m$ songs compute a divergence ratio:

$$R(i) = \frac{D_{KL}(i, s)}{D_{KL}(i, e)} \quad (3)$$

4. compute step width for playlist:

$$step = \frac{R(s) - R(e)}{p + 1} \quad (4)$$

5. compute p ideal positions (i.e. ideal divergence ratios) $\hat{R}(j), j = 1, ..., p$:

$$\hat{R}(j) = R(s) + j * step \quad (5)$$

6. select the p real songs S_j that best match the ideal divergence ratios $\hat{R}(j), j = 1, ..., p$:

$$S_j = \arg \min_{i=1,...,m} |\hat{R}(j) - R(i)| \quad (6)$$

The main part of our algorithm is the computation of divergence ratios $R(i)$. Songs which are closer to the start song S_s than to the end song S_e will have a divergence ratio $R(i) < 1$. Songs which are closer to the end song S_e than to the start song S_s will have a divergence ratio $R(i) > 1$. Songs which have about the same divergence to both songs will have a divergence ratio $R(i)$ around 1. Songs which have a big divergence to both start and end song will therefore also have a divergence ratio $R(i)$ around 1 and therefore might end up as part of the middle section of a playlist. This is of course not as desired since only songs which are close to either or both the start and end song should be part of the playlist. Songs too distant from both start and end song appear as outliers to the listeners. Therefore we discard songs which are distant to both start and end song during step 2 of the above algorithm. The amount of songs we discard is controlled with the parameter d. In initial experiments we found out that $d = 95\%$ works well for this data set.

The playlist is then computed in the divergence ratio space: $R(s)$ serves as the starting position and $R(e)$ as the end position of the list. The aim is to find p songs which are at equally spaced positions between these start and end positions. This is done by computing a step width in step 4 of the algorithm, computing ideal positions for the playlist songs in the divergence ratio space in step 5 and finally finding songs that best match these ideal positions in step 6.

4 RESULTS

4.1 Objective evaluation

One possibility to achieve an objective evaluation is to use the genre labels as indicators of music similarity. For a playlist with start song belonging to genre A and end song belonging to genre B we formulate the following hypotheses:

- the playlist should contain mostly songs from genres A and B
- at the beginning of the playlist, most songs should be from genre A, at the end from genre B and from both genres in the middle

		nearest neighbour classification					
		HiHo	Regg	Funk	Elec	Pop	Rock
t	HiHo	**73**	8	0	11	4	4
r	Regg	35	**33**	2	13	5	12
u	Funk	20	5	**29**	16	13	18
e	Elec	13	7	3	**56**	8	13
	Pop	22	3	4	13	**26**	32
	Rock	4	1	1	3	4	**87**

Table 2. Confusion matrix of genre classification results (nearest neighbour classification vs. true genre label). Results are given in percentages separately per genre in each row. Genres are Hip Hop, Reggae, Funk, Electronic, Pop and Rock.

The success of such an approach depends strongly on how well the genre labels actually indicate music similarity. This can be measured by looking at the genre classification results. Table 2 gives a confusion matrix for a 10-fold cross-validation experiment with one-nearest neighbour classification using the divergences D_{KL}. Results are given in percentages separately per genre in each row. Some of the genres can be classified very well (Hip Hop: 73%, Rock: 87%), others somewhat well (Electronic: 56%) and some quite badly (Reggae, Funk and Pop are all around 30%). Consequently, any playlist evaluation relying on the genre information should do quite well on genres Hip Hop, Rock and maybe Electronic. But it would show the same confusion of labels for all other genres.

We randomly chose 50 songs from each of the six genres as candidates for start and end songs. Since our playlist algorithm gives identical members of playlists in reversed order when start and end songs are exchanged, we need to look at only $(6 \times (6 - 1))/2 = 15$ possible combinations of our six genres. For each combination of two genres A and B, we compute all possible 50×50 playlists using the candidate songs as start and end songs. This yields 37500 playlists altogether. The length of each playlist is nine songs excluding start and end songs. We divide all playlists in three sections (first, middle and last three songs) and report distribution of songs across genres in the playlists. Instead of showing results for all possible 15 combinations of genres we concentrate on a number of examples showing the range of quality one can expect.

Table 3 shows the results for playlists starting at Hip Hop and ending at Rock. Both genres dominate (33% and 38%) the beginning of the playlists (Sec1). Whereas Hip Hop quickly diminishes to 5% and 2%, Rock rises to 81% and 88% at the end.

The results for playlists starting at Hip Hop and ending at Electronic (Tab. 4) as well as for playlists starting at Electronic and ending at Rock (Tab. 5) work equally well. The respective genres dominate the beginning of the playlists.

	HiHo	Regg	Funk	Elec	Pop	**Rock**
Sec1	33	5	2	15	8	38
Sec2	5	1	2	7	4	81
Sec3	2	0	3	4	2	88

Table 3. Distribution of songs across genres in playlists starting at Hip Hop and ending at Rock. Results given for first, middle and last section of playlists (Sec1 to Sec3).

	HiHo	Regg	Funk	**Elec**	Pop	Rock
Sec1	30	5	2	35	8	19
Sec2	6	2	3	66	5	18
Sec3	2	2	3	70	4	18

Table 4. Distribution of songs across genres in playlists starting at Hip Hop and ending at Electronic. Results given for first, middle and last section of playlists (Sec1 to Sec3).

The start genres diminish quickly and the end genres are most prominent in the last sections (Sec3). Tables 3 to 5 give results for the three genres which also achieve the best classification results (see Tab. 2). The results are basically in line with the two hypotheses we formulated at the beginning of this section. Only the fact that the end genre is already very prominent at the beginning of the playlists (Sec1) is a bit surprising. This might be due to the fact that the end genres in Tables 3 to 5 are also the most numerous in our data base (Electronic 36% and Rock 45% of all songs in the data base, see Tab. 1).

Table 6 shows the results for playlists starting at Reggae and ending at Rock. The amount of songs from genre Rock rises from 38% to 80% to 88% going from Sec1 to Sec3 as expected. Genre Reggae is somewhat under-represented in all sections of the playlists. Going back to the genre classification confusion matrix in Tab. 2, it is clear that there is a lot of mix-up between genres Reggae and Hip Hop. Consequently, Tab. 6 shows a considerable amount of Hip Hop in Sec1, diminishing towards Sec3.

The results for playlists starting at Funk and ending at Pop given in Tab. 7 are even less satisfactory. The genre classification confusion matrix in Tab. 2 shows that genre

	HiHo	Regg	Funk	**Elec**	Pop	**Rock**
Sec1	13	3	2	42	6	34
Sec2	8	2	2	8	5	77
Sec3	5	1	3	3	3	85

Table 5. Distribution of songs across genres in playlists starting at Electronic and ending at Rock. Results given for first, middle and last section of playlists (Sec1 to Sec3).

	HiHo	**Regg**	Funk	Elec	Pop	**Rock**
Sec1	26	7	2	20	7	38
Sec2	6	1	2	7	4	80
Sec3	3	0	2	4	2	88

Table 6. Distribution of songs across genres in playlists starting at Reggae and ending at Rock. Results given for first, middle and last section of playlists (Sec1 to Sec3).

	HiHo	Regg	**Funk**	Elec	**Pop**	Rock
Sec1	19	3	8	28	13	29
Sec2	17	4	4	20	19	36
Sec3	12	3	4	22	16	42

Table 7. Distribution of songs across genres in playlists starting at Funk and ending at Pop. Results given for first, middle and last section of playlists (Sec1 to Sec3).

Funk is confused with almost all other genres and genre Pop strongly with genre Rock. As a result, the only visible trend in Tab. 7 is a rising amount of songs from genres Pop and Rock going from Sec1 to Sec3. This clearly indicates the limits of our approach to objective evaluation of playlist generation. Such an evaluation only makes sense with reliable genre label information.

The amount of songs which are being excluded from becoming members of the playlist because of being too dissimilar from both start and end song was set to $d = 95\%$ for all experiments (see step 2 in Sec. 3.2). Relaxing this constraint to smaller values leads to less clear distribution of genres (i.e. less songs in the playlists have the same genre label as the start and end songs).

4.2 Subjective evaluation

Our playlist generation algorithm can be utilised by users of the FM4 Soundpark website [2] to create their own digital "mixtapes" online. Therefore the best evaluation would be a user study with people actually using this service on the internet. Such a user study is planned for the future. During the development phase of the project, we decided to do an internal form of user study by having one of the authors listen to a number of playlists and judge their quality. This one person has considerable experience with popular music for having been a record collector and DJ for about two decades. While this approach has the problem of being highly subjective it does have the advantage of actually judging the "raw" playlists instead of a certain implementation and user interface.

As pointed out in Sec. 4.1, our playlist algorithm gives identical members of playlists in reversed order when start

[2] http://fm4.orf.at/soundpark

and end songs are exchanged. Therefore, we look at only $(6 \times (6-1))/2 = 15$ possible combinations of our six genres (see two leftmost columns in Tab. 8). For each combination of two genres A and B, we randomly choose three of the 50×50 playlists computed as described in Sec. 4.1. This gives 45 playlists for evaluation. Our evaluator listened to all the playlists using the "XMMS 1.2.10 - Cross platform multimedia player" [3] . He would first listen to the start song, then the end song and then the songs in between in the correct order. The evaluator was allowed to freely move through a song by skipping parts and moving back and forth in time. He was also allowed to re-listen to songs in the playlist if necessary.

For each playlist, the evaluator was asked to answer the following two questions which are tightly connected to our two hypotheses formulated in Sec. 4.1:

- How many outliers are in the playlist which do not fit the overall flavour of the playlist?
- Is the order of songs in the playlist from the start to the end song apparent?

The first question should allow to judge whether all the songs in a playlist really are similar to either the start or the end song, or are located somewhere in the intended middle. The second question aims at the sequential ordering of the songs. Songs at the beginning should be more similar to the start song, songs at the end to the end song. The second question can be answered with either "yes", "somewhat" or "no".

The results of the evaluation are given in Tab. 8. For each combination of genres, the average number of outliers is given (average taken over three playlists). It is also indicated how the second question has been answered for the three playlists of a certain combination of genres. Each "x" in a column stands for the respective answer given for one playlist. So for each row (i.e. combination of genres) three "x" indicate three answers to the second question. At the bottom row, the average number of outliers is given as well as the percentages of different answers to the question about the sequential order of the playlists is given.

The average number of outliers in a playlist is quite low at 1.1 out of possible 9. This means that on average, a user might want to delete one song from an automatically created playlist. While for a lot of combinations of genres this number is 0 and therefore perfect, for some genre combinations the number of outliers is quite high. E.g. for playlists starting at Hip Hop and ending at Reggae, an average of 4.7 songs are rated as outliers. The reasons seems to be that for a listener, the defining part of a Reggae song is the offbeat percussion which is not well conserved in our timbral representation of music. Instead, the rhythm guitar seems to dominate the models giving rise to high similarities with certain types of rock songs. Other sources of mistakes are recordings of poor acoustic quality which are found to be similar to each other no matter what the genres of the songs are. The sequential order of the playlists seems to work very well with it being apparent in 71% of all playlists and "somewhat" apparent in another 17.8%. One problem with the sequential ordering that we noticed is a kind of "tilting"-effect at the middle of playlists: the first half would be very close to the start song, the second half to the end song but a sort of smooth transition is missing. This was sometimes the case if start and end songs are very different and the data base might not even contain songs fitting in between. Another problem are too many outliers obscuring the overall order of a playlist.

As with the objective evaluation in Sec. 4.1, relaxing the amount of songs which are being excluded from becoming members of the playlist below $d = 95\%$ (see step 2 in Sec. 3.2) results in more outliers and less clear sequential order of the playlists.

Genres		# of	order apparent		
from	to	outliers	yes	somewhat	no
HiHo	Regg	4.7		x	xx
HiHo	Funk	1.7	xx	x	
HiHo	Elec	1.3	xxx		
HiHo	Pop	2.7		xx	x
HiHo	Rock	0	xxx		
Regg	Funk	0.7	xx	x	
Regg	Elec	1.3	xxx		
Regg	Pop	1.3	xxx		
Regg	Rock	0.3	xx		x
Funk	Elec	1.0	xx	x	
Funk	Pop	1.7	xx		x
Funk	Rock	0	xx	x	
Elec	Pop	0	xxx		
Elec	Rock	0	xx	x	
Pop	Rock	0	xxx		
average		1.1	71.1%	17.8%	11.1%

Table 8. Results of the subjective evaluation. For each combination of genres, the average number of outliers and the answers to the question concerning the order in the playlist is given. At the bottom row, average number of outliers as well as the percentages of different answers to the question about order are given.

[3] http://www.xmms.org

5 CONCLUSION

We have presented a new approach for the generation of playlists using start and end songs and showing inherent sequential order. Our approach is based on audio similarity and requires very little user interaction. Both objective evaluation based on genre labels of songs and subjective evaluation based on listening tests showed that the concept works well.

Our playlist generation algorithm can be utilised by users of the website of a public radio station to create their own digital "mixtapes" online. Since our evaluation showed that, on average, at least one song does not fit the overall playlist, an editing functionality might be added to the user interface.

6 ACKNOWLEDGEMENTS

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MULTI-FEATURE MODELING OF PULSE CLARITY: DESIGN, VALIDATION AND OPTIMIZATION

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ABSTRACT

Pulse clarity is considered as a high-level musical dimension that conveys how easily in a given musical piece, or a particular moment during that piece, listeners can perceive the underlying rhythmic or metrical pulsation. The objective of this study is to establish a composite model explaining pulse clarity judgments from the analysis of audio recordings. A dozen of descriptors have been designed, some of them dedicated to low-level characterizations of the onset detection curve, whereas the major part concentrates on descriptions of the periodicities developed throughout the temporal evolution of music. A high number of variants have been derived from the systematic exploration of alternative methods proposed in the literature on onset detection curve estimation. To evaluate the pulse clarity model and select the best predictors, 25 participants have rated the pulse clarity of one hundred excerpts from movie soundtracks. The mapping between the model predictions and the ratings was carried out via regressions. Nearly a half of listeners' rating variance can be explained via a combination of periodicity-based factors.

1 INTRODUCTION

This study is focused on one particular high-level dimension that may contribute to the subjective appreciation of music: namely *pulse clarity*, which conveys how easily listeners can perceive the underlying pulsation in music. This characterization of music seems to play an important role in musical genre recognition in particular, allowing a finer discrimination between genres that present similar average tempo, but that differ in the degree of emergence of the main pulsation over the rhythmic texture.

The notion of pulse clarity is considered in this study as a subjective measure that listeners were asked to rate whilst listening to a given set of musical excerpts. The aim is to model these behavioural responses using signal processing and statistical methods. An understanding of pulse clarity requires the precise determination of *what* is pulsed, and *how* it is pulsed. First of all, the temporal evolution of the music to be studied is usually described with a curve – denominated throughout the paper *onset detection curve* – where peaks indicate important events (considered as pulses, note onsets, etc.) that will contribute to the evocation of pulsation. In the proposed framework, the estimation of these primary representations is based on a compilation of state-of-the-art research in this area, enumerated in section 2. In a second step, the characterization of the pulse clarity is estimated through a description of the onset detection curve, either focused on local configurations (section 3), or describing the presence of periodicities (section 4). The objective of the experiment, described in section 5, is to select the best combination of predictors articulating primary representations and secondary descriptors, and correlating optimally with listeners' judgements.

The computational model and the statistical mapping have been designed using *MIRtoolbox* [11]. The resulting pulse clarity model, the onset detection estimators, and the statistical routines used for the mapping, have been integrated in the new version of *MIRtoolbox*, as mentioned in section 6.

2 COMPUTING THE ONSET DETECTION FUNCTION

In the analysis presented in this paper, several models for onset or beat detection and/or tempo estimation have been partially integrated into one single framework. Beats are considered as prominent energy-based onset locations, but more subtle onset positions (such as harmonic changes) might contribute to the global rhythmic organisation as well.

A simple strategy consists in computing the root-mean-square (RMS) energy of each successive frame of the signal ("rms" in figure 1). More generally, the estimation of the onset positions is based on a decomposition of the audio waveform along distinct frequency regions.

- This decomposition can be performed using a bank of filters ("filterbank"), featuring between six [14], and more than twenty bands [9]. Filterbanks used in the models are Gammatone ("Gamm." in table 1) and two sets of non-overlapping filters ("Scheirer" [14] and "Klapuri" [9]). The envelope is extracted from each band through signal rectification, low-pass filtering and down-sampling. The low-pass filtering ("LPF") is implemented using either a simple auto-regressive fil-

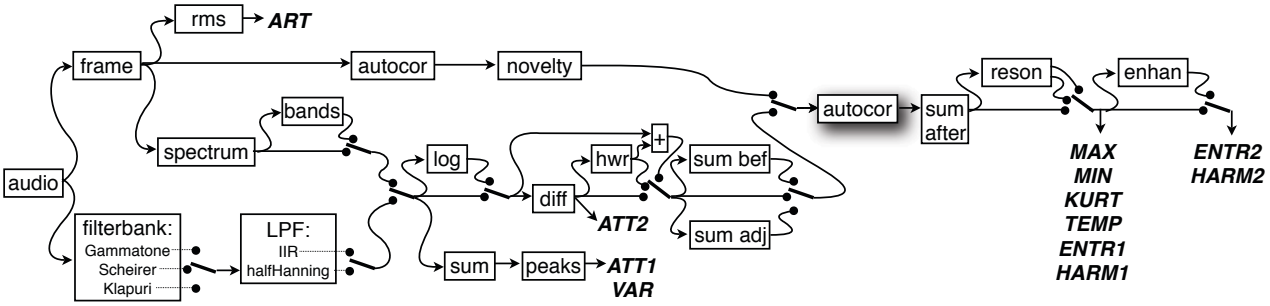

Figure 1. Flowchart of operators of the compound pulse clarity model, where options are indicated by switches.

ter ("IIR") or a convolution with a half-Hanning window ("halfHanning") [14, 9].

- Another method consists in computing a spectrogram ("spectrum") and reassigning the frequency ranges into a limited number of critical bands ("bands") [10]. The frame-by-frame succession of energy along each separate band, usually resampled to a higher rate, yields envelopes.

Important note onsets and rhythmical beats are characterised by significant rises of amplitude in the envelope. In order to emphasize those changes, the envelope is differentiated ("diff"). Differentiation of the logarithm ("log") of the envelope has also been advocated [9, 10]. The differentiated envelope can be subsequently half-wave rectified ("hwr") in order to focus on the increase of energy only. The half-wave rectified differentiated envelope can be summed ("+" in figure 1) with the non-differentiated envelope, using a specific λ weight fixed here to the value .8 proposed in [10] ("λ=.8" in tables 1 and 2).

Onset detection based on spectral flux ("flux" in table 1) [1, 2] – i.e. the estimation of spectral distance between successive frames – corresponds to the same envelope differentiation method ("diff") computed using the spectrogram approach ("spectrum"), but usually without reassignment of the frequency ranges into bands. The distances are hence computed for each frequency bin separately, and followed by a summation along the channels. Focus on increase of energy, where only the positive spectral differences between frames are summed, corresponds to the use of half-wave rectification. The computation can be performed in the complex domain in order to include phase information [1] [2].

Another method consists in computing distances not only between strictly successive frames, but also between all frames in a temporal neighbourhood of pre-specified width [3]. Inter-frame distances [2] are stored into a similarity matrix, and a "novelty" curve is computed by means of a convolution along the main diagonal of the similarity matrix with a Gaussian checkerboard kernel [8]. Intuitively, the novelty curve indicates the positions of transitions along the temporal evolution of the spectral distribution. We notice in particular that the use of novelty for multi-pitch extraction [16] leads to particular good results when estimating onsets from violin solos (see Figure 2), where high variability in pitch and energy due to vibrato makes it difficult to detect the note changes using strategies based on envelope extraction or spectral flux only.

[1] This last option, although available in *MIRtoolbox*, has not been integrated into the general pulse clarity framework yet and is therefore not taken into account in the statistical mapping presented in this paper.

[2] In our model, this method is applied to frame-decomposed autocorrelation ("autocor").

3 NON-PERIODIC CHARACTERIZATIONS OF THE ONSET DETECTION CURVE

Some characterizations of the pulse clarity might be estimated from general characteristics of the onset detection curve that do not relate to periodicity.

3.1 Articulation

Articulation, describing musical performances in terms of *staccato* or *legato*, may have an influence in the appreciation of pulse clarity. One candidate description of articulation is based on Average Silence Ratio (ASR), indicating the percentage of frames that have an RMS energy significantly lower than the mean RMS energy of all frames [7]. The ASR is similar to the low-energy rate [6], except the use of a different energy threshold: the ASR is meant to characterize significantly silent frames. This articulation variable has been integrated in our model, corresponding to predictor "*ART*" in Figure 1.

3.2 Attack characterization

Characteristics related to the attack phase of the notes can be obtained from the amplitude envelope of the signal.

- Local maxima of the amplitude envelope can be considered as ending positions of the related attack phases. A complete determination of each attack phase requires therefore an estimation of the starting position,

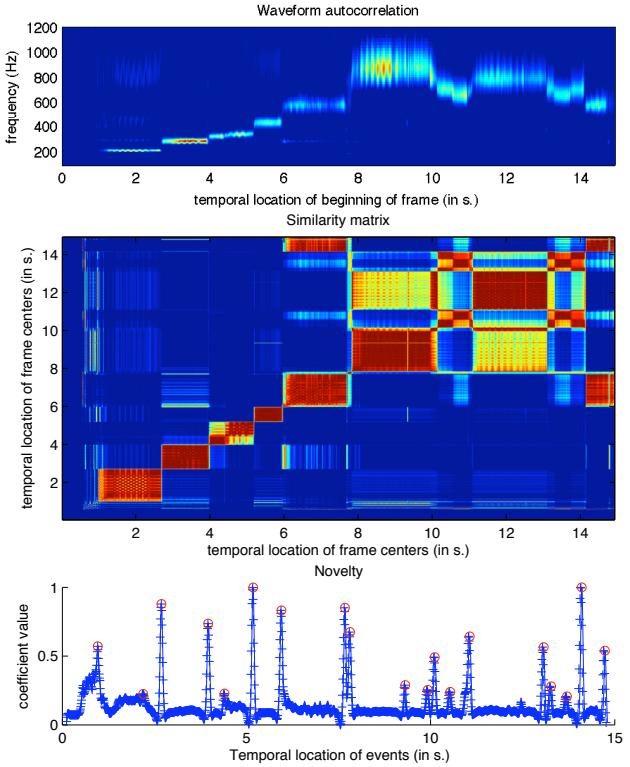

Figure 2. Analysis of a violin solo (without accompaniment). From top to bottom: 1. Frame-decomposed generalized and enhanced autocorrelation function [16] computed from the audio waveform; 2. Similarity matrix measured between the frames of the previous representation; 3. Novelty curve [8] estimated along the diagonal of the similarity matrix with onset detection (circles).

through an extraction of the preceding local minima using an appropriate smoothed version of the energy curve. The main slope of the attack phases [13] is considered as one possible factor (called "*ATT1*") for the prediction of pulse clarity.

- Alternatively, attack sharpness can be directly collected from the local maxima of the temporal derivative of the amplitude envelope ("*ATT2*") [10].

Finally, a variability factor "*VAR*" sums the amplitude difference between successive local extrema of the onset detection curve.

4 PERIODIC CHARACTERIZATION OF PULSE CLARITY

Besides local characterizations of onset detection curves, pulse clarity seems to relate more specifically to the degree of periodicity exhibited in these temporal representations.

4.1 Pulsation estimation

The periodicity of the onset curve can be assessed via autocorrelation ("autocor") [5]. If the onset curve is decomposed into several channels, as is generally the case for amplitude envelopes, the autocorrelation can be computed either in each channel separately, and summed afterwards ("sum after"), or it can be computed from the summation of the onset curves ("sum bef."). A more refined method consists in summing adjacent channels into a lower number of wider band ("sum adj."), on each of which is computed the autocorrelation, further summed afterwards ("sum after") [10].

Peaks indicate the most probable periodicities. In order to model the perception of musical pulses, most perceptually salient periodicities are emphasized by multiplying the autocorrelation function with a resonance function ("reson."). Two resonance curve have been considered, one presented in [15] ("reson1" in table 1), and a new curve developed for this study ("reson2"). In order to improve the results, redundant harmonics in the autocorrelation curve can be reduced by using an enhancement method ("enhan.") [16].

4.2 Previous work: Beat strength

One previous study on the dimension of pulse clarity [17] – where it is termed *beat strength* – is based on the computation of the autocorrelation function of the onset detection curve decomposed into frames. The three best periodicities are extracted. These periodicities – or more precisely, their related autocorrelation coefficients – are collected into a histogram. From the histogram, two estimations of beat strength are proposed: the *SUM* measure sums all the bins of the histogram, whereas the *PEAK* measure divides the maximum value to the main amplitude.

This approach is therefore aimed at understanding the global metrical aspect of an extensive musical piece. Our study, on the contrary, is focused on an understanding of the short-term characteristics of rhythmical pulse. Indeed, even musical excerpts as short as five second long can easily convey to the listeners various degrees of rhythmicity. The excerpts used in the experiments presented in next section are too short to be properly analyzed using the beat strength method.

4.3 Statistical description of the autocorrelation curve

Contrary to the beat strength strategy, our proposed approach is focused on the analysis of the autocorrelation function itself and attempts to extract from it any information related to the dominance of the pulsation.

- The most evident descriptor is the amplitude of the main peak ("*MAX*"), i.e., the global maximum of the curve. The maximum at the origin of the autocorrelation curve is used as a reference in order to normal-

ize the autocorrelation function. In this way, the actual values shown in the autocorrelation function correspond uniquely to periodic repetitions, and are not influenced by the global intensity of the total signal. The global maximum is extracted within a frequency range corresponding to perceptible rhythmic periodicities, i.e. for the range of tempi between 40 and 200 BPM.

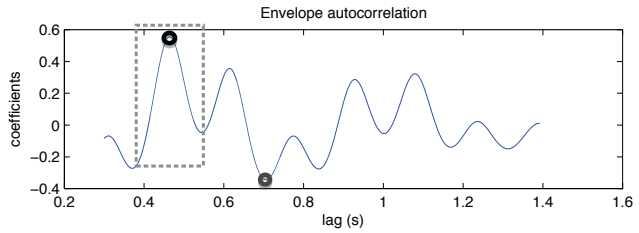

Figure 3. From the autocorrelation curve is extracted, among other features, the global maximum (black circle, *MAX*), the global minimum (grey circle, *MIN*), and the kurtosis of the lobe containing the main peak (dashed frame, *KURT*).

- The global minimum ("*MIN*") gives another aspect of the importance of the main pulsation. The motivation for including this measure lies in the fact that for periodic stimuli with a mean of zero the autocorrelation function shows minima with negative values, whereas for non-periodic stimuli this does not hold true.

- Another way of describing the clarity of a rhythmic pulsation consists in assessing whether the main pulsation is related to a very precise and stable periodicity, or if on the contrary the pulsation slightly oscillates around a range of possible periodicities. We propose to evaluate this characteristic through a direct observation of the autocorrelation function. In the first case, if the periodicity remains clear and stable, the autocorrelation function should display a clear peak at the corresponding periodicity, with significantly sharp slopes. In the second and opposite case, if the periodicity fluctuates, the peak should present far less sharpness and the slopes should be more gradual. This characteristic can be estimated by computing the kurtosis of the lobe of the autocorrelation function containing the major peak. The kurtosis, or more precisely the excess kurtosis of the main peak ("*KURT*"), returns a value close to zero if the peak resembles a Gaussian. Higher values of excess kurtosis correspond to higher sharpness of the peak.

- The entropy of the autocorrelation function ("*ENTR1*" for non-enhanced and "*ENTR2*" for enhanced autocorrelation, as mentioned in section 4.1) characterizes the simplicity of the function and provides in particular a measure of the peakiness of the function. This measure can be used to discriminate periodic and non-periodic signals. In particular, signals exhibiting periodic behaviour tend to have autocorrelation functions with clearer peaks and thus lower entropy than non-periodic ones.

- Another hypothesis is that the faster a tempo ("*TEMP*", located at the global maximum in the autocorrelation function) is, the more clearly it is perceived by the listeners. This conjecture is based on the fact that fast tempi imply a higher density of beats, supporting hence the metrical background.

4.4 Harmonic relations between pulsations

The clarity of a pulse seems to decrease if pulsations with no harmonic relations coexist. We propose to formalize this idea as follows. First a certain number N of peaks [3] are selected from the autocorrelation curve. Let the list of peak lags be $P = \{l_i\}_{i\in[0,N]}$, and let the first peak l_0 be related to the main pulsation. The list of peak amplitudes is $\{r(l_i)\}_{i\in[0,N]}$.

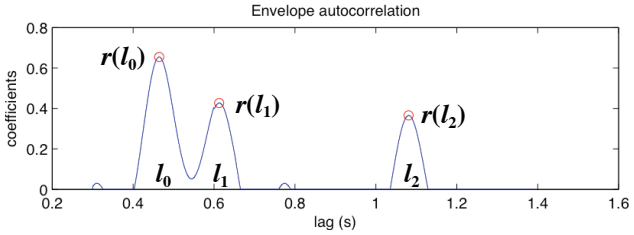

Figure 4. Peaks extracted from the enhanced autocorrelation function, with lags l_i and autocorrelation coefficient $r(l_i)$.

A peak will be inharmonic if the remainder of the euclidian division of its lag l_i with the lag of the main peak l_0 (and the inverted division as well) is significantly high. This defines the set of inharmonic peaks $\overline{H}$:

$$\overline{H} = \left\{ i \in [0,N] \;\middle|\; \begin{array}{ll} l_i \in [\alpha l_0, (1-\alpha) l_0] & (\mathrm{mod}\ l_0) \\ l_0 \in [\alpha l_i, (1-\alpha) l_i] & (\mathrm{mod}\ l_i) \end{array} \right\}$$

where α is a constant tuned to 0.15 in our implementation.

The degree of harmonicity is thus decreased by the cumulation of the autocorrelation coefficients related to the inharmonic peaks:

$$HARM = \exp\left(-\frac{1}{\beta} \frac{\sum_{i\in\overline{H}} r(l_i)}{r(l_0)} \right)$$

where β is another constant, initially tuned [4] to 4.

[3] By default all local maxima showing sufficient contrasts with respect to their adjacent local minima are selected.

[4] As explained in the next section, an automated normalization of the

5 MAPPING MODEL PREDICTIONS TO LISTENERS' RATINGS

The whole set of pulse clarity predictors, as described in the previous sections, has been computed using various methods for estimation of the onset detection curve [5] . In order to assess the validity of the models and select the best predictors, a listening experiment was carried out. From an initial database of 360 short excerpts of movie soundtracks, of 15 to 30 second length each, 100 five-second excerpts were selected, so that the chosen samples qualitatively cover a large range of pulse clarity (and also tonal clarity, another high-level feature studied in our research project). For instance, pulsation might be absent, ambiguous, or on the contrary clear or even excessively steady. The selection has been performed intuitively, by ear, but also with the support of a computational analysis of the database based on a first version of the harmonicity-based pulse clarity model.

25 musically trained participants were asked to rate the clarity of the beat for each of one hundred 5-second excerpts, on a nine-level scale whose extremities were labeled "unclear" and "clear", using a computer interface that randomized the excerpt orders individually [12]. These ratings were considerably homogenous (Cronbach alpha of 0.971) and therefore the mean ratings will be utilized in the following analysis.

Table 1. Best factors correlating with pulse clarity ratings, in decreasing order of correlation r with the ratings. Factor with cross-correlation κ exceeding .6 have been removed.

var	r	κ	parameters
MIN	.59		Klapuri, halfHanning, log, hwr, sum bef., reson1
KURT	.42	.55	Scheirer, IIR, sum aft.
HARM1	.40	.53	Scheirer, IIR, log, hwr, sum aft.
ENTR2	-.4	.54	Klapuri, IIR, log, hwr(λ=.8), sum bef., reson2
MIN	.40	.58	flux, reson1

The best factors correlating with the ratings are indicated in table 1. The best predictor is the global minimum of the autocorrelation function, with a correlation r of 0.59 with the ratings. Hence one simple description of the autocorrelation curve is able to explain already $r^2 = 36$ % of the variance of the listeners' ratings. For the following variables, κ indicates the highest cross-correlation with any factor of better r value. A low κ value would indicate a good independence of the related factor, with respect to the other factors considered as better predictors. Here however, the cross-correlation is quite high, with $\kappa > .5$. However, a stepwise regression between the ratings and the best predictors, as indicated in table 2, shows that a a linear combination of some of the best predictors enables to explain nearly half (47%) of the variability of listeners' ratings. Yet 53% of the variability remains to be explained...

Table 2. Result of stepwise regression between pulse clarity ratings and best predictors, with accumulated adjusted variance r^2 and standardized β coefficients.

step	var	r^2	β	parameters
1	*MIN*	.36	.97	Klapuri, halfHanning, log, hwr, sum bef., reson1
2	*TEMP*	.43	-.5	Gamm., halfHanning, log, hwr, sum aft., reson1
3	*ENTR1*	.47	-.55	Klapuri, IIR, log, hwr(λ=.8), sum bef.

6 *MIRTOOLBOX* 1.2

The whole set of algorithms used in this experiment has been implemented using *MIRtoolbox* [6] [11]: the set of operators available in the version 1.1 of the toolbox have been improved in order to incorporate a part of the onset extraction and tempo estimation approaches presented in this paper. The different paths indicated in the flowchart in figure 1 can be implemented in *MIRtoolbox* in alternative ways:

- The successive operations forming a given process can be called one after the other, and options related to each operator can be specified as arguments. For example,

```
a = miraudio('myfile.wav')
f = mirfilterbank(a,'Scheirer')
e = mirenvelope(f,'HalfHann')
```
etc.

- The whole process can be executed in one single command. For example, the estimation of pulse clarity based on the *MIN* heuristics computed using the implementation in [9] can be called this way:

```
mirpulseclarity('myfile.wav',
                'Min','Klapuri99')
```

distribution of all predictions is carried out before the statistical mapping, rendering the fine tuning of the β constant unnecessary.

[5] Due to the high combinatory of possible configurations, only a part has been computed so far. More complete optimization and validation of the whole framework will be included in the documentation of version 1.2 of *MIRtoolbox*, as explained in the next section.

[6] Available at *http://www.jyu.fi/music/coe/materials/mirtoolbox*

- A linear combination of best predictors, based on the results of the stepwise regression can be used as well. The number of factors to integrate in the model can be specified.

- Multiple paths of the pulse clarity general flowchart can be traversed simultaneously. At the extreme, the complete flowchart, with all the possible alternative switches, can be computed as well. Due to the complexity of such computation [7], optimization mechanisms limit redundant computations.

The routine performing the statistical mapping – between the listeners' ratings and the set of variables computed for the same set of audio recordings – is also available in version 1.2 of *MIRtoolbox*. This routine includes an optimization algorithm that automatically finds optimal Box-Cox transformations [4] of the data, ensuring that their distributions become sufficiently Gaussian, which is a prerequisite for correlation estimation.

7 ACKNOWLEDGEMENTS

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[7] In the complete flowchart shown in figure 1, as many as 4383 distinct predictors can be counted.

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QUANTIFYING METRICAL AMBIGUITY

Patrick Flanagan

ABSTRACT

This paper explores how data generated by meter induction models may be recycled to quantify metrical ambiguity, which is calculated by measuring the dispersion of metrical induction strengths across a population of possible meters. A measure of dispersion commonly used in economics to measure income inequality, the Gini coefficient, is introduced for this purpose. The value of this metric as a rhythmic descriptor is explored by quantifying the ambiguity of several common clave patterns and comparing the results to other metrics of rhythmic complexity and syncopation.

1 INTRODUCTION

This study initially grew out of an interest in modeling listener responses of a tapping study in which listeners tapped to the beat of sections of Steve Reich's *Piano Phase* [17]. For some sections, tapping rate and phase varied widely among listeners, and for others they displayed relative consistency. Accordingly, the goal of the study was to model metrical ambiguity as the degree of variety of listener responses to a given rhythm.

This task, however, would be complicated by a lack of simplified experimental data; the experiments of [17] involved pitched music, which added a layer of complexity I wanted to avoid. I therefore turned my attention to a corollary kind of metrical ambiguity, which is the amenability of a rhythm to gestalt flip, in which a listener reevaluates the location of the beat. This amenability has been convincingly advanced as one reason for the pull of African and African-derived rhythms on listeners [7]. Data on metrical gestalt flips is no easier to come by, but orientating the study around the proverbial average listener instead of groups of listeners allowed it to interface with the voluminous literature on syncopation. While metrical ambiguity is not equivalent to syncopation, the results of this study suggest that it roughly correlates to syncopation and offers an interesting rhythmic descriptor in its own right that could be used for theoretical analysis, genre classification, and the production of metrically ambiguous music.

To describe the ambiguity of the metrical scene, the ambiguity model recycles data that existing models of meter and beat induction often discard, the data about the induction strengths of *all* potential meters. The central idea of the model is that a more even distribution of induction strengths across a group of potential meters will cause greater ambiguity in the perception of meter. The model of ambiguity, therefore, requires a model of meter induction that, for a given rhythm, produces numerical values for an array of potential meters, with each meter defined by its period and phase relation to some arbitrary point in the rhythm. For the sake of simplicity, this study uses a hybrid model based on older models of meter induction that operate on quantized symbolic data, but the theory of ambiguity presented here is extensible to any model, including those that work with audio, that produces such an array.

In this paper the term "meter" refers to an isochronous, tactus-level pulse, characterized by a period and phase expressed in discrete integer multiples of some smallest unit of musical time. The more common term for this isochronous pulse is, of course, "the beat." "Meter," however, is the preferable term because this paper uses the conventionally understood downbeat of musical examples as a reference point for the tactus-level pulse.

2 METER INDUCTION MODEL

2.1 Influences on the Model

While the intent of this paper is not to advance a new model of meter induction, the existing models are either too complex for the purposes of this study or otherwise flawed. I therefore employ a hybrid meter induction model that calculates an induction strength for each candidate meter by summing the values of phenomenal accents that a given meter encounters, a strategy inherited from [12; 9; 2; 10]. Parncutt's model is the most sophisticated of the four in that it incorporates durational accent and tempo preference. For this reason, I largely adopt Parncutt's model.

Parncutt's model, however, has one flaw that prevents its full deployment in this study. In calculating what he called "pulse-match salience," which is the degree of match between phenomenal accents and an isochronous pulse stream, Parncutt summed the *product* of the phenomenal accents that occur on neighboring pulses. The problem with multiplying the phenomenal accents that occur on neighboring pulses is that some rhythms contain no inter-onset intervals (IOIs) that map onto candidate meters that we know to be viable meters. In this case, the metrical induction score, or pulse-match salience, of a viable meter turns out to be 0. For example, consider the Bossa Nova pattern in Figure 2.

The rhythm contains no inter-onset intervals of 4 tatums that begin on what, according to musical practice, is the

beat. There is one IOI of 4 tatums in the middle of the pattern, but it is shifted by 2 tatums off the beat. The pulse-match salience as calculated by Parncutt's formula for what is conventionally understood as the meter of this rhythm is 0, which clearly is not a desirable result. This problem did not crop up in Parncutt's article because the rhythmic patterns he investigated are relatively simple and unsyncopated and therefore have at least one IOI that coincides with a reasonable metrical candidate. As discussed below, the model presented here avoids this problem by summing the phenomenal accents that coincide with a candidate meter regardless of their adjacency.

2.2 The Meter Induction Algorithm

The model first describes a rhythmic surface as a series of values indexed to the tatums of the rhythm, with 0s signifying a rest or continuation of a previously attacked note and non-zero values signfiying phenomenal accents. The model follows that of Parncutt and uses only durational accent in modeling phenomenal accent and is calculated as

$$A_d\left(T\right) = \left(1 - \exp\left(\frac{-IOI\left(T\right)}{r}\right)\right)^i \qquad (1)$$

where T refers to the index of an event in tatums, exp is the natural exponential function, *IOI(T)* is the duration in milliseconds of the IOI from the onset at tatum *T* to the next onset, *r* is the saturation duration, and *i* is the accent index. Calculating *IOI(T)* requires fixing the speed (i.e., tatum duration), which is set to 200 ms for the examples in this paper. Saturation duration is the threshold of short term auditory memory, beyond which point lengthening the duration of an IOI will increase only marginally its durational accent. Parncutt, by optimizing the value of *r* to fit his experimental results, arrived at 500 ms, and I adopt this value in the model. The free parameter *i* magnifies the difference in accent between long and short durations. I follow Parncutt in setting it to 2.

The next step is generating the meters that will be tested against the accent vector. The model follows [11] in using only those meters with a period that is a factor of the length of the accent vector and excludes meters with extremely fast or slow tempos. Each candidate meter is further characterized by its phase, which is measured from index 0 of the accent vector. The preliminary induction strength *S* for meter *m*, characterized by period *P* and phase *Q*, over A_d, can then be calculated as

$$S_m = \frac{1}{n}\left(\sum_{i=0}^{n-1} A_d\left(Pi + Q\right)\right) \qquad (2)$$

where *n* is the number of beats of the meter that occur in the pattern. It is equal to the pattern length divided by the period of the meter. Multiplying the sum of accents by the reciprocal of *n* normalizes the induction strengths with respect to period length. As Parncutt noted, normalizing for period length allows us to explicitly introduce a model of human tempo preference, which numerous researchers have fixed at 600 ms [5]. Parcutt formalized tempo preference as

$$S_p = \exp\left(-\frac{1}{2}\left(\frac{1}{\sigma}\log_{10}\left(\frac{p}{u}\right)\right)^2\right) \qquad (3)$$

where S_p is the induction strength for period *p* of the meter and *u* and σ are the free parameters representing the preferred tempo and the standard deviation of its logarithm. Here I depart from Parncutt and use the commonly observed values of 600 ms and 0.2, respectively.

Incorporating the model of tempo preference into the induction strength model yields

$$S'_m = S_m S_p \qquad (4)$$

Applying (4) to the Bossa Nova pattern of Figure 2, assuming a speed of 200 ms, yields induction strengths for 16 different candidate meters, which are presented in the bar chart of Figure 1, ordered from lowest induction strength to highest. The next task is to quantify the evenness of this distribution.

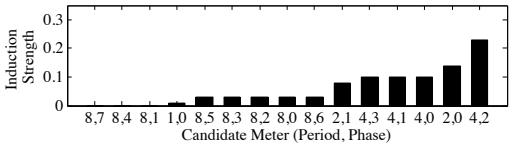

Figure 1. Induction strengths of the Bossa Nova pattern, ordered lowest to highest, with speed set to 200 ms.

3 THE GINI COEFFECIENT

Statisticians have developed a number of tools for quantifying the evenness of a distribution of values. One of them, the Gini coefficient, has gained popularity in economics as a measure of income inequality because it is easy to interpret and its value is independent of the scale of the data (i.e., unit of measurement) and the sample size. For this study, the inequality of the dispersion of induction strengths across a population of candidate meters is calculated. In the formula's discrete version, for a population *P* of n candidate meters with induction strengths calculated from (4), the Gini coefficient can be calculated as

$$G(P) = \frac{\sum_{i=1}^{n}\sum_{j=1}^{n} \mid x_i - x_j \mid}{2n^2\mu} \qquad (5)$$

where μ is the mean induction strength, and x_i and x_j are induction strengths of the candidate meters. It produces

values with a range of 0 to 1, where 1 signifies perfect inequality and 0 signifies perfect equality. In music theoretical terms, lower values denote greater metrical ambiguity, and vice versa. In practice, values for metrical ambiguity range from the high 0.20s for high metrical ambiguity up to the mid 0.70s for low metrical ambiguity. With the necessary tools in hand, we may now proceed to assess the metrical ambiguity of some rhythms.

4 METRICAL AMBIGUITY AND CLAVE RHYTHMS

4.1 Comparison to Syncopation Measures

Thus far the Gini coefficient has been described as a measure of metrical ambiguity, which is not necessarily synonymous with rhythmic complexity or syncopation. A fast isochronous pulse stream is metrically ambiguous because it is amenable to subjective rhythmization, but it is also rhythmically simple. Nevertheless, it is fruitful to compare the Gini coefficient to existing measures of syncopation.

Toussaint has tested several metrics of complexity and syncopation against clave rhythms in [16; 15], with particular attention paid to six of the most popular 4/4 timelines: the Bossa Nova, Gahu, Rumba, Soukous, Son, and Shiko patterns (Figure 2). Histograms of the induction strengths according to (4), ordered lowest to highest, of each rhythm with a moderate tatum duration of 200 ms are presented in Figure 3 (except for the Bossa Nova pattern, the histogram of which is presented in Figure 1). Note that the bars are labelled by their period and phase, and that the candidate meter that is most commonly understood to orgranize the rhythm in musical practice is 4,0. A simple visual inspection of the histograms reveals variation in the evenness of the distribution of induction strengths; to the eye, Shiko is the most uneven, and Bossa Nova the most even. As one would expect, these differences are reflected in the Gini coefficients of each rhythm, which are displayed in the first row of Table 1 alongside several metrics of rhythmic complexity and syncopation. In reading Table 1, be mindful that the Gini coefficient measures inequality of metrical induction strengths and therefore is inversely related to metrical ambiguity. A low value implies higher metrical ambiguity.

A word about the different metrics of complexity in Table 1. "Gini Syncopation" is derived from the Gini coefficient as explained in section 4.2. Michael Keith's measure of syncopation assigns each note to one of four categories: unsyncopated, hesitation, anticipation, and syncopation. Each category is associated with a different value or degree of syncopation, and the total syncopation of a passage equals the sum of its constituent notes' values. The reader is referred to [15] for more information. Pressing's measure [13] is similar in that it parses a rhythm into six basic units and sums their values.

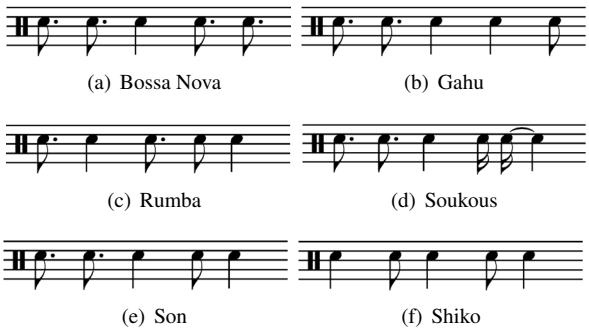

Figure 2. Six common 4/4 timelines.

	Bossa Nova	Gahu	Rumba	Soukous	Son	Shiko
Gini	0.583	0.719	0.679	0.683	0.679	0.803
Gini Syncopation	0.894	0.841	0.733	0.876	0.571	0.492
Keith	3	3	2	3	2	1
Off-Beatness	2	1	2	2	1	0
Pressing	22	19.5	17	15	14.5	6
Metrical Complexity	6	5	5	6	4	2
LHL	6	5	5	6	4	2
WNBD	4	3.6	3.6	3.6	2.8	1.2

Table 1. A comparison of measures of rhythmic complexity and syncopation for six clave rhythms.

Two different measures of rhythmic syncopation were proposed in [15]. The first is off-beatness, which is equal to the number of onsets that occur at indices of the measure that share no factors other than 1 with the length of the measure (a measure in usually indexed in 16th notes). The second measure of syncopation, called "metrical complexity," employs the metrical accent grid of [6]. In this scheme, the level of syncopation is related to the coincidence of onsets and shallow levels in Lerdahl and Jackendoff's metrical accent grid. The Longuet-Higgins Lee syncopation measure (LHL) also employs a metrical accent grid while considering whether an onset is followed a rest (or continuation) or another onset [8]. Finally, the Weighted Note-to-Beat Measure (WNBD) relates syncopation to the distance from each onset to the nearest beat [4].

The extreme Gini values in Table 1 correspond well to the other metrics of syncopation and to intuition; Bossa Nova is the most syncopated pattern and Shiko the least. Most of the metrics, however, have different interpretations of Gahu, Soukous, Son, and Rumba. According to [16], the patterns should be ordered in decreasing degree of syncopation as Bossa Nova, Gahu, Rumba/Soukous, Son, and Shiko, which is exactly the order of Pressing's model. I would have

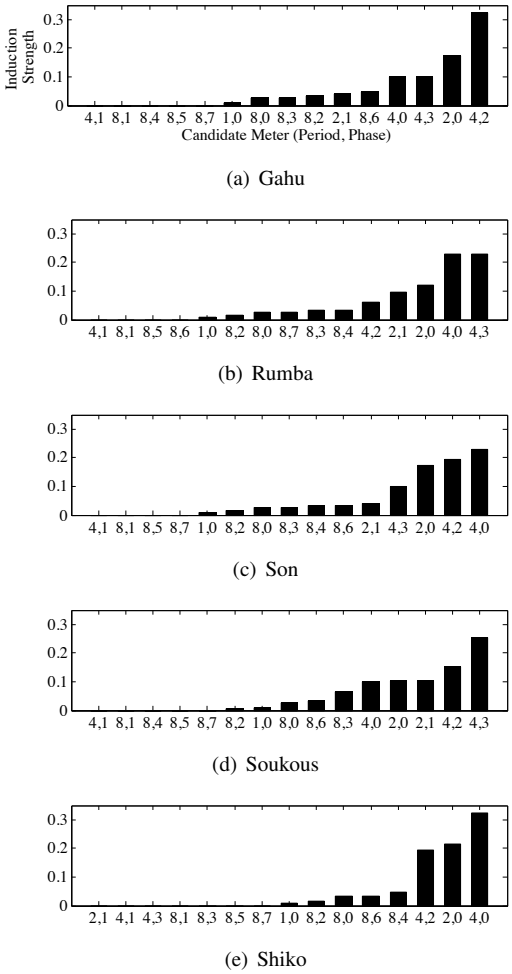

Figure 3. Histograms of the induction strengths of potential meters of five additional clave patterns. The two numbers below each bar represent the period and phase of a candidate meter.

to quibble with the assesment of Gahu as the second most syncopated pattern. The adjacency of the last two notes of Soukous, a uniquely tight spacing among the patterns, makes the pattern sound more syncopated by accenting the last onset, which is not on the beat, in two ways. First, the tight spacing of the fourth and fifth onsets and the relatively long IOI before the fourth onset makes the fourth onset function as a kind of pick-up to the fifth. Second, the last onset receives additional accent by virtue of the long inter-onset interval that follows it, which at five tatums, is the longest IOI of any of the patterns. The combined effect of the fourth onset functioning as a pick-up and the durational accent of the first onset lend strength to hearing the fifth onset as on the beat, an effect which, because the fifth onset is not on the beat, gives the rhythm a strongly syncopated feel. Gahu, by contrast, confirms the location of its "true" meter by placing its shortest IOI of two tatums between the last and first onsets. This eighth note IOI provides a pick-up into the real downbeat.

Clearly, experiments should be carried out to settle the matter, but for the sake of argument, let us bump Soukous ahead of Gahu on the syncopation scale. It would then read Bossa Nova, Soukous, Gahu, Rumba, Son, and Shiko. This slight rearrangement does nothing to help the Gini coefficient as a model of syncopation because it still ranks Son and Rumba as more complex than Soukous and Gahu. The reason it does this, however, is that it does not measure the metrical ambiguity of these particular rhythms but of the beat-class set types to which they belong. The Gini coefficient has no conception of a rhythm in a particular metrical context, whereas the other metrics assess a rhythm against a predetermined metrical framework, which listeners in real musical situations may infer from cues such as harmony and melody or other instruments. To this extent, comparing the Gini coefficient to the other metrics is comparing apples and oranges. As we shall see, using the Gini coefficient to describe beat-class sets is illuminating in certain contexts, but a metric of rhythmic complexity or syncopation should be able to distinguish a series of quarter notes that are on the beat from a series of quarter notes that are off the beat. The Gini coefficient does not do this.

4.2 From Metrical Ambiguity to Syncopation

There is a solution to this problem that, while adding additional mathematical complexity to an equation that is already far removed from the surface phenomena of the music, does at least correspond to our intuitions while offering several advantages over the existing metrics of rhythmic complexity and syncopation. The reason Gahu and Soukous are incorrectly assessed by the model is that it pays no heed to the relatively low induction score given to the meter that contextualizes the rhythm in musical practice (by definintion, the candidate meter 4,0). Indeed, one striking feature of Figures 2 and 4 is that for three of the rhythms (Bossa Nova, Gahu, and Soukous), the meter 4,0 is not even close to having the highest induction strength. A metric of syncopation should reflect this.

What is needed, then, to transform the Gini coefficient into a metric of syncopation in the mold of the others is a way of incorporating information about the strength of the meter understood to be the real meter relative to the other candidate meters. Furthermore, to aid comparison to other metrics, the score should increase with syncopation. The relative strength of the actual meter, 4,0, in comparison to the others can be neatly measured as its ranking, r, in the order list of induction strengths, and is set equal to the number of other meters that have an induction strength greater than or equal to that of the actual meter. Thus for Bossa Nova and Soukous, for example, r equals 5. We can use the reciprocal of r and the Gini coefficient to calculate a measure of

syncopation. For an accent vector A_d that produces a population *P* of meter induction strengths, its syncopation *Sync* may be described as

$$Sync(A_d) = 1 - \left(G(P)(1 - e^{-\frac{1}{r}})\right) \quad (6)$$

Inputting the values for the Gini coefficient and meter induction strengths calculated above yields the scores in the second row of Table 1.

These values correspond with what I have argued above is the most reasonable ranking of the patterns. This metric enjoys two other advantages over the others. The first is that it is invariant to scale. The other metrics invariably increase with the length of the rhythm being measured, making it impossible to compare rhythms of different lengths. The syncopation measure proposed here, on the other hand, always produces a value between 0 and 1. Second, it has a much finer grain of discrimination than the others. On the downside, it depends on a large number of free parameters, a circumstance that prevents fixing a final value for the syncopation of a rhythm conceived in the abstract; there is no single score for the Bossa Nova pattern. This fault could, however, be turned into an advantage if it allowed analysts to probe the effects of tempo and various kinds of phenomenal accent on syncopation.

4.3 Correspondence to Experimental Results

In [14], Spearman rank correlations were calculated between the experimental data of [12] and [3], which measured the difficulty of performing a set of rhythms, and the values produced by various models of syncopation for the same rhythms. The correlations give an indication of how well the models predict performance difficulty, which is assumed to be a reasonable proxy for syncopation. For the models tested, which were those listed in Table 1 (excluding the Gini coefficient models presented here), correlations ranged from 0.224 to 0.787. The procedure of [14] was repeated for the Gini coefficient and the Gini syncopation metrics, with the following results: The Gini coefficient showed correlations of 0.51 and 0.42 for the data of [12] and [3], respectively. The Gini syncopation measure produced correlations of 0.56 and 0.51.

These figures should treated cautiously. The Essens study [3] was a small experiment with only six participants and high *p*-values (i.e., low confidence). That said, the results indicate that the Gini coefficient and Gini syncopation measure are decent but not good predictors of performance difficulty. Considering that the best performing models in [14], the LHL and metrical complexity models, operate in reference to a full metrical hierarchy, one could speculate that the performance of the Gini measures could be improved by replacing the modified Parncutt beat induction model with one that uses a metrical hierarchy. It is also noteworthy that these same two models produced rankings of the six clave rhythms in Table 1 that agree with my own assessment and that of the Gini syncopation measure.

5 METRICAL AMBIGUITY OF BEAT-CLASS SET TYPES

The concept of the beat-class set was (re)introduced in [1] in order to analyze the properties and transformations of rhythmic material in Steve Reich's phase-shifting music (the concept originated in Babbitt's timepoint system). At issue in [1] was the transformational potential of rhythmic material, a topic that the Gini coefficient addresses because it describes how amenable a beat-class (bc) set type might be to various metrical interpretations. To investigate what the Gini coefficient might reveal about metrical ambiguity and beat-class sets, Gini coefficients for all mod 12 bc set types at three different speeds were calculated.

The most striking feature of the results is that the bc types with the highest ambiguity are those formed by one of the generator cycles of the modulo 12 system, the 5- (or 7-) cycle, the generator of the diatonic scale. In another interesting twist, if one skips applying rules for phenomenal accent and calculates induction strengths over an accent vector of 1s and 0s, the Gini coefficients of bc set types are invariant under M operation. Under M operation, the 5 cycle transforms into the 1-cycle, which means that, at least when accent rules are ignored, the two most famous scales of Western music, when transformed into rhythms, also turn out to be, at least according to the model developed here, the most metrically ambiguous bc set types. It might seem surprising that the dispersion of values produced by this model of meter induction, which was designed to reflect human preference for aligning isochronous downbeats with note onsets, would intersect so directly with set theory, but the explanation is almost intuitive.

By definition, the candidate meters have periods that are factors of the length of the meter. The generators cycles, by definition, have periods that are prime relative to the length of the meter and therefore prime relative to the periods of the candidate meters. The relative primeness of the candidate meters and the cycle generators ensures that the generator will cycle through all of the meters of a different phase and the same period before coinciding with the same meter twice. This cycling over candidate meters produces the most even distribution of onsets among the candidate meters, which in turn assures a low Gini coefficient.

It should be obvious from the status of cycle 1 generated beat-class sets as highly ambiguous that maximal ambiguity is hardly a guarantor of musical interest. As Cohn [1] noted in a parenthetical aside, these sets are "inherently less interesting for other reasons as well, although to articulate these reasons is not an easy task." One possible reason is that while bc sets generated from the 1-cycle are metrically ambiguous, their grouping structure is all too obvious. The

maximally even dispersion of onsets in the 5-cycle generated sets creates the potential for ambiguity of grouping as well as metrical structure, allowing grouping and metrical structures to interact in interesting ways. Alternatively, perhaps the maximally even sets are more interesting because they constantly flip the beat around, forcing the listener to retrospectively reevaluate the location in the metrical grid of what has already been heard. In this sense, the metrical ambiguity of maximally even rhythms is like that of the familiar visual phenomenon, in which the image of two faces opposite one another can be flipped to the background to produce an image of a vase. The perceptual gestalt is fragile but clearly delineated. The 1-cycle generated rhythms are more like looking at a fuzzy image. The gestalt is not so much ambiguous as amorphous. To be precise, their metrical structure is amorphous while the larger scale grouping structure is clear. One is confronted with a blob of notes followed by silence, a sonic image both blunt and vague.

6 CONCLUSION

The use of the Gini coefficient to quantify metrical ambiguity holds promise as a means to assess the syncopation of short, repeated rhythmic patterns and the perceptual qualities of beat-class set types. While a model of meter induction, based on older models, was introduced here for the purpose of creating input data for the Gini coefficient, the principle of using a dispersion of probabilities over candidate meters to calculate metrical ambiguity is in no way dependent on this model of meter induction. It may be fruitful to apply this principle to data from the more recent models that work with audio in order to use metrical ambiguity as a rhythmic descriptor of audio tracks. As it stands, the Gini coefficient may help analysts discuss rhythmic material in repetitive compositions and possibly serve as a utility for composers and other creators of rhythmically complex music.

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A MUSIC DATABASE AND QUERY SYSTEM FOR RECOMBINANT COMPOSITION

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ABSTRACT

We propose a design and implementation for a music information database and query system, the MusicDB, which can be used for Music Information Retrieval (MIR). The MusicDB is implemented as a Java package, and is loaded in MaxMSP using the mxj external. The MusicDB contains a music analysis module, capable of extracting musical information from standard MIDI files, and a search engine. The search engine accepts queries in the form of a simple six-part syntax, and can return a variety of different types of musical information, drawing on the encoded knowledge of musical form stored in the database.

1. INTRODUCTION

Inspired by the analysis techniques developed by David Cope for his work in the field of algorithmic composition by "music recombinance", the MusicDB [11] was originally designed as an analysis and data-retrieval back-end for an interactive music composition system. One of the primary motivations behind the design of this system was to allow the user to create original musical works, drawing on the musical language exemplified by their existing works, but maintaining as much flexibility as possible. We were drawn to Cope's notion of music recombinance because of its proven capacity to replicate musical style, which we felt would allow the user to easily incorporate the software into their existing compositional practice, and provide the greatest continuity between new works composed with the system and existing works. However, since the principle of paraphrasing which underlies the notion of "strict" recombinance can be somewhat of a limitation on originality, we felt it was important to develop a system that could promote musical continuity without relying exclusively on verbatim quotations of musical fragments for its base materials.

As the music recombinance approach to composition is clearly "data-driven" [6], the fundamental design of the MusicDB is well suited for use in the field of MIR. The software was designed to parse, analyse, and store information from symbolic musical representations—specifically, from standard MIDI files of 'scored' musical works—in a way which acknowledges similarities across multiple source works, but which also retains a formal 'map' of the original structure of each individual work. Single MIDI files of works are analysed by the software, and the analysis output is saved in the form of proprietary ".speac" data files. These files can then be loaded into a "session", consisting of an arbitrary number of analysed works. The database built during a session represents the compilation all of the analysed musical material into a single data structure, making it possible for the system to reveal inter-relationships between the elements of a potentially large body of analysed works.

1.1. Organization by Hierarchy

The MusicDB uses Cope's "SPEAC analysis" system to build a hierarchical analysis of each source work. A more detailed description of SPEAC is given later in the paper, and can also be found in numerous publications [3, 4, 5, 6]. This hierarchical analysis builds a tree-like structure for the representation of the musical form, in which the formal development of the music is segmented according to continuous changes observed over a variety of analysis parameters. In the MusicDB, this type of analysis is applied to each individual work, and is also used as an organizational system for the final session database. The end result is a data structure that associates the content of all the analysed works according to their overall pattern of formal development, regardless of their unique surface details. That is, the introductory, developmental, and concluding formal sections of all the stored works will be grouped together, regardless of the specific style or characteristics of each individual work. This allows the user to look at the formal contour of works, and find formal correlations between works, regardless of superficial and/or perceptual differences in musical content.

1.2. MusicDB Objectives

The MusicDB differs from Cope's work in that we have sought to develop a *component* for data analysis, storage, and recall, to be used as a tool by algorithmic composition

system designers. In particular, we have designed the MusicDB to accept input searches, and provide output data, at varying degrees of musical abstraction. While the system *can* provide complete, verbatim quotations of musical fragments from the analysed source works, it is not limited to such quotations. It can also be queried for more abstract representations, like melodic contour [12], chroma [13], kinesis [6], periodicity [10], and so on, thus offering the possibility of using musical knowledge provided by example works as a formal guideline for composition systems which are not otherwise based in music recombinance. In this sense, the MusicDB could be used strictly as a music analysis tool, or as a parameter control module for an agent-based [1, 7, 8], or fuzzy logic-based, system [2]—i.e., if such a system took kinesis (general activity level) and harmonic tension as parameters, such parameters could be provided by the MusicDB in a way which modelled the formal development exhibited by the analysed works.

2. IMPLEMENTATION

2.1. Music Analysis

The MusicDB breaks music apart using a hierarchical structuring of objects, which describe the score both vertically and horizontally. Beginning at the lowest level in the horizontal hierarchy, there are Events, Chords, and VoiceSegments. The vertical view of the score is described primarily by Group objects (though Chords obviously imply a vertical aspect as well), and at the top of the hierarchy both horizontal and vertical elements of the score are combined into a composite object called a Phrase (see Figure 1).

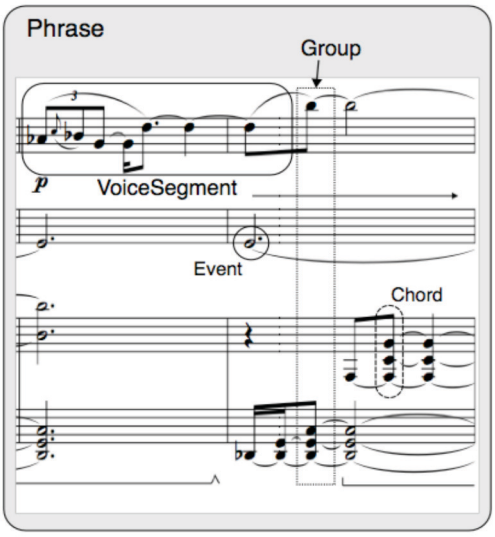

Figure 1. Music descriptors used by the MusicDB

An *Event* is a singular musical sound, occurring at a specific point in time[1]. *Chords* are two or more events, which occur at the same onset time (ED)[2], and a *VoiceSegment* is a horizontal sequence of Events and/or Chords, confined to a single MIDI channel. The decision to keep the voices in polyphonic parts together, where such parts are confined to a single MIDI channel, was a consequence of our desire to preserve, wherever possible, any special characteristics of idiomatic instrumental writing present in the source music. In the current implementation, VoiceSegments are confined to durations of 3 to 8 Chords or Events.

Groups constitute vertical 'slices' of the score, marked by static harmonic structures. In the MusicDB, the smallest Group duration is 1/8th-note. If the harmony remains static, Groups will extend for the entire duration of that harmony, and will always be lengthened in 1/8th-note increments. Groups are identified by an *entryPitch vector*, which is a sorted set of all pitches in the Group's harmonic structure.

Phrases are the largest component objects in the MusicDB, and are generally made from a combination of VoiceSegments and Groups. As the name suggests, a Phrase is a complete, semantically integrated, segment of music. Phrases are determined as sequences of Groups[3], and also hold references to all VoiceSegments found to intersect their Group sequence.

Cope's technique of SPEAC analysis [6] provides a segmentation of symbolic, or 'scored' music[4], based on the relative tensions of Groups. Groups are given a tension score derived from a set of analysis parameters, and are then assigned SPEAC 'labels' (**S**tatement, **P**reparation, **E**xtension, **A**ntecedent, **C**onsequent) based on their relative scores. A new segment, or Phrase, is created at each **A** to **C** label transition. The same process is then applied to the segmentation of the entire sequence of "foreground" Phrases, thus deriving a sequence of "background" Phrases (see Figure 4) and so on. The resulting SPEAC hierarchy provides a clear formal topography for the analysed work, giving the MusicDB access to the musical materials therein at the level of individual Groups, Phrases (and their component VoiceSegments), or sequences of Phrases (i.e., 'sections'), and also as a complete formal 'tree.' The hierarchical nature of SPEAC analysis is exploited by the MusicDB's search engine, which will be discussed later.

[1] The event's time is taken directly from the MIDI file, and is defined as the event's location in MIDI ticks (i.e., "timestamp").

[2] We use the concept of Entry Delay (ED), to identify the onset time of an event, calculated as the time passed *since* the previous event's onset.

[3] Phrases can also be made from sequences of other Phrases, as will be shown later in the paper.

[4] The SPEAC system is well-suited to the analysis of symbolic representations of music, not to audio recordings.

The MusicDB performs horizontal segmentation of Groups into Phrases (and Phrases into background Phrases) following the above SPEAC method, however, unlike in Cope, the MusicDB also uses a SPEAC-like system for segmenting sequences of events and/or chords on a single MIDI channel to form VoiceSegments. For this purpose, in addition to Cope's analysis parameters of rhythmic tension, duration tension, and "approach tension" (melodic interval tension), we've added measures for dynamic stress and pitch symmetry. Dynamic stress is simply a scaling of the current MIDI velocity value against the running mean velocity, in order to draw attention to strongly accented events. Pitch symmetry is a scaled interval measurement, which returns the absolute value of the current interval divided by 12. The intention is to show the "leapiness" of the melodic line, where low values indicate step-wise movement or repetitions, and values closer to 1.0 indicate leaps (intervals greater than an octave are set to 1.0).

2.2. Data Structure

The three primary objects used in the MusicDB's data structure—VoiceSegments, Groups, and Phrases—are tightly linked by object referencing. VoiceSegments hold a reference to the Phrase in which they originally appeared (their "root" Phrase), and also hold references to the sequence of Groups found to occur over their musical duration. Further, each VoiceSegment stores references to its "preceding" and "target" (following) VoiceSegment. In a similar manner, Groups hold a reference to their root Phrase, references to preceding and target Groups, and a list of VoiceSegments found to be playing during the vertical 'slice' from which the Group was taken. Phrases store references to all VoiceSegments active during their total duration, and the sequence of either Groups (in the case of "foreground" Phrases), or Phrases (in the case of "background" Phrases) from which they are made. Phrases also store references to their parent Phrases, where appropriate. This proliferation of references makes it computationally trivial to locate and extract a great deal of information about the musical form of a piece, given even the smallest aspect of its content. For example, it would be possible, using an iterative search process, to extract an entire instrumental part from a single musical work in the database, given only a short sequence of pitches, rhythmic values (EDs), chords, etc., as input. It is this interconnectivity by object referencing that is exploited by the MusicDB during the search process. As a convenience, all of the above objects also store the name of the source work in which they originally appeared, and VoiceSegments additionally store the instrument name assigned to the MIDI channel from which the VoiceSegment was derived.[1]

We would like to draw attention to the manner in which pitch-related searches are carried out in the MusicDB. For search processes involving specific pitch content—Pitch Lists and Chords—we have adopted Cope's scale of harmonic tension, rather than using Euclidean distance, or some other linear distance metric, as a measure for what we call "pitch distance." Cope assigns float values to each of the intervals within one octave, as follows [6]:

Interval	Value
0	0.0
1	1.0
2	0.8
3	0.225
4	0.2
5	0.55
6	0.65
7	0.1
8	0.275
9	0.25
10	0.7
11	0.9

All intervals greater than one octave are constrained to pitch-classes before their tension value is calculated. This sequence of values is based on the relative consonance of intervals within the octave, roughly following the harmonic series. It should not be viewed as a set of strictly 'tonal' relationships, though tonality generally follows a similar pattern, but rather as a measure of the degree to which a new note could be substituted for a given note, in an existing musical setting. It should be clear, therefore, that this is a metric chosen to apply to a broad range of *compositional* situations, but which would not be appropriate for most serial procedures, or for set theoretic analysis. Our application of the above harmonic tension scale as a pitch distance metric has the positive effect of relating searched items by a more acoustically founded notion of distance[2] than would be reflected by a linear distance measure. For example, using the above scale to find a match for the Pitch List {60, 62, 67} would return the list {67, 69, 60} as a "closer" match than the list {61, 64, 66}, in spite of the fact that the numeric values are clearly more distant, and the melodic contour is not the same. Assuming that this searched Pitch List was extracted from a supporting harmonic context, such a result would likely offer a more suitable alternative than would be provided by a linear distance metric. A search

[1] The instrument name is taken from the "name" metadata item in the MIDI file.

[2] Again, we would like to stress the fact that we sought to find a solution for a wide number of cases, and we are aware that serial and set theoretic compositional approaches will benefit less from the use of this distance metric.

for Pitch Contour, on the other hand, would use Euclidean distance to compare the Pitch Lists, and would thus have the opposite result.

2.3. Query System

Data is extracted from the MusicDB using a six-part query syntax (see Figure 2).

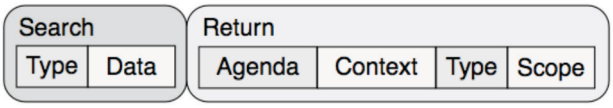

Figure 2. Format of a search query

Searches are carried out on VoiceSegment, Group, and Phrase objects, which hold the following fields[1]:

- **VoiceSegment**: Pitch List, Melodic Interval List, Pitch Contour, Chroma, ED List, ED Contour, Kinesis, Periodicity
- **Group**: Harmonic Interval List, Harmonic Tension, Harmonic Motive
- **Phrase**: Harmonic Tension, Kinesis, Periodicity, Chroma, Pitch Grid

The search **Type** can be chosen from a number of options:

1) PitchList: an ordered list of pitch values
2) MelodicIntervalList: an ordered list of interval values (signed integers)
3) PitchContour: a series of indices giving the relative "height" of each unique pitch in a Pitch List
4) Chroma: a 12-member vector indicating the occurrence-rate of pitch classes
5) EDList: an ordered list of ED values
6) EDContour: contour applied to an ED List
7) Kinesis: the general activity level (0. to 1.)
8) Periodicity: the rhythmic regularity of an ED list (0. to 1.)
9) Chord: an ascending list of pitches
10) HarmonicIntervalList: the intervals between adjacent pitches in a chord (i.e., a major triad is [0, 4, 3])
11) HarmonicTension: the interval tension of a given Chord[2] (0. to 1.)

The search **Data** is a vector of float values, used to represent the search **Type**.

The return **Agenda** (Figure 3) indicates the formal 'motivation' for the search. There are three options: Preceding, Current, and Following. A setting of Current will cause the search to return the requested field (indicated by the return Type) from the object containing the match, while Preceding and Following will return the same field from the preceding or following object, respectively. Beyond returning the matched field itself, the flexibility of the Agenda system makes it possible to extrapolate formal 'causes' and 'effects' relating to the searched data.

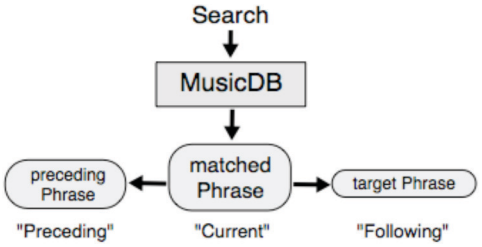

Figure 3. Search Agenda

The return **Context** indicates the immediate musical context of the object holding the requested field. There are three options: Element, Accompaniment, and Setting. A return Context of Element will return only the requested field from the matched object. The Setting option will attempt to provide data for a complete Phrase (i.e., a musical "setting") and will thus return the requested field from all objects referenced by the *root Phrase* holding the matched object[3]. If the requested field is held by a Group object, the *Group sequence* of that object's root Phrase will be returned[4]. For fields held by Groups the Accompaniment option is handled identically to Setting, but for fields held by VoiceSegments, Accompaniment removes the VoiceSegment in which the match was found, returning only the fields from the 'accompanying' VoiceSegments.

Return **Types** include all of the Search Types, with the addition of three more: PitchGrid, HarmonicMotive, and StatisticalRepresentation. A PitchGrid is a 128-member vector indicating the rate of occurrence for each available pitch. A HarmonicMotive is a sequence of Chords (these correspond to the *entryPitch vectors* used to identify Groups[5]), and a StatisticalRepresentation is a vector of statistical analysis values, used internally by the database to measure the similarity between instances of a given class (VoiceSegment, Group, or Phrase).

The final argument, **Scope** (Figures 4 and 5), controls the way in which searches are constrained by the formal structure of the database. A setting of Static (Figure 4) will

[1] All objects also hold a field for their Statistical Representation.

[2] The precise weighting of interval values used to calculate Harmonic Tension is taken from Cope [9].

[3] This is the case whenever the requested field belongs to a VoiceSegment or Group. If it belongs to a Phrase, the Phrase itself provides the return.

[4] The result is equivalent to a "Harmonic Motive".

[5] This is different from the Chord *object*, used by the VoiceSegment class for storing simultaneous pitch events. The term "chord" is used here to fit easily into musical parlance.

search the entire database, beginning at the top of the SPEAC hierarchy, and working its way through the tree. A setting of Progressive (Figure 5) will limit the scope of the search to a particular branch of the tree[1], and thus to a certain 'section' of the encoded musical form. With each completed Progressive search, the focus of the search 'steps forward', moving through the formal plan of the database. Static scoping can be useful for finding the best possible match for a desired search, while Progressive scoping would generally by used to move through a large-scale musical form.

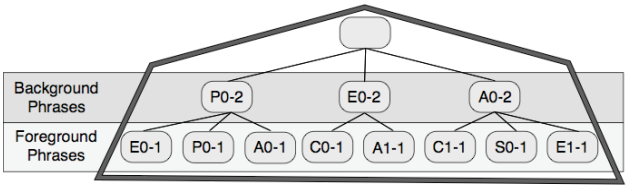

Figure 4. Static scoping of SPEAC hierarchy[2]

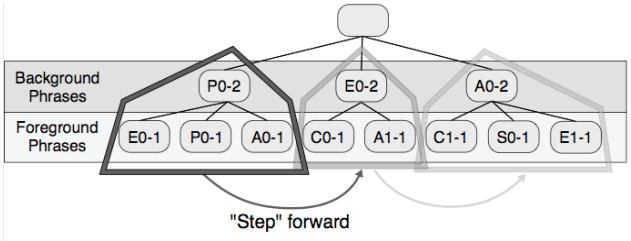

Figure 5. Progressive scoping of SPEAC hierarchy

Figure 6 demonstrates the way in which the MusicDB would handle searches with an input type of PitchList. It will be noticed that the actual data used for the return is always extracted in *relation* to the match found for the searched type. In the example, because PitchList is a field of the VoiceSegment object, the search for a best matching PitchList is carried out on VoiceSegments. From there, different return types (Chord, Contour, EDList) can be called, in relation to the VoiceSegment holding the match. In this case, the Contour and ED List are both stored by the VoiceSegment itself, whereas the Chord is stored by a Group, which is *referenced* by the VoiceSegment.

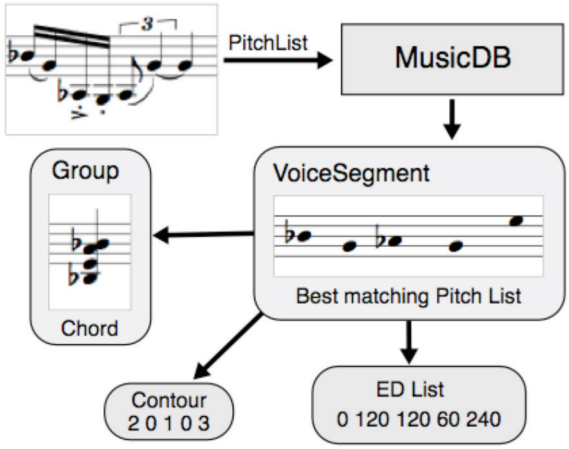

Figure 6. The system of references used to extract information from a given query

2.4. User Experience

Because input and output types are not required to match, the system offers great flexibility in the manner in which data is found and extracted. Take the following query as an example:

"PitchList 60 64 70 61 Current Accompaniment PitchContour Progressive"

This query would return all the Pitch Contours which accompanied the VoiceSegment that best matched the input Pitch List {60, 64, 70, 61}, given the current Scope, and would step 'forward' in the database once the search was returned. On the other hand, a query such as:

"PitchList 60 64 70 61 Following Element HarmonicMotive Static"

would search the *entire* database (due to the Static scoping) and return the harmonic sequence used by the Phrase *following* the VoiceSegment with the best matching Pitch List. Figure 7 shows a basic MaxMSP patch for querying the MusicDB.

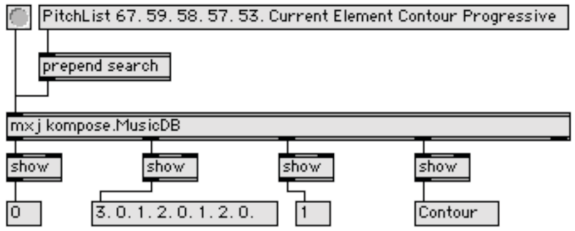

Figure 7. The MusicDB in MaxMSP. The second outlet reports the return data, the fourth return type, and outlets 1 and 3 provide Group count and voice number

[1] The branch chosen is dependent upon the last leaf to return a search. In the Figure 4 hierarchy, A0-1 would be followed by E0-2.

[2] This representation shows the tree-like pattern provided by SPEAC analysis. The label given to each node indicates the Phrase's formal function [9].

Composite searches are also possible, by entering multiple search queries *before* sending out the result. When queries are concatenated in this way, each query serves to narrow the scope of the search for the following query.

3. FUTURE DIRECTIONS

The MusicDB has been designed as a component for a larger system with the working title "ManuScore." This system will be a music notation-based application for interactive composition using principles of music recombinance. We will be working to extend the MusicDB itself by including the capacity for adding new material to the stored database dynamically, during the composition process. The other area we are working to develop concerns the discrimination of "quantized" versus "performed" rhythm. In particular, we are interested in refining the system in such a way as to make it more sensitive to the *differences* between quantized and performed rhythm, so that it may be given the capacity to apply a performed quality to otherwise quantized music, or to match the performance quality of fragments from two or more source works. One way we have thought to do this is by storing an event's ED as a *pair* of values, rather than a single value. The first value would indicate the quantized rhythmic position, and the second would indicate the offset of the performed rhythmic position from the quantized position. Analysis could be carried out using the quantized values, as in the current implementation, and the offsets could be used to reconstruct the rhythmic quality of the original performance. We are also considering the idea of integrating search categories more directly related to set theoretical analysis—perhaps a return type of "Set Similarity", for example—which could offer more flexibility in searching for materials based on specific pitch content.

Further, in order to introduce a more legitimate form of musical *inference* to the ManuScore system[1], we are also currently investigating the integration of Hierarchical Temporal Memory networks, modelled after those proposed by Jeff Hawkins, Dileep George, and Bobby Jaros at Numenta Inc. [9]. It is our feeling that the structural organization of the MusicDB could provide a useful pre-processing step in building the data representations used to train HTM networks, and could also provide category information for supervised training of the HTMs.

[1] Strictly speaking, the MusicDB is incapable of inference, being limited only to retrieving explicitly requested information from the data stored during analysis.

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COMBINING FEATURE KERNELS FOR SEMANTIC MUSIC RETRIEVAL

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ABSTRACT

We apply a new machine learning tool, *kernel combination*, to the task of semantic music retrieval. We use 4 different types of *acoustic content* and *social context* feature sets to describe a large music corpus and derive 4 individual kernel matrices from these feature sets. Each kernel is used to train a support vector machine (SVM) classifier for each semantic tag (e.g., 'aggressive', 'classic rock', 'distorted electric guitar') in a large tag vocabulary. We examine the individual performance of each feature kernel and then show how to learn an optimal linear combination of these kernels using convex optimization. We find that the retrieval performance of the SVMs trained using the combined kernel is superior to SVMs trained using the best individual kernel for a large number of tags. In addition, the weights placed on individual kernels in the linear combination reflect the relative importance of each feature set when predicting a tag.

1 INTRODUCTION

A significant amount of music information retrieval (MIR) research has focused on signal processing techniques that extract features from *audio content*. Often, these feature sets are designed to reflect different aspects of music such as timbre, harmony, melody and rhythm [15, 9]. In addition, researchers use data from online sources that places music in a *social context* [6, 16]. While individual sets of audio content and social context features have been shown to be useful for various MIR tasks (e.g., classification, similarity, recommendation), it is unclear how to combine multiple feature sets for these tasks.

This paper presents a principled approach to combining multiple feature sets by applying a useful machine learning technique: *kernel combination* [7]. A *kernel function* for each feature set measures the similarity between pairs of songs. Kernel functions can be designed to accommodate heterogeneous feature representations such as vectors [8], sets of vectors [4], distributions [10], or graphs [1]. We use the kernel function to compute a *kernel matrix* that encodes the similarity between all pairs of songs. From multiple feature sets, we compute multiple kernel matrices for a set of songs. We find the optimal linear combination of the kernel matrices using quadratically-constrained quadratic-programming (i.e., convex optimization). The result is a single kernel matrix that is used by a support vector machine (SVM) to produce a decision boundary (i.e., a classifier).

In this paper, we use kernel combination to integrate music information from two audio content and two social context feature sets. For each semantic tag (e.g., "aggressive", "classic rock", "distorted electric guitar") in our vocabulary, we learn the linear combination of the four kernel matrices that is optimal for SVM classification and rank-order songs based on their relevance to the tag. The weights that are used to produce the combined kernel reflect the importance of each feature set when producing an optimal classifier.

In the following subsection, we discuss related work on music feature representations, kernel methods in MIR research, and the origins of kernel combination. We continue in Section 2 with an overview of kernel methods and the kernel combination algorithm. Section 3 introduces a number of features that capture information about musical content and context and describes how to build song-similarity kernels from these features. Section 4 describes our experiments using individual and combined heterogenous feature kernels for semantic music retrieval. Section 5 concludes with a discussion of our findings.

1.1 Related Work

McKinney and Breebaart [9] use 4 different feature sets to represent audio content and evaluate their individual performance on general audio classification and 7-way genre classification. They determine that features based on the temporal modulation envelope of low-level spectral features are most useful for these tasks but suggest that intelligent combinations of features might improve performance. Tzanetakis and Cook [15] present a number of content-based features and concatenate them to produce a single vector to represent each song for use in genre classification.

Knees et al. [6] use semantic data mined from the results of web-searches for songs, albums and artists to generate a *contextual* description of the music based on large-scale, social input, rather than features that describe the audio *content*. Using this context data alone achieves retrieval results comparable to the best content-based methods. Whitman and Ellis [16] also leverage web-mined record reviews to develop an unbiased music annotation system.

Flexer et al. [3] combine information from two feature sources: tempo and MFCCs. They use a nearest-neighbor classifier on each feature space to determine an independent class-conditional probability for each genre, given each feature set. Using a naïve Bayesian combination, they multiply these two probabilities and find that the resulting probability improves 8-way genre classification of dance music.

Lanckriet et al. [7] propose a more sophisticated method for combining information from multiple feature sources. They use a kernel matrix for *each* feature set to summarize similarity between data points and demonstrate that it is possible to learn a linear combination of these kernels that optimizes performance on a discriminative classification task. They show that, for a protein classification task, an optimal combination of heterogenous feature kernels performs better than any individual feature kernel.

Discriminative SVM classifiers have been successfully applied to many MIR tasks. Mandel and Ellis [8] use patterns in the mean and co-variance of a song's MFCC features to detect artists. Meng and Shawe-Taylor [10] use a multivariate autoregressive model to describe songs. Using Jebara et al.'s probability product kernels [5], they kernelize the information contained in these generative models and then use an SVM to classify songs into 11 genres.

2 KERNEL METHODS

A simple solution for binary classification problems is to find a linear discriminant function that separates the two classes in the data vector space. In practice, the data may not lie in a valid vector space or may not be linearly separable. A linear discriminant may still be learned using a kernel function to map the data items into a new kernel-induced vector (Hilbert) space, called the feature space *F*.

If we consider mapping a pair of data items $\mathbf{x}_i, \mathbf{x}_j \in \Omega$ with $\Phi : \Omega \rightarrow F$, kernel methods only require the inner product of the data features $\Phi(\mathbf{x}_i), \Phi(\mathbf{x}_j) \in F$. The kernel function $K(\mathbf{x}_i, \mathbf{x}_j) = \langle \Phi(\mathbf{x}_i), \Phi(\mathbf{x}_j) \rangle$ completely specifies this inner product relationship. Thus, a data set $\{\mathbf{x}_i, i \in 1, ..., n\}$ can be specified in the feature space with the kernel matrix $\mathbf{K}$ where each element (i, j) is obtained by the kernel function $K(\mathbf{x}_i, \mathbf{x}_j)$. See [12] for details.

If we use a valid kernel function, the resulting kernel matrix $\mathbf{K}$ must be positive semi-definite. Henceforth, kernel matrix and kernel function will be referred to as "kernel" with the meaning being clear from the context. Many different kernels are used in practice and we have experimented with several of them. A natural question that arises is which kernel to use. Our work and the work of Lanckriet et al. [7] suggests that settling for a single kernel may not be optimal and that a combination of kernels can improve results.

2.1 Kernelizing Feature Data

2.1.1 Radial Basis Function

The Radial Basis Function (RBF) kernel is one of the most commonly used kernels. It is defined as:

$$\mathbf{K}_{i,j} = \exp(-\frac{\|\mathbf{x}_i - \mathbf{x}_j\|^2}{2\sigma^2}).$$

The hyper-parameter σ is learned by cross validation [12].

2.1.2 Probability Product Kernel

The Probability Product Kernel (PPK) is a natural candidate for data items that have been modeled with probability distributions [5]. We have found experimentally that the PPK tends to outperform other distribution-capturing kernels (such as the Kullback-Leibler divergence kernel space[11]), and is easier and more elegant to compute. We model each data item $\mathbf{x}_i$ with a distribution $p(\mathbf{x}; \theta_i)$ specified by parameters θ_i and compute the PPK as:

$$\mathbf{K}_{i,j} = \int p(\mathbf{x}; \theta_i)^\rho p(\mathbf{x}; \theta_j)^\rho d\mathbf{x},$$

where $\rho > 0$. When $\rho = 1/2$, the PPK corresponds to the Bhattacharyya distance between two distributions. This case has a geometric interpretation similar to the inner-product between two vectors. That is, the distance measures the cosine of the angle between the two distributions. Another advantage of the PPK is that closed-form solutions for Gaussian mixture models (GMM's) are available [5], whereas this is not the case for the Kullback-Leibler divergence.

2.1.3 Normalized Kernels

It can also be useful to "normalize" the kernel matrix. This projects each entry in the kernel matrix onto the unit sphere and entries can be geometrically interpreted as the cosine of the angle between the two data vectors. Furthermore, when combining multiple kernels, normalized kernels will become "equally" important. We normalize $\mathbf{K}$ by setting:

$$\mathbf{K}_{i,j}^{norm} = \frac{\mathbf{K}_{i,j}}{\sqrt{\mathbf{K}_{i,i}\mathbf{K}_{j,j}}}.$$

The resulting kernel matrix has all ones on its diagonal, helping ensure numerical stability in the combination algorithm below. Henceforth all kernels are assumed to be normalized.

2.2 Support Vector Machine (SVM)

The SVM learns the separating hyperplane that maximizes the margin between two classes. We present the SVM here to establish notation and clarify the formulation of kernel combination. The algorithm considered is the 1-norm soft

margin SVM which can be cast as the following optimization problem:

$$\min_{\mathbf{w},b,\zeta \succeq 0} \mathbf{w}^T\mathbf{w} + C\sum_{i=1}^{n}\zeta_i \quad \text{subject to: } y_i(\mathbf{w}^T\Phi(\mathbf{x}_i)+b) \geq 1-\zeta_i, \forall i, \tag{1}$$

where $\mathbf{x}_i$ is the data point we wish to classify and $y_i \in \{+1,-1\}$ is its corresponding label. $\Phi(\mathbf{x}_i)$ is the data point mapped to the feature space. $\mathbf{w}$ is the normal vector to the hyperplane that defines the discriminant boundary in the feature space and b is the hyperplane offset. C is a regularization parameter that penalizes misclassifications or within-margin points and ζ_i are slack variables that "relax" the hard-margin constraints of the SVM.

The the dual formulation of 1-norm soft margin optimization in Equation 1 is:

$$\max_{0\leq\alpha\leq C,\alpha^T\mathbf{y}=0} 2\alpha^T\mathbf{e} - \alpha^T\text{diag}(\mathbf{y})\mathbf{K}\text{diag}(\mathbf{y})\alpha, \tag{2}$$

where $\mathbf{e}$ is an n-vector of ones and $\alpha \in \mathcal{R}^n$ is the vector of Lagrange multipliers. The solution to this dual problem is a function of K and results from the point-wise maximum of a series of affine functions in alpha. Because the point-wise maximum of affine functions is convex, the solution to the dual problem is convex in K.

The SVM classifies a new, unlabeled data point $\mathbf{x}_{new}$ based on the sign of the linear decision function:

$$f(\mathbf{x}_{new}) = \mathbf{w}^T\Phi(\mathbf{x}_i) + b = \sum_{i=1}^{n}\alpha_i\mathbf{K}(\mathbf{x}_i,\mathbf{x}_{new}) + b. \tag{3}$$

For example, the classification task might be to determine whether a given song represents the tag "guitar" or not. It is common to associate positive examples (e.g. those classified as "guitar") with the label "+1" and negative examples with "-1." This labeling is achieved by taking the sign of the decision function $sign(f(\mathbf{x}))$.

2.3 Kernel Combination SVM

Lanckriet et al. [7] propose a linear combination of m different kernels, each encoding different features of the data:

$$\mathbf{K} = \sum_{i=1}^{m}\mu_i\mathbf{K}_i,$$

where $\mathbf{K}_i$ are the individual kernels formulated via feature extraction methods described in Section 3. A useful property of kernel matrices is that, since they are positive semi-definite, a positive linear combination of a set of kernels will also be a positive semi-definite kernel [12]. That is,

$$\mathbf{K} = \sum_i \mu_i\mathbf{K}_i,\ \mu_i > 0,\ \mathbf{K}_i \succeq 0 \quad \forall i \quad \Rightarrow \quad \mathbf{K} \succeq 0.$$

The kernel combination problem reduces to learning the set of weights, μ, that combine the feature kernels, $\mathbf{K}_i$, into the "optimum" kernel. To ensure the positive semi-definiteness of $\mathbf{K}$, the weights are restricted to be positive, $\mu_i \geq 0$.

We wish to ensure that the weights μ sum to one. This prevents any weight from growing too large, forces the kernel matrices to be combined convexly and gives interpretability to the relative importance of each kernel. We can achieve this with the constraint $\mu^T\mathbf{e} = 1$. The optimum value of the dual problem in Equation 2 is inversely proportional to the margin and is convex in $\mathbf{K}$. Thus, the optimum $\mathbf{K}$ can be learned by minimizing the function that optimizes the dual (thereby maximizing the margin) with respect to the kernel weights, μ.

$$\min_{\mu}\left\{\max_{0\leq\alpha\leq C,\alpha^T\mathbf{y}=0} 2\alpha^T\mathbf{e} - \alpha^T\text{diag}(\mathbf{y})\mathbf{K}\text{diag}(\mathbf{y})\alpha\right\} \quad \text{subject to: } \mu^T\mathbf{e}=1,\ \mu_i \geq 0 \quad \forall i=1,...,m, \tag{4}$$

where $\mathbf{K} = \sum_{i=1}^{m}\mu_i\mathbf{K}_i$.

Since the objective function is linear in μ, and hence convex, we can write this min-max problem equivalently as a max-min problem [2]. We may recast this problem in epigraph form and deduce the following quadratically constrained quadratic program (QCQP)[7]:

$$\max_{\alpha,t} 2\alpha^T\mathbf{e} - t \quad \text{subject to: } t \leq \alpha^T\text{diag}(\mathbf{y})\mathbf{K}_i\text{diag}(\mathbf{y})\alpha, i=1,\ldots,m,\ 0\leq\alpha\leq C, \alpha^T\mathbf{y}=0. \tag{5}$$

The solution to this QCQP will return the optimum convex combination of kernel matrices. For a more complete derivation of the kernel combination algorithm, see [7].

3 KERNEL CONSTRUCTION FROM AUDIO FEATURES

We experiment with a number of popular MIR feature sets which attempt to represent different aspects of music: timbre, harmony and social context.

3.1 Audio Content Features

Audio content features are extracted directly from the audio track. Each track is represented as a set of feature vectors, $\mathcal{X} = \{\mathbf{x_1},...,\mathbf{x_T}\}$, where each feature vector $\mathbf{x_t}$ represents an audio segment, and T depends on the length of the song. We integrate the set of feature vectors for a song into a single representation by estimating the parameters of a probability distribution, approximated with a Gaussian mixture model, over the audio feature space. Finally, we compute a kernel matrix using set of song GMMs with the probability product kernel technique [5] described in Section 2.1.2.

3.1.1 Mel-Frequency Cepstral Coefficients

Mel-frequency cepstral coefficients (MFCCs) are a popular feature for a number of music information retrieval tasks [8, 10, 3]. For a 22050Hz-sampled, monaural song, we compute the first 13 MFCCs for each half-overlapping short-time ($\sim$23 msec) segment and append the first and second instantaneous derivatives of each MFCC. This results in about 5,000 39-dimensional **MFCC+delta** feature vectors per 30 seconds of audio content.

We summarize an entire song by modeling the distribution of its MFCC+delta features with an 8-component GMM. We sample 5,000 feature vectors from random times throughout a song, learn a MFCC+delta GMM for each song and convert the GMM parameters of every song into a probability product kernel matrix. For our semantic music retrieval task, we find that using GMMs with diagonal covariances results in better performance than using single, full-covariance Gaussian distributions.

3.1.2 Chroma

Chroma features [4] represent of the harmonic content of a short-time window of audio by computing the spectral energy present at frequencies that correspond to each of the 12 notes in a standard chromatic scale. We extract a 12-dimensional chroma feature every $\frac{1}{4}$ second and, as with the MFCCs above, model the distribution of a song's chroma features with a GMM and create a probability product kernel matrix. We also investigated features that describe a song's key (estimated as the mean chroma vector) and tempo (in beats per minute) but found that these features performed poorly for semantic music retrieval.

3.2 Social Context Features

We can also summarize each song in our dataset with a *semantic* feature vector. Each element of this vector indicates the relative strength of association between the song and a semantic tag. We propose two methods for collecting this semantic information: social tags and web-mined tags. Given these semantic feature vectors, we compute an RBF kernel for each data source as described in Section 2.1.1. The semantic feature vectors tend to be *sparse* as most songs are annotated with only a few tags. When tags are missing for a song, we set that dimension of the semantic feature to zero. In addition to being sparse, many semantic context features are *noisy* and do not always reflect an accurate semantic relationship between a song and a tag. Songs which have not been annotated with any tags are assigned the average semantic feature vector (i.e., the estimated prior probabilities of the tags in the vocabulary).

3.2.1 Social Tags

For each song in our dataset, we attempt to collect two lists of social (raw-text) tags from the Last.fm website (www.last.fm). The first list relates the *song* to a set of tags where each tag has a score that ranges from 0 (low) to 100 (high) as a function of both the number and diversity of users who have annotated that song with the tag. The second list associates the *artist* with tags and aggregates the tag scores for all the songs by that artist. We find the scores for all relevant song and artist tags as well as their synonyms (determined by the authors). For example, a song is considered to be annotated with "down tempo" if it has instead been annotated with "slow beat". We also allow wildcard matches for tags so that, for example, "blues" matches with "delta electric blues" and "rhythm & blues". To create the Last.fm semantic feature vector, we add the song and artist tag scores into a single vector.

3.2.2 Web-Mined Documents

We extract tags from a corpus of web documents using the relevance scoring (RS) algorithm [6]. To generate tags for a set of songs, RS works by first repeatedly querying a search engine with each song title, artist name and album title to obtain a large corpus of relevant web-documents. We restrict the search to a set of musically-relevant sites (see [13] for details). From these queries, we retain the (many-to-many) mappings between the songs and the documents. Then we use each tag as a query string to find all the relevant documents from our corpus, each with an associated relevance weight. By summing the relevance weights for the documents associated with a song, we calculate a score for each song-tag pair. The semantic feature vector for a song is then the vector of song-tag scores for all tags in our vocabulary.

4 SEMANTIC MUSIC RETRIEVAL EXPERIMENTS

We experiment on the CAL-500 data set [14]: 500 songs by 500 unique artists, each annotated by a minimum of 3 individuals using a 174-tag vocabulary representing genres, instruments, emotions and other musically relevant concepts. The kernel combination algorithm requires at least 50 data points (and often more) from each class to learn a reliable classifier so, for the experiments reported here, we require that each tag be associated with at least 50 songs and remove some redundant tags, reducing the vocabulary to 61 tags [1].

Given a tag (e.g., "jazz"), the goal is to rank all songs by their relevance to this query (e.g. jazz songs at the top). We learn a decision boundary for each tag (e.g., a boundary between jazz / not jazz) and rank all test songs by their distance (positive or negative) from this boundary, calculated using Equation 3. The songs which most strongly embody the query tag should have a large positive distance from

[1] For the list of tags used, see http://cosmal.ucsd.edu/cal

the boundary. Conversely, less semantically relevant songs should have a small or negative distance from the boundary.

We compare the SVM results to the human-annotated labels provided in the CAL-500 dataset where a song is positively associated with a tag if 80% of test subjects agree that the tag is relevant. We quantify the ranking returned by the discriminative SVM classifier using the area under the receiver operating characteristic (ROC) curve. The ROC compares the rate of correct detections to false alarms at each point in the ranking. A perfect ranking (i.e., all the relevant songs at the top) results in an ROC area equal to one. Ranking songs randomly, we expect the ROC area to be 0.5. We also compute mean average precision (MAP) by moving down the ranked list of test songs and averaging precisions at every point where we correctly identify a new song.

One benefit of using ROC area as a metric to evaluate rankings is that it is immune to differences in the tags' prior probabilities. The tag frequencies in the data set roughly follow an exponential distribution with most terms having far more negative than positive examples. The average prior probability over all tags is 16.9%. While measuring binary classification performance would have to overcome a bias towards the negative class, evaluating ranking performance (using ROC area) is a better measure of what's important for real MIR applications and does not suffer from this imbalance. For example, 339 of the 500 songs were annotated as having "Male Lead Vocals" while only 56 songs were judged to be from the "Electronica" genre. In the latter case, a classifier could achieve 88% accuracy by never labeling any songs as "Electronica" while its average ROC area would be 0.5 (random).

4.1 Single Kernel Results

For each of the four features described in Section 3, we construct a kernel and train a one-vs-all SVM classifier for each tag where the negative examples are all songs not labeled with that tag. We train SVMs using 400 songs, optimize regularization parameter, C, using a validation set of 50 songs and use this final model to report results on a test set of 50 songs. The performance of each kernel, averaged over all tags and 10-fold cross validation (so that each song appears in the test set exactly once), is shown in Table 1. The rightmost column of Table 1 shows the number of tags for which each of the different features resulted in the best single kernel for ranking songs. The bottom row of Table 1 shows the theoretical upper bound that an 'oracle' could achieve by choosing the single best feature for each tag, based on test-set performance.

Table 1 demonstrates that the MFCC content features are most useful for about 60% of the tags while social context features are best for the other 40%. However the facts that all kernels individually perform well above random and that three of four perform best for many tags indicate that all features can be useful for semantically ranking songs.

Feature	ROC area	MAP	Best Kernel
MFCC	0.71	0.50	37
Chroma	0.60	0.40	0
Last.fm	0.68	0.49	16
Web Docs	0.67	0.48	8
Single Oracle	0.73	0.54	

Table 1. Individual feature kernel SVM retrieval results and the number of tags for which each kernel was the best.

4.2 Kernel Combination Results

The kernel combination SVM can integrate the four sources of feature information to enhance our music retrieval system. In practice, we find that some tags do not benefit from this integration since they have reached maximum performance with their individual best feature kernel. With appropriate setting of the regularization parameter, C, it is generally possible to get the combined kernel to correctly place all weight on this single best kernel and the result in these cases is the same as for the single best kernel. 31 of the 61 tags (51%) show an improvement in ROC area of 0.01 or more while 14 tags (23%) show no significant change. 16 tags (26%) get worse, we suspect due to insufficient data to train and optimize the parameters of the combined kernel SVM.

Table 2 shows the average retrieval ROC area and MAP results for the combined kernel as well as those that could be obtained if the single best feature were known in advance. We break down results by tag category and show how many tags from each category benefit from kernel combination. Despite the facts that some tags perform worse using kernel combination and that knowing which single feature is best requires looking at the test set, the combined kernel improvement is significant (paired t-test, $N = 61, \alpha = 0.05$).

Though all kernels are normalized, the absolute value of a weight does not give direct insight into the relevance of that feature to the classification task. If its weight is zero (as happens if we include a randomly-generated kernel), the feature is truly useless but even a small positive weight can allow a feature kernel to have a large impact. However, examining the relative feature weights in Figure 1 for the 45 words where the algorithm succeeds does allow us to make some interesting comparisons of how the features' utility varies for different tags. The simplex in Figure 1 shows that the MFCC kernel tends to get most weight, unsurprising since it is the single most reliable feature. We also see a preference for Last.fm over Web Docs and both get more weight than Chroma, in accordance with their individual performances in Table 1. The spectral MFCC features are most useful for genres like "Rock" as well as the ubiquitous "Drum Set" which most web users neglect to tag. The human-derived context information gets more weight for nebulous tags like "Powerful" as well as both "Male" & "Female Lead Vocals" which, while only represented in a few MFCC vectors, are easily identified by human listeners.

Tag Category	Single Oracle ROC	Single Oracle MAP	Kernel Combo ROC	Kernel Combo MAP	Improve / Total
All tags	0.73	0.54	0.74	0.54	31 / 61
Emotion	0.71	0.51	0.72	0.52	14 / 28
Genre	0.80	0.58	0.82	0.58	3 / 5
Instrument	0.74	0.55	0.76	0.56	8 / 10
Song	0.73	0.60	0.73	0.60	5 / 15
Usage	0.70	0.34	0.69	0.35	0 / 1
Vocals	0.69	0.37	0.69	0.39	1 / 2

Table 2. Music retrieval results for 61 CAL500 tags. "Single Oracle" picks the single best feature for each tag, based on test set performance. "Kernel Combo" learns the optimum combination of feature kernels from the training set.

It may be possible to enhance these results by adding even more feature kernels. In practice however, as the number of kernels grows, over-fitting can be a problem. Keeping C low ensures the optimum solution is well regularized. In the experiments reported here, optimum results for both single and combined kernel SVMs are found by optimizing a different value of C for the positive and negative class examples.

5 DISCUSSION

This paper uses a novel machine learning tool to combine a number of existing MIR features in a manner that optimizes retrieval performance for a variety of semantic tags. The results show significant improvement in retrieval performance for 51% of the 61 tags considered. Although discriminative SVM classification may not be the best solution for semantic retrieval problems, the results reported here (ROC area of 0.71 for MFCC+delta, 0.73 for the single best feature, 0.74 for kernel combination) compare favorably to the generative framework in [14] which reports retrieval ROC area of 0.71 on the CAL-500 data using the MFCC+delta features (although that work used a larger vocabulary of 174 tags).

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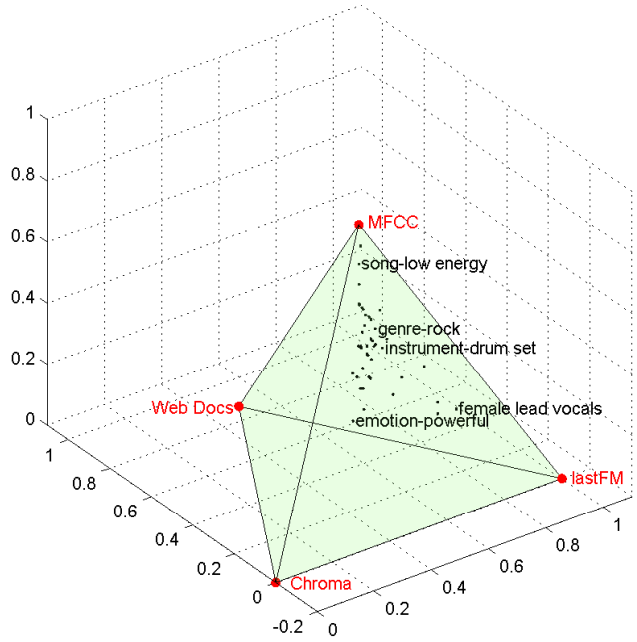

Figure 1. Optimal feature weights for 45 tags from the CAL-500 data set. Five tags are named for illustrative purposes.

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CLUSTERING MUSIC RECORDINGS BY THEIR KEYS

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ABSTRACT

Music key, a high level feature of musical audio, is an effective tool for structural analysis of musical works. This paper presents a novel unsupervised approach for clustering music recordings by their keys. Based on chroma-based features extracted from acoustic signals, an inter-recording distance metric which characterizes diversity of pitch distribution together with harmonic center of music pieces, is introduced to measure dissimilarities among musical features. Then, recordings are divided into categories via unsupervised clustering, where the best number of clusters can be determined automatically by minimizing estimated Rand Index. Any existing technique for key detection can then be employed to identify key assignment for each cluster. Empirical evaluation on a dataset of 91 pop songs illustrates an average cluster purity of 57.3% and a Rand Index of close to 50%, thus highlighting the possibility of integration with existing key identification techniques to improve accuracy, based on strong cross-correlation data available from this framework for input dataset.

1. INTRODUCTION

Musical key which specifies the tonal center (also called tonic), describes the hierarchical pitch relationship in a composition. Tonic refers to the most stable pitch in a music piece, upon which all other pitches are referenced and scale implies pitch set which occur in a passage and interval between them. Therefore, key is extremely important for music representation and conveys semantic information about a composition. Automatic key estimation can be applied to many problems in content-based analysis of music, such as structure analysis and emotion detection, and also in music retrieval & recommendation systems.

Although considerable work can be found in the literature addressing the problem of estimating music key from audio signal automatically, it is still a challenging task. Major difficulties lie in the fact that key is a high level feature and difficult to extract from audio signals based on complexities of polyphonic audio analysis. Krumhansl [9] proposed a Maximum key-profile correlation(MKC) method that compares spectrum of music piece with key profiles which are derived by probe tone rating task and represent perceived stability of each chroma within the context of a particular key. The key that provides the maximum correlation with the music piece is considered as the solution[13]. Some modifications of MKC are involved, such as [19] which introduces Bayesian probabilistic model to infer key profiles from a pattern of notes. In [2], Chew described a mathematical model called Spiral Array, where pitch class, chord and key are spatially aligned to points in 3D space using a knowledge-based approach. The distance from tonal center of the music piece to the key representation acts as a likely indicator, and key estimation is performed through finding the nearest neighbor of the music piece. İzmirli[6] calculated similarity of tonal evolution among music pieces via dynamic time warping. Martens *et. al.*[12] compared decision tree methods with distance-based approaches. However, supervised classification approaches need a large scale of annotated data to train a model and new samples can't be reliably detected. Shenoy and Wang[16] adopted a rule-based method that combines higher level musical knowledge with lower level audio features. Based on a chromagram representation for each beat spaced frame of audio, triads are detected and matched against templates of key patterns, to identify the key with the highest ranking. Zhu and Kankanhalli[23] attempted to extract precise pitch profile features for key finding algorithms, considering the interference of pitch mistuning and percussive noise. Gómez and Herrera [4] extracted kinds of descriptors which can represent tonal content of a music piece and introduced machine learning models into key estimation. HMM based models have also been used to identify keys[11,14] and to detect key changes[1] by modeling timing and context information during a music performance. Most of the techniques discussed above suffer from the fact that complexities in polyphonic audio analysis make it rather difficult to accurately determine the actual template or position for each key.

In this paper, we propose a novel, unsupervised method to cluster music recordings into several categories without performing actual key identification. We believe that

This work was performed in National University of Singapore

elimination of training process, along with not having a dependency on template patterns of musical keys, makes this unsupervised framework applicable across a broad range of musical styles and can be fairly easily ported to other forms of clustering by changing input feature set. The performance of existing key identification approaches in the literature, though high, is observed to drop as dataset increases or musical recordings do not satisfy set criteria, like the presence of a strong beat and time signature of 4/4. Cluster information can thus serve as a valuable input to increase accuracy of key identification, on the basis of strong cross correlation of songs in a specific cluster, and the perceived stability of cluster purity over a large data set, demonstrated later in the evaluation section.

The rest of paper is organized as follows: the overview of the proposed approach is introduced in Section 2. The details of chroma-based extraction and inter-recording dissimilarity measurement are presented in Section 3. Section 4 describes the approach for cluster generation and number of clusters estimation. Experimental results are discussed in Section 5. Finally, conclusions are drawn in Section 6.

2. METHOD OVERVIEW

2.1. Problem Formulation

Given a dataset of N musical recordings, each one performed in one of P different keys, where P, the actual number of keys in the specific dataset, is unknown. Our aim is to produce a partitioning of the N recordings into M clusters such that M=P, and each cluster consists exclusively of recordings associated with the same key. Existing key identification algorithms could then potentially yield an improved performance from this cluster information. This schematic for the proposed framework is shown in Figure1 below.

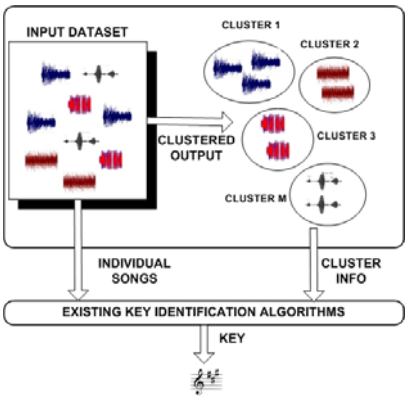


Figure 1. Clustering music recordings by their keys

2.2. System Configuration

As illustrated in Figure 2, the proposed clustering system consists of four major components: chroma-based feature extraction, computation of inter-recording dissimilarities, cluster generation and estimation of cluster number.

In the phase of feature extraction, pitch is estimated from the audio signal by spectral analysis technique and mapped into a chromagram. The dissimilarity computation based on chromagram is designed to produce small values for dissimilarities between recordings associated with the same key and large values for dissimilarities between recordings associated with different keys. Then, clusters are generated in a bottom-up agglomerative manner, followed by an automatic cluster number estimation algorithm.

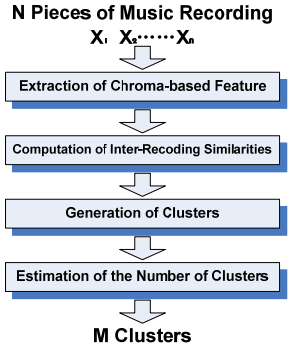


Figure 2. Framework for music clustering by keys

3. DISSIMILARITY MEASURE

Inter-recording dissimilarity computation is the most critical task in our work. The dissimilarity between recordings serves as a distance metric which imparts position and distribution of samples in the space. A good distance metric helps to gather similar samples together and make them easy to cluster. This section provides a viable approach to measure dissimilarity between music recordings, where spectrum divergence as well as harmonic center is taken into account.

3.1. Chroma-based Feature Extraction

Chroma-based[17] feature is a musical representation of audio data, where spectrum is reduced to 12 bins, corresponding to pitch classes in music theory. Intensity of frequencies is analyzed from audio and projected onto chroma scale. Two pitches separated by an integral number of octaves are mapped to the same element in a chroma vector. The output feature for each frame is a 12-dimensional vector, called chromagram, which stores the distribution of the energy for each of twelve semitones.

In our current system, input audio data is divided into half-overlapping 32ms long frames with Hanning widow. In each frame, spectrum is obtained via FFT. Then energy of pitches sharing the same pitch class are summed up and assigned to the corresponding bin in chroma vector.

3.2. SKL Divergence across Chroma Distributions

Kullback–Leibler divergence[10] in statistics, is a measure of difference between probability distributions. Its symmetrical version, SKL divergence, is proven to be effective for evaluation of distance between spectral envelops of audio signal[8,22]. Based on chroma vectors discussed in the previous section, we calculate expectations of chroma components and normalize them by total energy in each music piece. This spectral envelope can be interpreted as probability distribution of chroma components in a music piece. Thus, SKL divergence is utilized to measure difference between two chroma distributions with respect to musical keys.

3.3. Center of Effect

The spiral array[2] is a computational geometric representation for modeling tonality where pitches are mapped to points along a spiral in 3D coordinates. First, pitch classes are indexed by intervals of perfect fifths from a reference pitch, C for example. Then one increment in the index which stands for the interval of perfect fifth, leads to rotation of one quarter in horizontal plane as well as a height gain. And pitches with a major third apart (four increments of the index), result in vertical alignment with each other. This property satisfies the fact that interval of perfect fifth is responsible for the most consonant of the unison, while major third is the second. Then center of effect(ce) of a music piece is defined as the arithmetic mean of each pitch class weighted with its duration.

In our implementation, most of the configuration is similar to [2]. However, when calculating center of effect, we adopt energy in chromagram as the weight coefficient rather than duration and refined center of effect as:

$$ce = \sum_{i=1}^{12}\sum_{j=1}^{N}\frac{e_{ij}}{E} \times pitch_j \text{ where } E = \sum_{i=1}^{12}\sum_{j=1}^{N} e_{ij} \quad (1)$$

where $pitch_i$ denotes the coordinate of i^{th} pitch class and N is the total number of frames.

3.4. Inter-Recording Dissimilarity Measure

We utilize the linear combination of SKL divergence across chroma spectrum and Euclidean distance between center of effect to compute the overall dissimilarity between music recordings as follow:

$Div_{spectrum}$ denotes SKL divergence of the chroma spectrum envelope, while Dis_{ce} is the Euclidean distance between center of effect of two music recordings. Thus, the overall dissimilarity is

$$Dissim(i, j) = \alpha\beta Dis_{ce}(i, j) + (1-\alpha) Div_{spectrum} \quad (2)$$

where β is a normalization factor which normalizes both metrics into the same order, and α, a weighting coefficient, implies the bias between tonic and scale. They are set to 0.25 and 0.20 respectively by experience in real implementation. The linear combination of the two metrics satisfies the assumption that once one of the dissimilarity metrics increases, the overall dissimilarity will go up.

4. CLUSTER GENERATION

4.1. Agglomerative Clustering

After computing inter-recording dissimilarities, the next step is to assign the recordings deemed similar to each other to the same cluster. This is done by an hierarchical clustering method[7], which sequentially merges the recordings deemed similar to each other. The similarity is inferred from the metric described in Section 3.4. The algorithm consists of the following procedure:

```
Begin
    initialize M←N, and form clusters Ci←{Xi}, i=1, 2,..,N
    Do
        find the most similar pair of clusters, say Ci and Cj
        merge Ci and Cj
        M←M−1
    Until M = 1
End
```

Outcome of the agglomeration procedure is a cluster tree with the number of clusters ranging from 1 to N. The tree is then cut by optimizing number of cluster, which corresponds to an estimation of the number of keys actually occurring in the dataset.

4.2. Estimating Number of Clusters

According to the music theoretic basis of 24 keys (12 Major and 12 Relative Minor[1]) and the fact that not all these keys may be necessarily included in any sample data set, we limit the cluster number to be a maximum of 24 in our framework. However, in order to obtain a higher purity which is important for further processing, we can relax this limitation to a number, slightly higher than 24. Experiment shows that as long as the cluster number range is in a small neighborhood, the performance varies slightly. Additionally, we utilize a automatic cluster number estimation method, which has been used successfully to detect the number of speakers in the scenario of speaker clustering[21].

To evaluate the accuracy of clustering algorithm, here we follow the manner of [21], using two basic metrics: cluster purity[18] and Rand Index[5,15]. Cluster purity indicates the degree of agreement in a cluster. The purity for the m-th cluster C_m is defined as:

$$purity_m = \sum_{p=1}^{P}\left(\frac{n_{mp}}{n_{m*}}\right)^2 \quad (3)$$

where n_{mp} is the number of recordings in cluster C_m that are performed in the p-th key and n_{m*} is the number of

[1] We only consider major keys and natural minor keys here. In the rest of this paper, "minor key" refers to natural minor key except as noted.

recordings in the cluster of C_m. Deriving from Equation 3, *purity* follows $\frac{1}{n_{m*}} \le purity_m \le 1$ and is proportion to the probability that two music recordings in a cluster are in the same key. Specifically, the overall performance can be evaluated using average purity for all clusters:

$$\overline{purity} = \frac{1}{M}\sum_{m=1}^{M}(n_m \times purity_m) \quad (4)$$

The average purity is monotonically increasing with cluster number. This is based on the fact that as the number of clusters increases, average count of recordings in each cluster decreases, which leads to a higher purity. Hence, purity is not suitable for evaluating clustering performance, when the number of clusters is uncertain.

In contrast, Rand index, implying the extent of divergence of clustering result, is the number of incorrect pairs, actually performed in the same key but are placed in different clusters and vice versa.

Let n_{*p} denotes the number of recordings associated with the p-th key. Rand index can be calculated by:

$$R(M) = \sum_{m=1}^{M} n_{m*}^2 + \sum_{p=1}^{P} n_{*p}^2 - 2\sum_{m=1}^{M}\sum_{p=1}^{P} n_{mp}^2 \quad (5)$$

Rand index can also be represented as a mis-clustering rate:

$$R(M) \text{ in percentage} = \frac{\sum_{m=1}^{M} n_{m*}^2 + \sum_{p=1}^{P} n_{*p}^2 - 2\sum_{m=1}^{M}\sum_{p=1}^{P} n_{mp}^2}{\sum_{m=1}^{M} n_{m*}^2 + \sum_{p=1}^{P} n_{*p}^2} \times 100\% \quad (6)$$

It's obvious that the smaller the value of $R(M)$ is, the better the cluster performance will be. It has been proven that the approximately minimal value will be achieved, when the number of cluster is equal to the actual number of keys occurring in the dataset[21]. So our task is to search for a proper cluster number, such that Rand Index is minimized.

Recalling the Rand Index in Equation 5, the first term in the right side of the equation, $\sum_{m=1}^{M} n_{m*}^2$, can be computed based on the clustering result. Meanwhile the second term, $\sum_{p=1}^{P} n_{*p}^2$, is a constant irrelevant to clustering. The third term, $\sum_{m=1}^{M}\sum_{p=1}^{P} n_{mp}^2$ requires that the true key attribute of each recording is known in advance, which cannot be computed directly. To solve this problem, we represent it by

$$\begin{aligned}\sum_{m=1}^{M}\sum_{p=1}^{P} n_{mp}^2 &= \sum_{m=1}^{M}\sum_{p=1}^{P}[\sum_{i=1}^{N}\delta(h_i,m)\delta(o_i,p)]^2 \\ &= \sum_{m=1}^{M}\sum_{p=1}^{P}[\sum_{i=1}^{N}\delta(h_i,m)\delta(o_i,p)][\sum_{j=1}^{N}\delta(h_j,m)\delta(o_j,p)] \\ &= \sum_{m=1}^{M}\sum_{p=1}^{P}\sum_{i=1}^{N}\sum_{j=1}^{N}\delta(h_i,m)\delta(o_i,p)\delta(h_j,m)\delta(o_j,p) \\ &= \sum_{i=1}^{N}\sum_{j=1}^{N}\delta(h_i,h_j)\delta(o_i,o_j)\end{aligned} \quad (7)$$

where $\delta(\cdot)$ is Kronecker Delta function, h_i is the index of cluster where the i-th recording is located, and o_i is the true key attribute of the i-th recording. Note that h_i, $1 \leqslant i \leqslant N$, is an integer between 1 and M, if M clusters are generated. The term $\delta(o_i,o_j)$ in Equation 7 is then approximated by the similarity between Xi and Xj.

$$\delta(o_i,o_j) \leftarrow \begin{cases} 1, & \text{if } i = j \\ S(X_i,X_j), & \text{if } i \neq j \end{cases}$$

where $S(Xi, Xj)$ is a similarity measure between Xi and Xj, and $0 \leqslant S(Xi, Xj) \leqslant 1$, which derives from dissimilarity metric(Equation 2). Hence, the optimal set of cluster indices can be determined by

$$M^* = \arg\min R'(M) \quad (8)$$

where $R'(M) = \sum_{m=1}^{M} n_{m*}^2 + \Omega - 2\sum_{i=1}^{N}\sum_{j=1}^{N}\delta(h_i,h_j)S(X_i,X_j)$ is the estimated Rand Index.

5. EXPERIMENT

The evaluation of the framework has been carried out in two phases - effectiveness of the dissimilarity metrics, and the clustering algorithm. The test dataset consists of 91 pop songs, which include 21 out of 24 keys. The audio has been collected from CD recordings and contain the singing voice together with musical instrument accompaniment. The files are stored as 44 kHz, 16 bit, mono PCM waveform. Ground truth for the actual key information has been obtained from commercially available sheet music[1].

5.1. Dissimilarity efficiency validation

For convenience, we denote the relationship between music pieces as intra-class, relative-class, parallel-class and inter-class. Musical pieces with the same key are intra-class; share the same Major/Relative Minor combination of keys are relative-class (for example C Major and A Minor); share the same tonic but are in Major and Minor modes are parallel-class (e.g. A Major vs. A Minor), while inter-class means two pieces are in unrelated keys. The Dissimilarity evaluation is carried out using SKL Divergence of chroma spectrum and Euclidian Distance between center of effect, of two music recordings. The results are discussed below:

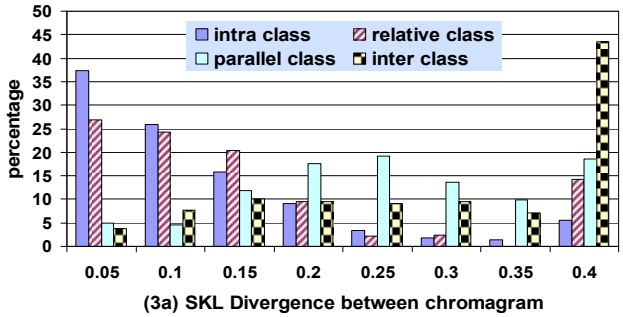
		SKL Divergence of chroma spectrum	Euclidian Distance between Center of Effect
Intra class	**average**	**0.1402**	**0.2917**
	Stat. lower bound[2]	**0.0004**	**0.0193**
	Stat. upper bound	**0.2177**	**0.4419**
Relative class	**average**	**0.1860**	**0.3344**
	Stat. lower bound	**0.0330**	**0.0675**
	Stat. upper bound	**0.4777**	**0.5224**
Parallel class	**average**	**0.2715**	**0.5160**
	Stat. lower bound	**0.0740**	**0.2317**

[1] http://www.musicnotes.com/, Commerical Sheet Music Archive
[2] Stat. lower/upper bound stands for the bound of interval, which contains 80 percentile of the total samples

Inter class	Stat. upper bound	0.4092	0.8670
	average	0.3681	0.5637
	Stat. lower bound	0.1292	0.2087
	Stat. upper bound	1.1042	1.6324

Table 1. Dissimilarity between music with various keys

From Table 1 it is seen that average SKL Divergence for inter-class samples (0.36) and parallel-class samples (0.27) is much higher than that of the intra-class (0.14). This difference can be further demonstrated by Figure 3a, which shows the percentage distribution of SKL Divergence. It is observed that the total percentage of the intra-class distances less than 0.2 is 78%, while 80% of inter-class and parallel-class distances are greater than 0.2. A similar trend is observed for Euclidian Distance between Centre of Effect in Table 1 and Figure 3b.



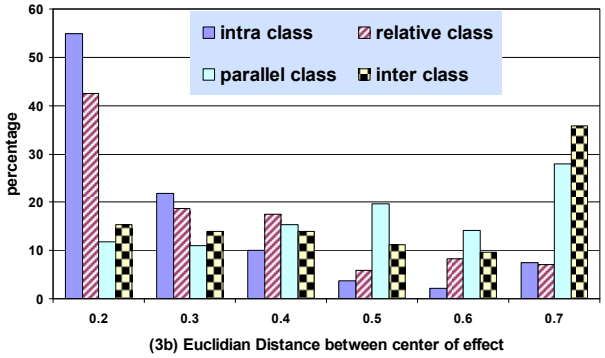


Figure 3. Proportion of dissimilarity metric in various intervals

These results corroborate significant confidence in distinguishing intra-class samples from parallel-class and inter-class samples. However, relative-class distances are observed to be much more difficult to distinguish as the SKL Divergence and Euclidian Distance, are both observed to be fairly close to the intra-distance. This can be explained by the music theoretic knowledge that relative keys share the same scale and similar harmonic structure. Thus, chroma features of relative-class pieces have similar distribution. In addition, modulations between relative modes are common in tonal music, which makes it harder to identity whether a song is primarily structured around a Major or its Relative Minor key (for example, a song with the verse sections in C Major and the Chorus sections in A Minor).

5.2. Clustering performance evaluation

The accuracy of clustering results is evaluated via cluster purity and Rand Index. Figure 4 reveals cluster purity and Rand Index as well as their correlation with the number of clusters. It can be observed that the cluster purity is always above 50%. On the other hand, the Rand index is relatively stable around 50% and reaches minima of 48% when the number of clusters is 21 (the actual number of keys occurring in the test dataset as per ground truth). This follows the discussion in Section 4.2 that Rand Index will reach its minimal value, when the number of cluster is equal to the actual number of keys occurring in the data.

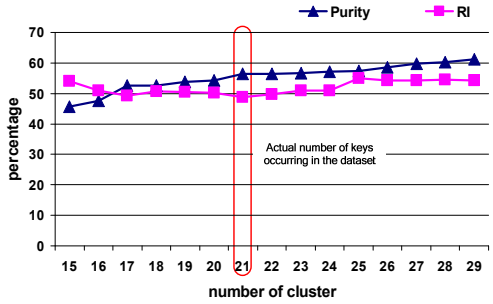


Figure 4. Clustering accuracy evaluation

The number of clusters predicted by our system for the test dataset is observed to be 24 based on a minimum of estimated Rand Index(computed by Equation 8), while the cluster purity is observed to be 57.2%. It can be seen that this is fairly close to the actual number of clusters - 21.

On further analysis of the clustering result, we find that quite a few errors are caused because of the Major/ Relative Minor ambiguity. A straightforward approach to reduce the confusion here would be to merge such keys into key groups[12], which implies that the signature for each cluster is a combination of 2 keys - the Major and its Relative Minor. In our experiments, the cluster purity has been observed to be as high as 70% with this change. Furthermore, errors are also caused because the algorithm is sometimes unable to distinguish the tonic from the dominant (perfect fifth interval). The overlap of harmonic components makes it difficult to identify the chroma component the harmonic belongs to. Besides, perfect fifth is a basic element when construction of triads is concerned in harmony. Similar errors were also observed in [3,12].

6. CONCLUSION

In this paper, a framework has been presented to cluster a higher level feature of music, the key, in an unsupervised way, discarding prior training information and music theoretic based rules. To the best of our knowledge this is the first attempt in this direction and hence there is no strong basis for evaluation against existing techniques. An empirical evaluation shows that accuracy of the existing

rule-based key estimation approach[16] suffers a decrease from over 80% to around 60% as the dataset has been scaled from 30 to 91 songs. From the discussion above, it is observed that the clustering performance is still stable as the data set grows because the dependency on specific higher level musical knowledge is not present. This gives us sufficient confidence that clustering, if involved as a preprocessing component in key detection tasks, will contribute in improving the accuracy for key estimation in large music databases. Future work will focus on integrating the clustering framework with key detection techniques to evaluate performance and scalability. Although the clustering framework is not yet sufficient to output the actual key assignment for each music recording or cluster, we believe it could provide useful information for further structure analysis of musical work, in addition to music retrieval & recommendation systems, and emotion recognition systems.

7. ACKNOWLEDGEMENT

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DETECTION OF STREAM SEGMENTS IN SYMBOLIC MUSICAL DATA

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ABSTRACT

A listener is thought to be able to organise musical notes into groups within musical streams/voices. A *stream segment* is a relatively short coherent sequence of tones that is separated horizontally from co-sounding streams and, vertically from neighbouring musical sequences. This paper presents a novel algorithm that discovers musical stream segments in symbolic musical data. The proposed algorithm makes use of a single set of fundamental auditory principles for the concurrent horizontal and vertical segregation of a given musical texture into stream segments. The algorithm is tested against a small manually-annotated dataset of musical excerpts, and results are analysed; it is shown that the technique is promising.

1. INTRODUCTION

Voice separation algorithms [6, 8, 12, 13, 14, 16, 17, 18] attempt to model computationally the segregation of polyphonic music into separate voices; music segmentation algorithms [1, 4, 5, 7, 15] on the other hand, segment music voices/streams into smaller coherent groups. A common assumption underlying computational models of musical structure is that, firstly voices (or at least the melody) have to be identified and, then, segmentation can be applied to individual voices. For instance, Temperley [17] states that 'once the voices of a texture are identified, the grouping problem becomes more tractable. … polyphonic grouping cannot be done without first grouping the notes into contrapunctal lines.' (pp.81, 83).

In the current paper, the concept of *stream segment* is introduced, i.e. a relatively small number of tones grouped together into a coherent 'whole' perceived independently from other adjacent tones. Such stream segments may be organised into longer streams in case streams are relatively stable for a period of time, or may remain independent structural units. The advantage of adopting the concept of stream segments is that they are meaningful in any type of music, not only when music has a relatively 'fixed' number of voices, as in fugues, choral music, string quartets (see example of stream segments in Figure 1). The proposed algorithm makes use of a single set of auditory principles for the concurrent horizontal and vertical segregation of a musical texture into stream segments.

An algorithm capable of detecting stream segments can be very useful as it enables the organisation of tones into coherent groups that are musically meaningful, allowing thus more efficient and higher quality analytic processing. Most music analytic methodologies (from traditional harmonic analysis to pc-set theory and semiotic analysis) rely on an initial breaking down of the music into relevant (hierarchic) groups/segments/spans (we would say stream segments). In terms of MIR, for instance, a melodic pattern should be identified not spread across different voices or melodic boundaries (perceptally implausible), but within meanigful musical units, or cover-song detection algorithms may be enhanced if they can identify pertinent structural similarities between stream segments (e.g. between melodic segments rather than accompanimental segments).

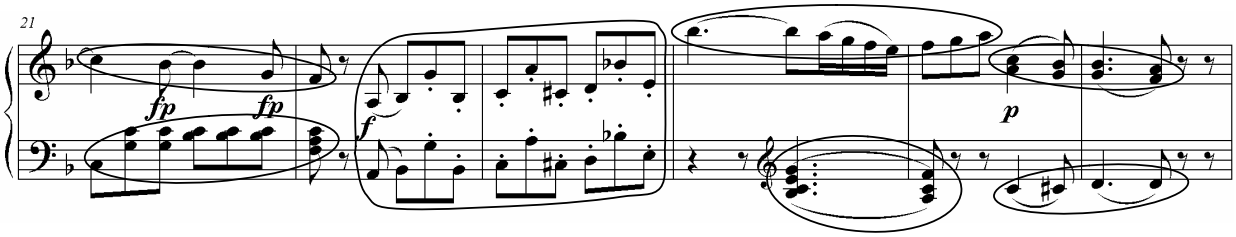

Figure 1 Excerpt from Mozart's Sonata K332, Allegro Assai. Potential stream segments are circled (other options: third segment broken into two pseudopolyphonic voices, and last two segments seen as a single homophonic segment).

2. STREAM SEGMENTS

Most voice separation algorithms such as algorithms [6, 8, 12, 13, 14, 16, 17, 18] model computationally the segregation of polyphonic music into separate voices. Such algorithms commonly assume that 'voice' is a monophonic sequence of successive non-overlapping musical tones. Karydis et al. [12] adopt a perceptual view of musical 'voice' that corresponds to the notion of auditory stream and develop a computational model that automatically splits a musical score into different voices/streams (fewer voices than the maximum number of notes in the greatest chord can be detected manually in [13]).

An underlying assumption of such algorithms is that a fixed number of voices/streams evolve throughout an individual piece of music (sometimes a voice may pause for a certain period and, then, reappear again later - see more on voice and stream in [3]). Most of these algorithms are tested against groundtruth that consists of musical pieces that have a 'fixed' number of voices/streams, such as fugues, choral music, string quartets, songs (melody and accompaniment).

The above assumption, however, is limiting. There exists a significant amount of music that is not composed of a steady number of voices/streams. In many musical works, homophonic, polyphonic, heterophonic elements are mixed together not allowing a listener to trace musical streams throughout the duration of a piece. This fact has led us to hypothesize a novel music theoretic concept, which we will refer to as a *stream segment* (we are not aware of any equivalent music theoretic notion). A stream segment is a relatively short coherent sequence of tones that is separated horizontally from co-sounding streams and, vertically from neighbouring musical sequences.

Grouping relies essentially on fundamental cognitive mechanisms that enable a listener to perceive individual entities as gestalts/wholes [2, 9, 10]. Such mechanisms are based on the principles of proximity and similarity: proximal or similar entities in terms of time, space, pitch, dynamics, timbre to be grouped perceptually together. Such principles are applied locally (e.g. a large pitch interval in-between smaller ones signifies a potential local boundary) or at a higher-level (e.g. a repeating pitch pattern gives rise to a higher level structural unit delimited by pattern boundaries) [4]. In this paper we will discuss segmentation only in relation to the application of gestalt principles at the local level.

In the temporal domain, the most important perceptual principles are the following:

T1 – Synchronous Note Principle: 'Notes with synchronous onsets and same IOIs (durations) tend to be merged into a single sonority.' ([12], p.446)

T2 – Principle of Temporal Continuity: 'Continuous or recurring rather than brief or intermittent sound sources' evoke strong auditory streams ([10], p.12).

In essence, these two principles indicate that concurrent sounds and sounds following closely one another tend to be merged into the same group/stream, and that asynchronous overlapping tones and sounds occurring at a distance from one another tend to be perceived as belonging to different groups.

In the pitch domain, the most important principles are:

P1 – Principle of Tonal Fusion: The perceptual independence of concurrent tones is weakened when they are separated by intervals (in decreasing order: unisons, octaves, perfect fifths…) that promote tonal fusion [10].

P2 – Pitch Co-modulation Principle: 'The perceptual union of concurrent tones is encouraged when pitch motions are positively correlated.' ([10], p.31)

P3 – Pitch Proximity Principle: 'The coherence of an auditory stream is maintained by close pitch proximity in successive tones within the stream.' ([10], p.24)

The first two of these principles apply in the case of synchronous concurrent tones, whereas the third principle applies for successive non-overlapping tones.

By combining the principles we get the following:

a. (T1 & P1 & P2) Synchronous notes tend to be merged; merging is stronger when the notes are separated by intervals that promote tonal fusion and when the pitch motions of concurrent notes are positively correlated,

b. (T2 & P3) Successive notes tend to be grouped together when they follow each other closely and the pitch intervals separating them are relatively small.

In all other cases, groupings are discouraged and potential segmentation boundaries are introduced.

3. THE ALGORITHM

Following the above discussion a stream segment detection algorithm has been developed. The algorithm is based on the *k*-nearest neighbor clustering algorithm, where the distances between notes are computed based on four of the above auditory perceptual organisation principles (only the Tonal Fusion principle is not taken into account at this stage). The algorithm accepts as input a musical piece in symbolic format (quantised piano-roll) and outputs a list of musical stream segments.

The stream segment detection algorithm is illustrated in Figure 3. Procedure `Segment` gets the list of notes `N` of the examined piece, and three parameters, `k` (number of nearest neighbors), `cL` (chain length; see below), and `w` (window length; see below). It first computes the distance matrix `D` between each pair of notes. For the computation of `D`, we examine three cases: (i) If two notes, n_i and n_j, fully coincide, they are checked by a procedure called `verifyCoincide`, which examines whether the vertical structure in the temporal neighborhood of these notes is strong or not (this case examines notes in accordance to the Synchronous Note and the Pitch Co-Modulation principles).

In particular, for each coinciding notes n_i and n_j, `verifyCoincide` forms two separate 'chains' centered at n_i and n_j, which contain preceding and following notes that coincide with each other and have the smallest absolute pitch difference with the center notes. Procedure `verifyCoincide` tries to form chains with length up to `cL`. Figure 2a illustrates an example of two such chains with length 3, centered at notes n_2 and n_5. If two such chains can be formed this means that there exists vertical homophonic structure, as several nearby notes coincide. Additionally, `verifyCoincide` examines the *direction* in the movement of chains, by checking the deviation in the pitch differences between the corresponding chain elements. This way, cases like the one depicted in Figure 2b, where the two chains containing concurrent notes move in non-parallel directions, are distinguished and merging is avoided according to the Pitch Co-modulation Principle.

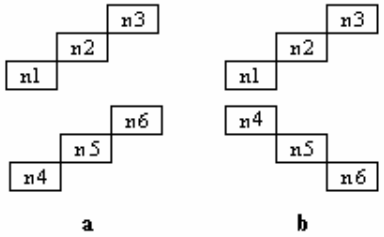

Figure 2 Parallel (a) and non-parallel (b) note chains.

(ii) The second case in the computation of distance matrix `D`, examines asynchronous notes that overlap in time; such notes are considered to have infinite distance. (iii) The final case examines notes that neither overlap nor coincide and their distance is set equal to their normalized pitch difference (`normPD`) plus their normalized IOI (`normIOI`), divided by two (in accordance to the Temporal Continuity and Pitch Proximity Principles). A normalized pitch difference (IOI, respectively) is taken by dividing with the maximum occurring pitch difference (IOI, respectively) between notes occurring within the same time window of length less than `w`.

Next, clustering is performed with procedure `kNNCluster`. This procedure implements the *k*-nearest neighbor clustering algorithm [11], which finds for each note, the list of its k-nearest neighbors (sorted in increasing order of distance). The `kNNCluster` procedure is able to detect clusters of various shapes and sizes. It assigns to each pair of notes n_i and n_j, a mutual neighborhood value (MVN) that is equal to p + q, where p is the position of n_j in the list of n_i and q is the position of n_i in the list of n_j. If n_j or n_i do not belong to the list of each other, then MVN equals infinity. Therefore, if MVN is small, then the corresponding notes are close to each other. For this reason, the procedure joins in the same cluster, notes with MVN = 2, 3, …, 2*k*.

The main characteristic of our stream segment detection algorithm (procedure `Segment`) is that it places a pair of objects (notes) in the same cluster, if they are mutual *k*-nearest neighbors. Due to this characteristic, as will be verified experimentally, this algorithm is able to detect segments of various shapes and sizes, as it is based only on the coherency within clusters. In contrast to several other clustering algorithms, the proposed algorithm does not require that the number of final clusters be determined a priori. This is very suitable for the case of musical stream segment detection as the number of clusters is difficult to be defined correctly in advance. The complexity of the algorithm is $O(N^2)$ for N notes.

```
Procedure Segment(N, k, cL, w)
  foreach ni ∈ N
    foreach nj ∈ N
      if coincide(ni,nj,cL)
        if verifyCoincide(ni,nj)
              D(ni,nj) = 0;
      else if nonoverlap(ni,nj)
      D(ni,nj) = 0.5*(normPD(ni,nj) +
             normIOI(ni,nj));
      else
        D(ni,nj) = ∞;
  kNNCluster(N, D, k);
end;

Procedure kNNCluster(N, D, k)
   foreach ni ∈ N
     L(ni) = kNN(k, N, D);
  foreach ni ∈ N
    foreach nj ∈ N
     if nj ∈ L(ni) and ni ∈ L(nj)
       MVN(ni,nj) = pos(nj,L(ni)) +
        pos(ni,L(nj));
     else
       MVN(ni,nj) = ∞;
   for m = 2 to 2*k
     find all ni,nj∈N with MVN(ni,nj)==m
     assign ni,nj to the same cluster
end;
```

Figure 3 Algorithmic description of the proposed method.

Regarding the tuning of parameters, for several types of musical pieces, like sonatas, mazurkas, waltzes, parameter *cL* takes small values (not more than 2), whereas for others, like fugues, it takes relatively high values (around 10). The reason is that *cL* is responsible for detecting vertical structure. Thus, in musical genres that contain vertical homophonic structures (like sonatas, mazurkas, waltzes), a relatively low value suffices. In contrast, for genres for which this does not hold (polyphonic music like fugues), a higher value is required to avoid false drops.

The tuning of parameter *k* depends on the characteristics of each particular piece, thus has to be tuned accordingly. In our experiments, values within the range 1 to 5 were adequate. In contrast, when considering the stream segment detection algorithm for voice/stream separation (for instance, in Figure 6 the algorithm finds two streams, i.e. melody and accompaniment), higher *k* values should be used. The reason is that, in order to detect entire voices, which can be considered as concatenations of

smaller segments, a higher *k* value enables the algorithm to combine more small segments into larger ones, and eventually to voices/streams. For the detection of smaller, localized segments, small values of *k*, as those mentioned previously, have to be used. Finally, the length of window *w* is set constant to a value equal to half a second, in order to capture local variance of pitch and IOI differences.

4. EXPERIMENTS AND RESULTS

The proposed algorithm has been tested on a small set of musical extracts for piano.[1] The dataset has been annotated by a music theory research student that was instructed to indicate non-overlapping low-level groups of notes that may be perceived as 'wholes' (not to indicate phrases, themes) - the example of figure 1 was discussed before preparing the groundtruth. This small dataset acted as groundtruth for testing the algorithm in this first experiment. Seven extracts were selected from different musical styles: from the openings of Bach's Fugue No.14 in F# major, WTCI, BWV859, Chopin's Mazurkas Op6, No.2 & Op.7, No.5 and Waltz Op.69, No.2 and Beethoven's Sonatas Op.2, No.1, Op.13 (Pathétique) & Op.31, No.3 (overall 1500 notes). The small size of this dataset allowed a more detailed qualitative analysis as well (see discussion below and Figures 4, 5, 6 and 7), in addition to a quantitative analysis.

We studied experimentally the performance of the proposed method (henceforth denoted as kNNClust). For comparison purposes, we also examined a segmentation method that is based on a hierarchical clustering algorithm (henceforth denoted as HierarchicalClust). We considered the following variations of hierarchical clustering: single link, complete link, and Ward's. We found significant differences in the performance of the variations. Thus, we examined all of them, and for each measurement we present the best result among them. Hierarchical methods run on the same distance matrix produced by Procedure Segment, whereas the number of clusters was tuned to give the best results for these methods.

For each music piece and for each segmentation method, we measured accuracy against the groundtruth in terms of the *Jaccard Coefficient*. Let f_{11} denote the number of notes that belong to the same segment in the groundtruth and have been assigned to the same segment by the segmentation method. In the same manner, f_{10} denotes the number of notes that belong to the same segment in the groundtruth but have not been assigned to the same segment by the segmentation method, whereas f_{01} denotes the number of notes that do not belong to the same segment in the groundtruth but have been assigned to the same segment by the segmentation method. The Jaccard Coefficient is computed as the ratio: $f_{11}/(f_{10}+f_{01}+f_{11})$

[1] These pieces were downloaded in Melisma format from the Kern collection (http://kern.humdrum.org)

Segments vary significantly in size, i.e., the number of notes they contain. Jaccard Coefficient tends to penalize excessively incorrect assignments of notes to larger segments and, in contrast, it tends to underestimate errors in smaller segments. To cope for this factor and, thus, to treat equally segments of various sizes, we examined a local Jaccard Coefficient, by computing the f_{11}, f_{10}, and f_{01} numbers only for notes that belong within a time-window. As mentioned, in the proposed method we use a time window to measure normalized pitch and IOI distances. Therefore, we naturally select the length of this window when measuring the local Jaccard Coefficient. Comparison results between kNNClust and HierarchicalClust in terms of the local Jaccard Coefficient are presented in Table 1. Clearly, due to its special characteristics, the proposed method kNNClust compares favorably against HierearchicalClust.

Title	kNNClust	HierClust
Beethoven, Sonata, Op31, No3	0.96	0.82
Beethoven, Sonata, Op2 No1	0.82	0.59
Beethoven, Sonata, Op13, No8	0.84	0.60
Chopin, Mazurka, Op7, No5	0.86	0.62
Chopin, Mazurka, Op6, No2	0.83	0.75
Chopin, Waltz Op.69, No.2	0.94	0.51
Bach, Fugue No14 BWV859	0.84	0.73

Table 1 Accuracy (0 worst, 1 best) by local Jaccard Coefficient.

The algorithm was also tested on a dataset of complete pieces similar to the dataset used as groundtruth in [12]. In this dataset only voices/streams are annotated - not stream segments. A different measure was, therefore, used to evaluate the performance of the algorithm, that rated how well the detected stream segments fall within the voices/streams of the groundtruth, i.e. stream segments are penalised if they are shared across different streams (stream segments are detected for each piece using similar parametric settings as in the previous test). We measured the *voice-crossing penalty* as the ratio of the number of note pairs that were assigned to the same segment whilst they belong to different voices, divided by the number of note pairs that were assigned to the same segment. The voice-crossing penalty is presented in Table 2. In all cases, the measured error is low.

Title	kNNClust
Bach, Fugue No. 7 in E-flat major, BWV 852	0.09
Bach, Fugue No. 11 in F major, BWV 856	0.10
Bach, Fugue No. 1 in C major, BWV 846	0.14
Bach, Fugue No. 14 in F#minor, BWV 859	0.09
Joplin, Harmony Club Waltz	0.02
Chopin, Mazurka in C Major, Op7, No5	0.00
Chopin, Mazurka in C-sharp Minor, Op6, No2	0.02
Chopin, Waltz in D-flat Major, Op. 64, No. 1	0.09

Table 2 Results for voice-crossing penalty (0 best, 1 worst).

The results were examined in detail (qualitative analysis) in order to understand the kinds of mistakes produced by the algorithm. In the example of Figure 4, the algorithm detects coherent groups of notes such as melodic segments, harmonic accompanimental fragments, homophonic passages. Sometimes the algorithm over-segments, e.g. right hand in mm. 1-2, 10-11, 18-19 (these can be seen as single segments due to motivic similarity), or left hand in mm. 8, 18-19 (isolated chords). In m.10, the first note should be grouped with the following notes due to motivic repetition (NB. the current algorithm does not incorporate procedures for detecting parallelism). In the example of Figure 5, it is interesting that the algorithm groups together in stream segments the rhythmically independent notes that are 'sandwiched' in-between the homophonic top and lower notes (note that the algorithm 'mistakenly' does not group the top notes with the lower notes at the end of the passage). In Figure 6, the accompaniment should be segmented at the beginning of m.9 (not beginning of m.8) due to the fact that mm.9-12 are essentially a repetition of mm.5-8 (note that the algorithm does not detect repetitions). There are also mistakes at the edges of stream segments as in the ending of the first segment. Finally, in Figure 7, the algorithm 'correctly' groups together different contrapunctal lines in m.10 (perceptually they may be seen as a single stream), but makes serious mistakes in m.9 where independent voices are merged in the same segments (this type of mistake is under investigation).

5. CONCLUSIONS AND FUTURE WORK

This paper introduces the notion of *stream segment,* i.e., a relatively short coherent sequence of tones that is separated horizontally from co-sounding streams and, vertically from neighbouring musical sequences. A novel algorithm has been proposed that discovers musical stream segments in symbolic musical data based on a set of fundamental auditory principles. The algorithm has been tested against a small manually-annotated dataset of musical excerpts, and the results have been analysed; it was shown that the technique generates promising results.

The proposed concept is new in the sense that it views streaming and segmentation as linked to the same fundamental perceptual principles and that a single algorithm can generate 'stream segments'. Further study, however, is required to make this concept clearer and to provide a crisp definition. Links to appropriate music theoretic notions are important, along with detailed theoretical and empirical analyses of a large number of actual musical examples. Such analyses can form the necessary ground truth on which alrgorithms are tested. Ambiguity, overlapping between stream segments, hierarchic segmentations have to be considered in the future studies of stream segments. We hope that this paper will initiate further research on this topic.

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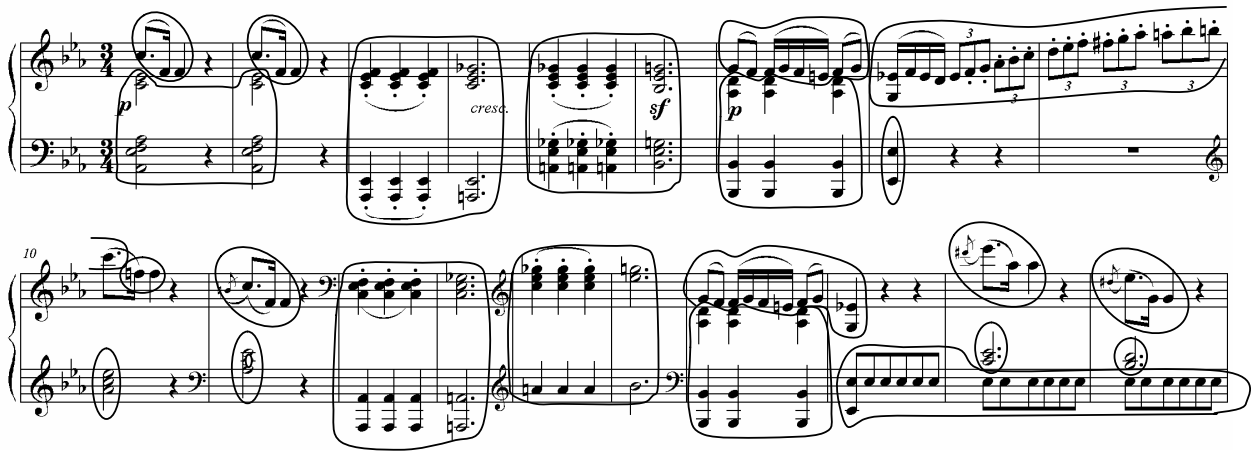

Figure 4 Stream segments detected by algorithm in the opening of Beethoven's Sonata Op.31, No.3 (see text for details).

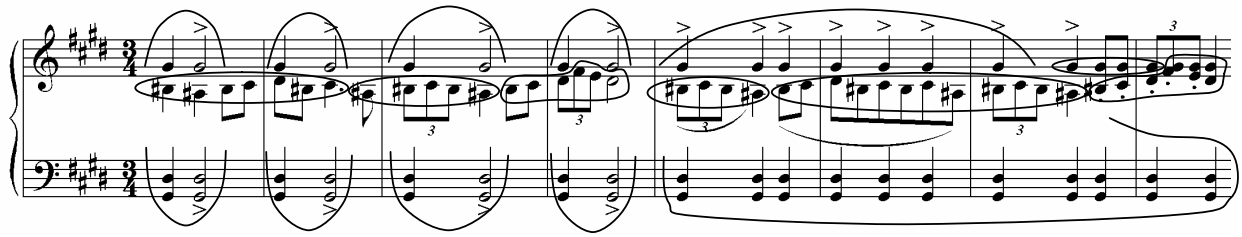

Figure 5 Stream segments detected by algorithm in the opening of Chopin's Mazurka Op.6, No.2 (see text for details)

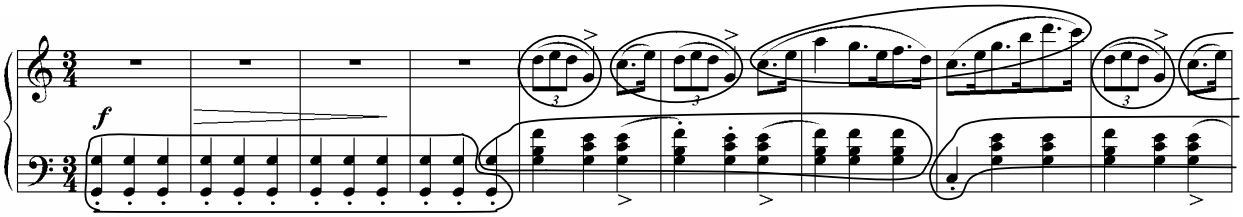

Figure 6 Stream segments detected by algorithm in the opening of Chopin's Mazurka Op.7, No.5 (see text for details)

Figure 7 Stream segments detected by algorithm in the opening of Bach's Fugue in F# major No.14, WTCI, BWV859 (see text for further details)

A MANUAL ANNOTATION METHOD FOR MELODIC SIMILARITY AND THE STUDY OF MELODY FEATURE SETS

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ABSTRACT

This paper describes both a newly developed method for manual annotation for aspects of melodic similarity and its use for evaluating melody features concerning their contribution to perceived similarity. The second issue is also addressed with a computational evaluation method. These approaches are applied to a corpus of folk song melodies. We show that classification of melodies could not be based on single features and that the feature sets from the literature are not sufficient to classify melodies into groups of related melodies. The manual annotations enable us to evaluate various models for melodic similarity.

1 INTRODUCTION

The long term goal of the WITCHCRAFT-project is to create computational methods that support folk song research. [1] This paper takes an essential step towards this goal by investigating the similarity of songs that have been classified by humans into groups of similar melodies.

For the computational modeling of melodic similarity numerous features of melody could be taken into account. However, for a specific problem such as classification only a few features might be sufficient. Hence, we need a means to evaluate which features are important. Once a similarity measure is designed that uses a single feature or a few features, we also need a means to evaluate that similarity measure.

Therefore, we have developed a manual annotation method that gathers expert judgments about the contribution of different musical dimensions to perceived similarity. We use this method to characterize the similarity of selected folk songs from our corpus. The human perception of melodic similarity is a challenging topic in cognition research (see e.g. [1] and [4]). The establishing of the annotation data in this paper is a first step to study the similarity as perceived by humans in the special case of similarity between melodies belonging to the same melody group. We evaluate in how far available computational features contribute to the characterization of similarity between these songs.

[1] http://www.cs.uu.nl/research/projects/witchcraft

Contribution: With these two methods we address the following questions:

1. Is there a small subset of features, or even one single feature, that is discriminative for all melody groups?
2. Is the membership of a melody group based upon the same feature for all member melodies?
3. Are the feature sets provided in earlier research sufficient for classification of the melodies?

1.1 Human classification of melodies

The Meertens Institute in Amsterdam hosts and researches folk songs of the corpus *Onder de groene linde* that have been transmitted through oral tradition. Musicological experts classify these songs into groups called *melody norms* such that each group is considered to consist of melodies that have a common historic origin. Since the actual historic relation between the melodies is not known from documentary evidence, the classification is based on similarity assessments. If the similarity between two melodies is high enough to assume a plausible genetic relation between them, the two melodies are assigned to the same melody norm. In the human process of assigning melody norms some melodies receive the status of a *prototypical* melody of their norms as the most typical representative. All other melody candidates are then compared to this prototypical melody in order to decide whether they belong to this norm.

The classification of melodies into groups of related melodies is a special case of human categorization in music. In order to be able to retrieve melodies belonging to the same melody norm we have to investigate whether all melodies belonging to a melody norm share a set of common features or vary in the number and kind of characteristic features they possess. Two different views of categorization are relevant for this.

The *classical* view on categorization goes back to Aristotle and defines a category as being constituted of all entities that posses a common set of features. In contrast to this, the *modern* view claims that most natural concepts are not well-defined but rather that individual exemplars may vary in the number of characteristic features they possess. The most prominent models according to this view are Wittgenstein's *family resemblance* model (see [10]) and Rosch's *prototype*

model (see [6]). Deliege in [2] and Ziv & Eitan in [11] provide arguments that the family resemblance and the prototype model are most appropriate to describe the categories built in Western classical music.

2 SIMILARITY ANNOTATIONS

The study of melodic similarity in this paper contributes to the development of a search engine for the collection of Dutch folk songs *Onder de groene linde*, which contains both audio data, metadata and paper transcriptions. The test collection employed consists of 1198 encoded songs (MIDI and **kern formats) segmented into phrases. The songs have been classified into melody norms. Three experts annotated four melody norms in detail. For each melody group one expert determined a reference melody that is the most prototypical melody. All other melodies of the group were compared to the reference melody.

The annotation data consists of judgements concerning the contribution of different musical dimensions to the similarity between the melody and the prototype of its melody. In daily practice, the experts mainly perform the similarity evaluation in an intuitive way. In order to analyze this complex and intuitive similarity evaluation, we specified the musical dimensions of the annotations in close collaboration with the experts. These dimensions are rhythm, contour, motifs, form and mode. [2] In order to be used as a ground truth for computational algorithms we standardized the human evaluation such that numeric values are assigned to most of the dimensions. We distinguish three different numeric values 0, 1 and 2: [3]

0. The two melodies are not similar, hence according to this dimension a relation cannot be assumed.
1. The two melodies are somewhat similar, a relation according to this dimension is not implausible.
2. The two melodies are obviously similar, a relation according to this dimension is highly plausible.

For each dimension we defined a number of criteria that the human decision should be based upon when assigning the numeric values. These criteria are as concrete as necessary to enable the musicological experts to give reliable ratings that are in accordance with their intuitive assignments.

2.1 Criteria for the similarity annotations

2.1.1 Rhythm

- If the two songs are notated in the same, or a comparable meter (e.g. 2/4 and 4/4), then count the number of transformations needed to transform the one rhythm into the other (see Figure 1 for an example of a transformation):
  - If the rhythms are exactly the same or contain a perceptually minor transformation: value 2.
  - If one or two perceptually major transformations needed: value 1.
  - If more than two perceptually major transformations needed: value 0.
- If the two songs are not notated in the same, or a comparable meter (e.g. 6/8 and 4/4), then the notion of transformation cannot be applied in a proper manner (it is unclear which durations correspond to each other). The notation in two very different meters indicates that the rhythmic structure is not very similar, hence a value of 2 is not appropriate.
  - If there is a relation between the rhythms to be perceived: value 1.
  - If there is no relation between the rhythms to be perceived: value 0.

Figure 1. Example of a rhythmic transformation: In the first full bar one transformation is needed to transform the rhythm of the upper melody into the rhythm of the lower melody.

In all cases "rhythm" refers to the rhythm of one line. Hence the songs are being compared line-wise.

2.1.2 Contour

The contour is an abstraction of the melody. Hence it remains a subjective decision which notes are considered important for the contour. From the comparison of the lines we cannot automatically deduct the value for the entire melody via the mean value. Therefore we also give a value for the entire melody that is based on fewer points of the melody and hence on a more abstract version of the melody than the line-wise comparison.

- For the line-wise comparison:
  - Determine begin (if the upbeat is perceptually unimportant, choose the first downbeat as begin) and end of the line and 1 or 2 turning points (extreme points) in between.
  - Based on these 3 or 4 points per line determine whether the resulting contour of the lines are very similar (value 2), somewhat similar (value 1) or not similar (value 0).
- For the comparison of the global contour using the entire song:

[2] Text often provides cues as to whether songs are recognized as being genetically related; hence this dimension is annotated too. Form is an auxiliary criterion in order to make reductions possible, such as from ABCC to ABC. Text and form are not discussed here; for their description see [8].

[3] Differentiating more than three values proved to be an inadequate approach for the musicological experts.

– Decide per line: if the pitch stays in nearly the same region choose an average pitch for this line; if not, choose one or two turning points.
– Compare the contour of the entire song consisting of these average pitches and turning points.
– If the melody is too long for this contour to be memorized, then choose fewer turning points that characterize the global movements of the melody.

2.1.3 Motifs

The decision to assign a certain norm to a melody is often based on the detection of single characteristic motifs. Hence it is possible that the two melodies are different on the whole, but they are recognized as being related due to one or more common motifs.

- If at least one very characteristic motif is being recognized: value 2.
- If motifs are shared but they are not very characteristic: value 1.
- No motifs are shared: value 0.

Characteristic in this context means that the motif serves as a basic cue to recognize a relation between the melodies.

2.1.4 Mode

We distinguish two groups of modes based on major vs. minor characteristics. Major/Ionian, Lydian and Mixolydian hence form one group, while Minor/Aeolian, Dorian and Phrygian from another group.

- If the two melodies have exactly the same mode: value 2.
- If the modes of the two melodies are different but belong to the same group: value 1.
- If the modes of the two melodies belong to different groups: value 0.

3 EXPERIMENT ON CREATING ANNOTATIONS

From the set of 1198 encoded melodies 4 melody norms containing 11–16 melodies each have been selected to be annotated by three musicological experts for an initial experiment on the similarity annotation. These are the melody norms *Frankrijk buiten de poorten 1* (short: *Frankrijk*), *Daar was laatst een boerinnetje* (short: *Boerinnetje*), *Daar was laatst een meisje loos 1* (short: *Meisje*) and *Toen ik op Neerlands bergen stond* (short: *Bergen*). For each melody norm one musicological expert determined the reference melody. Similarity ratings were assigned to all other melodies of the same norm with respect to the reference melody. In a first stage of the experiment *Frankrijk* and *Boerinnetje* were annotated, in a second stage *Meisje* and *Bergen*. After the first stage the results were discussed with all experts.

3.1 Agreement among the experts

Table 1 gives an overview of the agreement among the three experts for all musical dimensions using three categories. Category A counts the number of total agreement, i.e. all three experts assigned the same value. Categories PA1 and PA2 count the number of partial agreements such that two experts agreed on one value while the third expert chose a different value. In PA1 the difference between the values equals 1 (e.g. two experts assigned a 1 while one expert assigned a 2). In PA2 the difference between the values equals 2 (e.g. two experts assigned 0 while one expert assigned a 2). Category D counts the cases in which all experts disagree.

Melody Norm	A	PA1	PA2	D
Frankrijk	58.7	38.1	1.6	1.6
Boerinnetje	50.8	42.6	0.5	6.1
Meisje	70.4	27.6	1	1
Bergen	77.5	18.5	1.1	2.9
Average	64.3	31.7	1.1	2.9

Table 1. Comparison of agreement among three experts: A for total agreement, PA1 and PA2 for partial agreement D for disagreement (see section 3.1 for further details). Numbers are percentages.

Both the percentage of disagreement in category D and the percentage of partial agreement PA2 containing both values for *not similar* and *very similar* are quite low. The category of total agreement A comprises the majority of the cases with 64.3%. Moreover, comparing the values obtained for *Frankrijk* and *Boerinnetje* to those for *Meisje* and *Bergen* reveals that the degree of agreement is much higher within the second stage of the experiment after the discussion of the results of the first stage. Hence, this experiment indicates that the musical dimensions have been established in such a way that there is considerable agreement among the musical experts as to how to assign the similarity values.

3.2 Comparing dimensions across melody norms

Table 2 lists the distribution of the assigned values within each musical dimension for all melody norms. In three melody norms the dimension *mode* receives in 100% of the cases the value 2, since all melodies of the norm belong to the same mode. However, mode as an isolated dimension can hardly function as a discriminative variable for the classification of the melodies. In the following we study the values for the other musical dimensions.

Both *Frankrijk* and *Meisje* score highest for rhythm concerning the value 2, while *Boerinnetje* scores highest for motifs and *Bergen* for global contour. Hence the importance of the different musical dimensions regarding the similarity assignment of melodies belonging to one norm varies be-

Melody Norm	*Frankrijk*			*Boerinnetje*			*Meisje*			*Bergen*		
Value	0	1	2	0	1	2	0	1	2	0	1	2
Rhythm	0	1.3	98.7	11.2	51.6	37.2	3.3	8.2	88.5	3.5	15.8	80.7
Global contour	0	31.7	68.3	12.8	48.7	38.5	33.3	13.3	53.4	2.5	10.3	87.2
Contour per line	5.6	52.5	40.9	41.9	26.4	31.7	20.7	31.8	47.5	4.8	22.5	72.7
Motifs	0	36.6	63.4	0	20.5	79.5	13.3	16.7	70	0	17.9	82.1
Mode	13.3	13.3	83.4	0	0	100	0	0	100	0	0	100

Table 2. Distribution of the assigned values within each dimension per melody norm as percentages.

tween the norms. Moreover, in most of the cases single dimensions are not characteristic enough to describe the similarity of the melodies belonging to one melody norm.

The best musical feature (excluding mode) of *Boerinnetje* scores 79% for value 2, the other musical dimensions score below 40%. From this perspective, the melodies of *Boerinnetje* seem to form the least coherent group of all four melody norms. While *Frankrijk* receives the highest rating in a single dimension for value 2, all other dimensions score relatively low. *Bergen* scores in all dimensions above 72% for the value 2. Hence these melodies seem to be considerably similar to the reference melody across all dimensions. For *Meisje* two dimensions receive scores above 70% for value 2, on the other hand three dimensions have considerably high scores (between 13% and 33%) for the value 0. Hence this norm contains melodies with both very similar and very dissimilar aspects.

Comparing the contribution of the musical dimensions reveals that the contour scores for only one melody norm (*Bergen*) above 70% for value 2. Both rhythm and motifs score above 70% for value 2 in three out of four cases. Hence rhythm and motifs seem to be more important than contour for the human perception of similarity in these experiments.

3.3 Similarity within melody norm

As a measurement for the degree of similarity of each melody within the norm to the reference melody we calculated the average over the dimensions rhythm, global contour, contour per line and motifs. The results show that the degree of similarity within the norm can vary to a considerable amount. For instance, in the melody norm *Meisje* two melodies score higher than 95% for value 2, while two melodies score lower than 20% for value 2 with corresponding high scores for value 0.

The evaluation of single dimensions shows that also within these single features the degree of similarity to the reference melody varies. For instance, *Meisje* scores for the dimension rhythm on average 88.5% for value 2. However, one melody scores for rhythm only 42% for value 2 and 33% for value 0. Hence we conclude that there is not one characteristic (or one set of characteristics) that all melodies of a melody norm share with the reference melody.

3.4 Discussion

From sections 3.2 and 3.3 we conclude that both across and within the melody norms the importance of the musical dimensions for perceived similarity varies.

There is not one characteristic (or one set of characteristics) that all melodies of a melody norm share with the reference melody. Therefore, the category type of the melody norms cannot be described according to the classical view on categorization, but rather to the modern view. This agrees with the studies in [2] and [11] on categorization in Western classical music.

4 EVALUATING COMPUTATIONAL FEATURES

This section complements the preceding one by an evaluation of computational features related to melodic similarity.

4.1 Global Features

We evaluate the following three sets of features:

- 12 features provided by Wolfram Steinbeck [7].
- 40 features provided by Barbara Jesser [3].
- 40 rhythm, pitch and melody features implemented in jSymbolic by Cory McKay [5].

The sets of Steinbeck and Jesser were specifically assembled to study groups of folk songs within the Essen Folk Song Collection that are related through the process of oral transmission. Because our corpus consists of folk song melodies, the evaluation of especially these two feature sets is important to get an indication of the value of computational features in general. McKay's set has a general purpose. For the complete list of features we refer to [8].

All features for which absolute pitch is needed (e.g. Steinbeck's Mean Pitch) were removed because not all melodies in our corpus have the same key. Also the multidimensional features from the set of jSymbolic were removed because they are primarily needed to compute other features. Thus we have 92 features, which are characterized as 'global' because for each feature an entire song is represented by only one value.

These features can be considered aspects of the musical dimensions that were chosen for the manual annotations. For example, features like the fraction of descending minor seconds, the size of melodic arcs and the amount of arpeggiation contribute to contour, but they do not represent the holistic phenomenon of contour exhaustively.

4.2 Feature evaluation method

For the four melody norms that were examined in the previous sections, the discriminative power of each individual feature is evaluated. The songs are divided into two groups: one group contains the songs from the melody norm under consideration and the other group all other songs from the test collection. The intersection of the normalized histograms of both groups is taken as a measure for the discriminative power of a feature:

$$I_{mn} = 1 - \frac{\sum_{i=1}^{n} \left| H_{mn}[i] - H_{other}[i] \right|}{\sum_{i=1}^{n} H_{mn}[i]}$$

where $H_{mn}[i]$ is the value for bin i of the histogram of the songs belonging to the melody norm *mn* and H_{other} is the histogram for all other songs. Both histograms have n bins, with the same edges. For the nominal features n is the number of possible values, and for real valued features, $n = 11$, which is the size of the smallest class.

The smaller the intersection, the larger the discriminative power of the feature. The intersection therefore indicates whether a search algorithm that makes use of a certain feature could be successful or not retrieving the songs of the melody norm from the entire corpus.

Normalization of the histograms is needed for the intersection to get comparable values between 0 and 1. Because the four melody norms all have very few melodies compared to the entire corpus, this involves heavy scaling. As a consequence, the intersection value only serves as an indicator for the achievable recall of a retrieval system using the feature. If both $H_{mn}[i] > 0$ and $H_{other}[i] > 0$ the absolute number of songs in $H_{other}[i]$ is almost certainly larger. Therefore, to get an indication of the precision as well, the absolute values of H_{other} should be considered.

4.3 Results

Table 3 lists the best scoring features. For both *Boerinnetje* and *Meisje* none of the features have low values for the intersections. According to the annotation data the similarity of the melodies in these norms to their respective reference melody is less obvious; *Boerinnetje* is the least characteristic of all melody norms, while *Meisje* contains melodies with both very similar and dissimilar aspects.

Feature	I_F	I_B	I_M	I_N
JESdminsecond	**0.068**	0.764	0.445	0.686
STBAmbitus	0.739	0.720	0.622	**0.183**
Range	0.739	0.720	0.622	**0.183**
JESprime	**0.197**	0.575	0.574	0.719
Repeated_Notes	**0.197**	0.575	0.574	0.719
JESmeter	**0.211**	0.540	0.632	0.269

Table 3. I_{mn} for the six best scoring features sorted according to the smallest intersection (in bold) for any of the melody norms *Frankrijk* (F), *Boerinnetje* (B), *Meisje* (M) and *Bergen* (N). The prefixes JES- and STB- mean that the feature is in the set of Jesser or Steinbeck.

We observe that the best feature for *Frankrijk*, JESdminsecond, has quite high values for the other melody norms, which means that it is only discriminative for *Frankrijk*. This feature measures the fraction of melodic intervals that is a descending second. Apparently a large number of descending minor seconds is a distinctive characteristic of *Frankrijk*, but not of the other melody norms. Melodic samples are shown in Figure 1 and the histograms for this feature are shown in Figure 2. While for the normalized histograms the largest bin of $H_{Frankrijk}$ is much larger than the corresponding bin of H_{other}, the absolute values are 7 for H_{other} and 8 for $H_{Frankrijk}$. This means that a retrieval engine using only this feature would achieve a quite low precision.

The annotations suggest that rhythm contributes most to the similarity of the songs in the melody norm *Frankrijk*. Furthermore, the investigation of a set of melody norms using a rhythmic similarity approach in [9] indicates that the melodies of *Frankrijk* are rhythmically more similar to each other than to melodies of other norms. However, none of the rhythmic features of the three sets is discriminative.

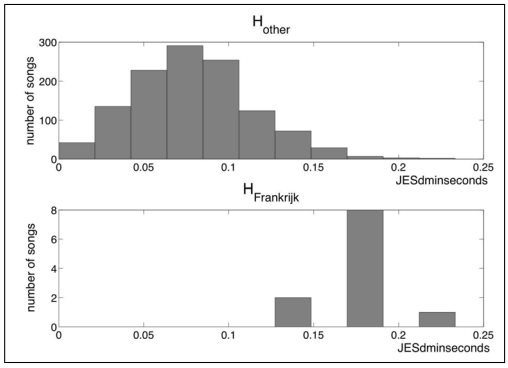

Figure 2. Unnormalized histograms for JESdminseconds for both *Frankrijk* and the other songs.

Most of the lowest values in Table 3 are for *Frankrijk*. STBAmbitus and Range (which are actually the same feature, but from different sets) receive low values for *Bergen*. According to the annotation data, *Bergen* is the only mel-

ody norm with high ratings for both the global contour and the line-wise contour. Range is an aspect of contour. The melodies of *Bergen* typically have a narrow ambitus. For all other features not shown in Table 3, $I_{mn} \geq 0.211$, which indicates that these are not discriminative.

4.4 Discussion

The evaluation of the individual features from the three feature sets shows that there is no single feature in the current set that is discriminative for all four melody norms. Most of the few features that proved discriminative are only so for *Frankrijk*. Therefore, it is not even the case that we find per melody norm a good feature. None of the three sets of features is sufficiently complete for this.

In the manual annotations we observed that motifs are important for recognizing melodies. There are many kinds of motifs: a rhythmic figure, an uncommon interval, a leap, a syncopation, and so on. Therefore it is not possible to grasp the discriminative power of motifs in only a few features. Besides that, global features are not suitable to reflect motifs, which are local phenomena. This is an important shortcoming of the approach based on global features.

It proves difficult to find clear links between the musical dimensions used in the manual annotations and the computational features. The two approaches reside on different levels of abstraction. Computational features have to be computed deterministically. Hence, low level and countable characteristics of melodies are more suited than the more intuitive and implicit concepts that are used by the human mind. Nevertheless, computational features provided complementary insights to the manual annotations, such as the characteristic descending minor second for *Frankrijk*.

5 CONCLUDING REMARKS

With the results of both approaches, we are able to provide answers to the questions stated in the introduction. First, there is no single feature or musical dimension that is discriminative for all melody norms. Second, it is not guaranteed that one single feature or musical dimension is sufficient to explain the similarity of each individual melody to the melody norm. Third, although two of the sets of computational features were specifically assembled for folk song melodies, none of the involved sets provides features that are generally useful for the classification task at hand. A next step would be to evaluate subsets of features instead of individual ones. Although these might prove more discriminative than single features, the importance of the dimension 'motifs' indicates strongly that local model-based features are needed rather than adding more global statistic ones.

The manual annotation of melodic similarity proved a valuable tool to analyze the complex and intuitive similarity assessment of the experts by specifying the constituent parts that contribute to the specific perception of melodic similarity that underlies folksong classification. Therefore a larger set of such annotations is now being created. The annotation data can also be used to evaluate similarity measures that are based on one or more of the musical dimensions.

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ACCESSING MUSIC COLLECTIONS VIA REPRESENTATIVE CLUSTER PROTOTYPES IN A HIERARCHICAL ORGANIZATION SCHEME

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ABSTRACT

This paper addresses the issue of automatically organizing a possibly large music collection for intuitive access. We present an approach to *cluster tracks in a hierarchical manner* and to *automatically find representative pieces of music* for each cluster on each hierarchy level. To this end, audio signal-based features are complemented with features derived via Web content mining in a novel way. Automatic hierarchical clustering is performed using a variant of the *Self-Organizing Map*, which we further modified in order to *create playlists* containing similar tracks.
The proposed approaches for playlist generation on a hierarchically structured music collection and finding prototypical tracks for each cluster are then integrated into the *Traveller's Sound Player*, a mobile audio player application that organizes music in a playlist such that the distances between consecutive tracks are minimal. We extended this player to deal with the hierarchical nature of the playlists generated by the proposed structuring approach.
As for evaluation, we first assess the quality of the clustering method using the measure of entropy on a genre-annotated test set. Second, the goodness of the method to find prototypical tracks for each cluster is investigated in a user study.

1 INTRODUCTION AND MOTIVATION

Increased storage capabilities, high-speed Internet access, and novel techniques for music compression have tremendously increased the size of private as well as commercial music collections over the past few years. Due to the large number of audio tracks in private repositories, it is virtually impossible for users to keep track of every piece. Hence, elaborate methods for organizing large music collections become increasingly important. The enormous economic success of high-capacity mobile music players, such as Apple's "iPod", necessitates intelligent methods for automated structuring of music collections.
The paper at hand presents a possible solution to this problem. The approach proposed here employs an unsupervised hierarchical clustering algorithm, more precisely the *Growing Hierarchical Self-Organizing Map* (GHSOM) [3], on audio features extracted from the music collection. Each cluster of the resulting hierarchical organization is then labeled automatically with a typical track from the cluster to facilitate exploration. To this end, we elaborated an original approach that not only relies on audio features, but incorporates Web-based information on the prototypicality of artists to determine a representative track for each cluster. The main motivation for this is the bad quality of straightforward methods to find a representative for a set of data items. Indeed, choosing for example the music track whose audio feature representation is closest to the average feature vector of the cluster may give a mathematically sound prototype for the cluster. However, in preliminary experiments that employed this selection approach, we encountered many representatives that were either songs by barely known artists (even though the cluster contained very well known artists) or barely known tracks by popular artists. In any case, using such tracks as representative for a cluster seems counterintuitive to guide the user in exploring his music collection.
The main contributions of this work are the novel approach to *determine cluster prototypes*, based not only on the audio features, but also on Web mining techniques, in order to omit "average" cluster representatives that are unknown to most users. In addition, we reimplemented the GHSOM algorithm and modified it in order to *generate playlists*. Furthermore, a novel version of the Traveller's Sound Player application, which uses the cluster prototypes to facilitate navigating the collection, is proposed. The approaches to generate playlists and finding cluster prototypes are further evaluated thoroughly.
The remainder is structured as follows. Section 2 briefly discusses related work. In Section 3, the acquisition of feature data to describe artists and pieces of music is elaborated. Our approaches to hierarchically cluster a given music collection and creating playlists from these clusters can be found in Section 4, whereas finding representative prototypes for each cluster is addressed in Section 5. The experiments conducted to evaluate the proposed approaches are detailed in Section 6. Finally, Section 7 gives some remarks on the integration of the presented approaches in the Traveller's Sound Player application, and in Section 8, conclusions are drawn.

2 RELATED WORK

This paper was mainly motivated by the work done in [9], where Pohle et al. employ different heuristics to solve a Traveling Salesman Problem on a track-level audio similarity matrix. The resulting tour represents a playlist with minimized distances between consecutive pieces of music. The playlist generated for a complete music collection is then made accessible via a special music player. In [8], this approach is modified in that the audio similarity matrix is adapted according to artist similarity estimations, which are derived from $TF \times IDF$ features computed on artist-related Web pages. In contrast to [9, 8], we do not model a Traveling Salesman Problem to generate playlists. Instead we follow a recursive approach that builds upon the GHSOM algorithm to hierarchically break down the tracks of a music collection until every track is represented by a sole map unit. The playlist is then generated by successively visiting each such unit.

As we build our playlist generation approach upon a variant of the GHSOM, a hierarchical version of the Self-Organizing Map, this work is certainly also related to [3, 4], where the GHSOM is presented.

3 FEATURE EXTRACTION

As we combine features extracted from the audio signal with cultural features from the Web in order to determine prototypical pieces of music for each cluster, we perform the following two feature extraction stages.

3.1 Audio-based Features

First, we extract audio features using a well-established approach. For each piece of music, Mel Frequency Cepstral Coefficients (MFCCs) are extracted on short-time audio segments to obtain a coarse description of the spectral shape. Details on this procedure can be found, for example, in [2, 1, 7]. Gaussian Mixture Models (GMMs) are then used to aggregate the MFCCs of each individual piece of music. The similarity between two arbitrary pieces is calculated as the inverse of the distance between their GMMs. For our experiments, we used parameters similar to those proposed in [1] (19 MFCCs, 30 clusters per GMM). Performing this procedure for a collection of n music pieces eventually yields an $n \times n$ distance matrix.

3.2 Web-based Features

In order to estimate the prototypicality of each artist in the collection, and thus his or her suitability to serve as representative for a cluster, we make use of information found on artist-related Web pages. To this end, we follow the approach *backlink/forward link ratio with exorbitant popularity penalization* as proposed in [11, 12]. The basic assumption underlying this approach is that the probability that a popular artist is mentioned on a Web page about a rather unknown artist is higher than vice versa. This seems reasonable if one thinks, for example, of references of the form "Band X sounds like band Y.". Such phrases are more likely to occur on a Web page of a rather unknown band X, which points to a popular band Y than vice versa.

For each artist, we retrieve up to 150 Web pages returned by Google for queries of the form "`artist name`" `+music+review`. This query scheme was originally proposed in [13] for retrieving artist-related Web pages. Subsequently, the Web pages of each artist are analyzed for occurrences of all other artist names in the collection, and the respective document frequency $df_{a,B}$ (i.e., the number of Web pages in the set B of retrieved pages for artist b on which the name of artist a occur) is stored for all combinations of artists.

To estimate the prototypicality of an artist a for a specific set of tracks given by cluster u [1] , we compare the document frequencies $df_{a,B}$ and $df_{b,A}$ for all b's in the same cluster u as a. We count for how many of these b's the inequality $df_{a,B} \geq df_{b,A}$ holds. The higher this count, the more prototypical artist a. Performing this calculation for all artists of a specific cluster u, we obtain an artist prototypicality ranking $r^u(a)$. However, this ranking is usually quite distorted as artist names like "Genius", "Kiss", or "Bush" are always overrated. Addressing this issue, a penalization term that downranks artists whose prototypicality is extremely high for all clusters (which actually cannot be the case) is introduced. This eventually gives a corrected ranking function $r^u_w(a)$. For a more detailed elaboration on the prototypicality ranking function, please see [12].

4 ADAPTING THE GHSOM FOR PLAYLIST GENERATION

Structuring the music collection under consideration is performed using an unsupervised neural network algorithm that automatically organizes a given data set in a hierarchical manner. More precisely, we reimplemented in Java a hierarchical extension of the Self-Organizing Map, namely the *Growing Hierarchical Self-Organizing Map* (GHSOM), as proposed in [3]. The GHSOM has been integrated in the CoMIRVA framework [10].

The GHSOM extends the widely used Self-Organizing Map [6] in a way that during training, new map units are automatically inserted into the existing grid between the map unit whose data items show the highest deviation from the map unit's model vector and its most dissimilar neighboring unit. This is performed until the mean quantization error of all map units is reduced to a certain threshold τ_1. Hence, τ_1 controls the final size of the individual SOMs. A second threshold τ_2 determines the quantization error a particular map unit must exceed in order to reorganize its data items on

[1] In our case, the clusters are given by the map units of the Self-Organizing Map that we use to organize the collection (cf. Section 4).

a newly generated SOM at a deeper hierarchy level. A more detailed explanation of the GHSOM algorithm is given, for example, in [3].

In order to suit our need for playlist generation, we had to modify the original GHSOM algorithm in two regards. First, we redefined the neighborhood function used in the training process. In particular, we defined a *cyclic neighborhood relation* that extends its range over the map's borders, i.e., considers the last row/column of the map whenever a map unit in the first row/column is modified and vice versa. Second, each time a map unit u is expanded to a new SOM at a deeper hierarchy level, this new SOM is initialized using the model vectors of u's neighboring units.

These two modifications allow for a very simple playlist generation approach in that we just have to train a *1-dimensional GHSOM* on the input data and ensure the recursive expansion of map units until each track is represented by its sole unit. Traversing the resulting GHSOM tree in preorder while successively storing all tracks represented by the leave nodes eventually yields a playlist containing all pieces of the collection. As for training the GHSOM, we use the audio similarity matrix, whose creation is described in Subsection 3.1. For performance reasons, however, we apply *Principal Components Analysis* (PCA) [5] to reduce the dimensionality of the feature space to 30. [2]

5 FINDING CLUSTER PROTOTYPES

One of the main goals of this work is determining prototypical tracks that can serve as representatives to describe each cluster, i.e., map unit, of the GHSOM. However, preliminary experiments showed that relying solely on audio-based information to calculate a representative piece of music for each cluster yields unsatisfactory results, as already explained in Section 1. To alleviate this problem, we incorporate artist prototypicality information extracted as described in Subsection 3.2 into an audio-based prototypicality ranking. The resulting ranking function thus combines audio features of all tracks in the cluster under consideration with Web-based artist prototypicality information as detailed in the following.

Denoting the map unit under consideration as u, the ranking function $r^u(p)$ that assigns a prototypicality estimation to the representation in feature space of each [3] piece of music p by artist a is calculated as indicated in Equation 1, where $r_s^u(p)$ denotes the audio signal-based part of the ranking function (cf. Equation 2) and $r_w^u(a)$ is the corrected Web-based ranking function from Subsection 3.2. In the signal-based ranking function, $\overline{u}$ denotes the mean of the feature vectors represented by u, $|\cdot|$ is the Euclidean distance, and $norm(\cdot)$ is a normalization function that shifts the range to $[1, 2]$. Finally, each piece of music p represented by map unit u is assigned a prototypicality value, which is the higher, the more prototypical p is for u.

$$r^u(p) = r_s^u(p) \cdot r_w^u(a) \tag{1}$$

$$r_s^u(p) = norm\left(\frac{1}{1 + \ln(1 + |p - \overline{u}|)}\right) \tag{2}$$

6 EVALUATION

We conducted two sets of evaluation experiments. The first aimed at comparing our GHSOM-based playlist generation approach to the ones proposed in [9, 8]. In accordance with [9], the quality of the playlist was evaluated by calculating the genre entropy for a number of consecutive tracks. Second, we assessed the quality of our approach to detecting prototypical tracks via a user study.

For evaluation, we used the same collection as used in [8]. This collection contains 2,545 tracks by 103 artists, grouped in 13 genres. For details on the genre distribution, please consider [8].

6.1 Entropy of the Playlist

To evaluate the coherence of the playlists generated by our GHSOM-based approach, we compute the entropy of the genre distribution on sequences of the playlists. More precisely, using each piece of the collection as starting point, we count how many of n consecutive tracks belong to each genre. The result is then normalized and interpreted as a probability distribution, on which the Shannon entropy is calculated, according to Equation 3, where $\log_2 p(x) = 0$ if $p(x) = 0$.

$$H(x) = -\sum_x p(x) \log_2 p(x) \tag{3}$$

In Figure 1, the entropy values for $n = 2 \ldots 318$ are given (plot "audio ghsom"), averaged over the whole playlist, i.e., each track of the playlist is chosen once as the starting track for a sequence of length n. [4]

Comparing the entropy values given by the GHSOM approach to the ones presented in [8] reveals that the GHSOM approach performs very similar to the SOM-based approach followed in [8]. This is obviously no surprise, but shows that the automated splitting of clusters by the GHSOM does not harm the entropy of the generated playlists, while at the same time offers the additional benefit of automatically structuring the created playlists in a hierarchical manner. Like those created by the recursive SOM approach in [8], the GHSOM-based playlists show quite good long-term entropy values (for large values of n), but rather poor values for short playlists.

[2] 30 dimensions seemed appropriate for our test collection of 2,545 pieces of music. Employing the approach on larger collections, most probably requires increasing this number.

[3] More precisely, "each" refers to those pieces of music that are represented by the map unit u under consideration.

[4] 318 tracks correspond to an angular extent of 45 degrees on the circular track selector in the Traveller's Sound Player application.

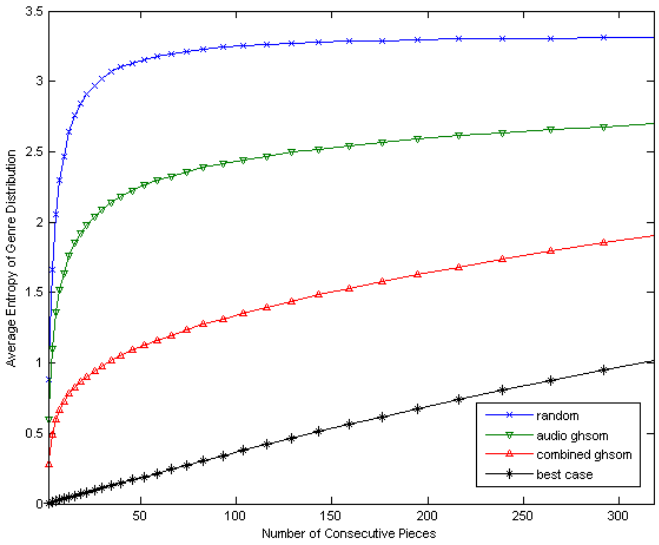

Figure 1. Genre entropy values for n consecutive tracks of the GHSOM-based playlist.

We also experimented with incorporating Web-based features directly into the playlist generation algorithm, as proposed in [8]. This improved on the one hand the entropy values (cf. plot "combined ghsom" in Figure 1). On the other hand, taking a critical look at the resulting playlists showed that using Web-based artist features as proposed in [8] yields "too perfect" playlists in that often one and the same artist contributes 20 or more consecutive tracks. Hence, this approach may be suited well for creating very consistent, but sometimes boring, playlists.

6.2 Quality of the Prototype Estimation

To evaluate the quality of the cluster representatives as determined by the approach proposed in Section 5, we performed a user study. The evaluation sets for the user study were created by training a 1-dimensional GHSOM with an initial number of 5 rows and gradient initialization. The growing threshold τ_1 was set to 0.5, the expansion threshold τ_2 to 0.1. We performed sequential training with a training length of 5 epochs. This setting resulted in a GHSOM with 11 clusters on the first level. [5]

We had 16 participants in the study, each of which evaluated 11 sets of tracks, consisting of 10 tracks each. 10 out of these 11 sets were constructed as follows. The most prototypical track of each cluster, according to our approach, was added to the corresponding set. The remaining 9 tracks were selected by drawing a random sample from their respective cluster. The 11^{th} set (for cluster 6) in contrast contained exactly the same pieces of music for all participants. [6]. The reason for this different creation method for cluster 6 is that we also wanted to take a qualitative look at some of the results, which would have been barely feasible for a large number of differing track sets, each of which is evaluated by only one user. We followed this rather complicated procedure of track set generation in order to raise the total number of evaluated sets. This procedure yielded 161 different track sets (16 participants · 10 sets + 1 invariant set for all participants). Each participant was instructed to select one track from each set that should be best suited to represent the corresponding set. We offered the participants the artist and song names as well as the audio files in case they were uncertain or unfamiliar with artist or track. In total, we received 176 prototype selections, which were analyzed as follows.

To obtain a general performance measure, we first calculated the overall concordance between the prototypes found by our automated approach and those indicated by the participants. Table 1 gives these overall concordance values over the 176 evaluated track sets. We assessed the concordances on different levels. The first row depicts the exact concordance of the tracks, i.e., in order to be considered concordant, the track given by our approach must be the same as the one selected by the user. This definition of concordance is weakened for rows two and three as our approach and the user judgment must, in this case, agree only on the artist or on the genre, respectively.

At first glance, these results may not seem very promising. However, taking the difficult nature of the user task into account (choosing between 10 probably quite similar tracks, many of which are by the same artists), it must be regarded a success that the concordance values considerably exceeded the baseline. The difficulty of the task is also reflected by the relatively high mean genre entropy of 1.95 over all track sets. This number is obviously higher than the entropy value given in Figure 1 for $n = 10$ because of the randomized selection process involved in the creation of the track sets.

Taking a closer look at the individual track sets, we often encountered the problem that the sets contained approximately equal numbers of tracks from two or three different genres. To analyze the influence this may have had on the evaluation results, Table 2 shows, for each group of sets (created from clusters 1 to 11 of the GHSOM), the absolute number of users that agreed with the prototype found by our approach (column "hits") and the mean genre entropy of the respective track set. Note that the entries are ordered by their average entropy. The baseline is 1.6 as each group was evaluated by 16 participants; a random guesser would have yielded a hit rate of 10% in the long run. From Table 2, we can see that high entropy values in fact tend to correspond to low hit rates and vice versa. The low hit rate of 2 for the group with the lowest entropy, however, can be explained by the fact

[5] Level 1 is the first distinctive level of the GHSOM as level 0 contains all data items.

[6] Cluster 6 also contained the prototype and a random sample of 9 other tracks, but this sample remained constant for all participants

level	baseline	concordance
track	10.00	17.61
artist	15.17	26.14
genre	35.34	50.57

Table 1. Overall concordance of the prototype finding algorithm and the user judgments, in percent.

that this group subsumed music by different, but all famous, metal bands. Furthermore, in most track sets of this group, 2 or 3 tracks by "Cannibal Corpse" were included, one of which was also rated the prototype by our approach. But the others were usually no less famous tracks by this band.

The identical composition of cluster 6 for all participants allowed us to take a qualitative look at the ranking results on the level of individual tracks. Table 3 summarizes these results. In this table, the evaluated tracks of cluster 6 are displayed in decreasing order of their prototypicality as determined by our approach. The leftmost column gives the number of times the respective track was selected as cluster representative by a participant. As for the composition of cluster 6, it mainly contained punk rock and folk rock songs, but also a metal song by "Pantera", which somehow also fits into this cluster, and a real outlier by the German electronic music producer "Scooter". From Table 3, it can be seen that about 56% (9 out of 16) of the user hits account for the top 3 rankings by our approach. On the other hand, also the lowest ranked two tracks by "Bad Religion" were chosen by 25% of the users. This is no surprise as this band may be seen as equally prototypical for the punk rock genre as the top ranked band "Offspring". The remaining 3 user hits were given to the tracks by "Subway to Sally". Since the randomly drawn sample included 3 tracks of this band in the evaluation set, it seems that some users considered the set best represented by a track by this artist. None of the participants, however, selected one of the outliers by "Pantera" or "Scooter" as representative.

To summarize the results, even though they certainly leave enough room for improvement, most users at least roughly agreed with the prototypicality ratings given by the proposed approach. Moreover, taking into account the especially difficult task the participants in the user study had to perform and the quite heavy mixtures of different genres in the track sets, the results are surprisingly good.

7 A NOVEL VERSION OF THE TRAVELLER'S SOUND PLAYER

The basic idea of the *Traveller's Sound Player* (TSP) presented in [9] is to arrange the tracks in a music collection around a wheel that serves as track selector. The authors of [9] aim at performing this arrangement in a way such that consecutive tracks are maximally similar. To this end, a large circular playlist is created by solving a Traveling Salesman Problem on an audio similarity matrix. A drawback of the original version of the TSP is, however, that it does not guide the user in finding a particular style of music he or she is interested in. Indeed, the user has to explore different regions of the playlist by randomly selecting different angular positions with the wheel. It is also very hard to exactly position the wheel at a specific track as thousands of tracks are placed around one wheel, which corresponds to an angular representation in the magnitude of tenths of one degree for each track.

Addressing these shortcomings, we propose to integrate the prototype information found by our GHSOM-based approach presented in Section 5 into the TSP. To this end, we elaborated an extension to the TSP that allows for hierarchically exploring the music collection by means of the cluster prototypes. More precisely, the novel user interface maps the structure of the collection, as given by the trained GHSOM, to the wheel. It thus reflects the hierarchical organization in that the user is first only presented the prototypes of the first level clusters. He or she can then either opt for playing the representative track or for descending to a lower hierarchy level. On each level, the user is presented a playlist containing the most prototypical tracks of all clusters (be them either single map units or other GHSOMs at a deeper level) that make up the GHSOM at the current level.

group	hits	entropy
4	2	0.9609
5	7	1.3385
6	**4**	**1.6855**
3	4	1.7516
2	2	2.0189
7	5	2.1068
8	2	2.1785
1	3	2.2763
0	2	2.2772
9	0	2.4343
10	0	2.5372

Table 2. Number of correctly found prototypes and mean entropy values for each of the 11 playlists.

hits	ranking	track name
4	0	offspring - hypodermic
2	1	offspring - leave it behind
3	2	blink 182 - don't leave me
0	3	pantera - strength beyond strength
0	4	scooter - back in the u.k.
2	5	subway to sally - mephisto
1	6	subway to sally - ohne liebe
0	7	subway to sally - böses erwachen
3	8	bad religion - let them eat war
1	9	bad religion - markovian process

Table 3. Detailed results of the user study for group 6, i.e., the group containing identical tracks for all participants.

8 CONCLUSIONS AND FUTURE WORK

We presented an approach to automatically cluster the tracks in a music collection in a hierarchical fashion with the aim of generating playlists. Each track is represented by similarities calculated on MFCC features, which are used as input to the clustering algorithm. To this end, we reimplemented and modified the Growing Hierarchical Self-Organizing Map (GHSOM). We further proposed an approach to determine, for each of the clusters given by the GHSOM, the most representative track. This is achieved by combining the MFCC-based audio features with artist-related information extracted from the Web. We finally integrated the two approaches into a music player software called the "Traveller's Sound Player", for which we elaborated an extension to deal with the hierarchical data.

We carried out evaluation experiments in two ways. First, we aimed at assessing the coherence of the playlists generated by the GHSOM approach by measuring their genre entropy. We compared these entropy values to those given by other playlist generation approaches and found that they largely correspond to a similar approach based on standard SOMs. In addition, our approach also yields a hierarchically organization of the collection and determines a representative for each cluster, while at the same time the good long-term entropy values of the (GH)SOM-based playlist generation are retained. Second, we conducted a user study to assess the quality of the found cluster prototypes. Even though the results are still improvable, we could exceed the baseline considerably. Especially when taking into account the challenging setup of the user study, the results are promising.

As for future work, we are thinking of integrating the artist distribution of the clusters into the ranking function, as the user study showed that users tend to prefer representative tracks by artists who occur often in the cluster under consideration. Similar considerations would probably also be suited to filter out outliers. Furthermore, we have to experiment with different user interfaces for hierarchically accessing music collections as the integration of the cluster representatives into the Traveller's Sound Player is currently done in a quite experimental fashion. Our special concern in this context will be the usability of the user interface for mobile devices.

9 ACKNOWLEDGMENTS

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USING BASS-LINE FEATURES FOR CONTENT-BASED MIR

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ABSTRACT

We propose new audio features that can be extracted from bass lines. Most previous studies on content-based music information retrieval (MIR) used low-level features such as the mel-frequency cepstral coefficients and spectral centroid. Musical similarity based on these features works well to some extent but has a limit to capture fine musical characteristics. Because bass lines play important roles in both harmonic and rhythmic aspects and have a different style for each music genre, our bass-line features are expected to improve the similarity measure and classification accuracy. Furthermore, it is possible to achieve a similarity measure that enhances the bass-line characteristics by weighting the bass-line and other features. Results for applying our features to automatic genre classification and music collection visualization showed that our features improved genre classification accuracy and did achieve a similarity measure that enhances bass-line characteristics.

1 INTRODUCTION

Content-based music information retrieval (MIR) is expected to be a key technology for developing sophisticated MIR systems. One of the main issues in content-based MIR is the design of music features to be extracted from music data. The effectiveness of various features has been examined by several researchers. For audio data, Tzanetakis et al., for example, used spectral shape features (e.g., the spectral centroid, rolloff, flux, and mel-frequency cepstral coefficients (MFCCs)), rhythm content features (e.g., the beat histogram), and pitch content features [1]. Pampalk used features as zero crossing rates, mel spectral features, and fluctuation patterns (the amplitude modulation of the loudness per frequency band) [2]. Aucouturier et al. used MFCCs with various pre- and post-processing techniques [3]. Berenzweig et al. also used MFCCs [4]. Some researchers used chroma features, also known as pitch-class profiles, in addition to or instead of MFCCs [5, 6].

Most studies handling audio data have used low-level features, as described above. Low-level features such as the MFCCs and spectral centroid can be easily calculated and can capture coarse characteristics of musical content to some extent, but they have a clear limit to their ability to capture fine characteristics of musical content; it is not clear, for example, which musical aspects, such as chord voicing and instrumentation, affect the similarity of two vectors of MFCCs. The limit of low-level features was also pointed out by Pampalk [2]. Hence, trying to discover new features beyond such low-level features is a common theme in effort to improve content-based MIR.

We focus on bass lines to design new features. Bass parts play an important role for two (rhythm and harmony) of the three basic elements of music. The base lines of each music genre proceed in their own style. Moreover, a method for extracting the pitch of bass lines has been proposed [7].

In this paper, therefore, we propose new audio features that can be extracted from bass lines and describe applications of them to automatic genre classification and music collection visualization. Section 2 discusses the characteristics of bass parts and the design of audio features to be extracted from bass lines. Section 3 describes an algorithm for extracting the features. Section 3 also describes the other conventional low-level features used in the experiments. Section 4 examines the effectiveness of the proposed features in automatic genre classification tasks. Section 5 describes an application of the proposed features to music collection visualization using Music Islands [2]. We generate different views of Music Islands by switching the weights given to the features. The differences are discussed.

2 DESIGNING BASS-LINE FEATURES

2.1 Characteristics of Bass Lines

Bass lines are contained in almost all genres of music and play important roles in both harmonic and rhythmic aspects. Bass lines usually emphasize the chord tones of each chord, while they work along with the drum part and the other rhythm instruments to create a clear rhythmic pulse [8].

Bass lines have a different style for each genre. In popular music, for example, bass lines often use "riffs", which are usually simple, appealing musical motifs or phrases that are repeated, throughout the musical piece [8]. In jazz music, a sequence of equal-value (usually quarter) notes, called a *walking bass* line, with a melodic shape that alternately rises and falls in pitch over several bars, is used [9]. In rock music, a simple sequence of eighth root notes is frequently used. In dance music, a repetition of a simple phrase that sometimes involves octave pitch motions is used.

2.2 Basic Policy for Designing Bass-line Features

As described above, bass lines for each genre have a different style in both pitch and rhythm. However, we design only pitch-related features due to technical limitations. We

use the PreFEst-core [7] for extracting bass lines. It detects the pitch of the bass part, specifically the most predominant pitch in a low-pitch range, for each frame. Because it does not contain the process of sequential (temporal) grouping, the rhythm of the bass line may not be recognized.

Our pitch-related bass-line features are divided into two kinds: features of pitch variability and those of pitch motions. We design as many features as possible because we reduce the dimensionality after the feature extraction. It is known that pitch estimation techniques sometimes generate octave errors (double-pitch or half-pitch errors). We therefore extract features not only from specific pitches but also from note names (pitch classes) to avoid the influence of octave pitch-estimation errors.

2.3 Features of Pitch Variability

The following 11 features are used:

- Fv1: Number of different pitches that appear in at least one frame in the musical piece.
- Fv2: Number of pitches from Fv1 excluding those with an appearance rate of less than 10% or 20%.
- Fv3: Temporal mean of the numbers of different pitches within a sliding short-term (2 s in the current implementation) window.
- Fv4: Percentage of appearance frequencies of the i most frequently appearing pitches ($i = 1, \cdots 5$).
- Fv5: Pitch interval between the two most frequently appearing pitches.
- Fv6: Period of the most frequently appearing pitch.

Fv1 was designed to distinguish rock music, which tends to use a comparatively small number of notes in bass lines, and jazz music, which tends to use passing notes frequently. Fv2 was prepared to improve the robustness to pitch misestimation. Fv3 was designed to distinguish electronic music, which tends to repeat short phrases moving in pitch, from rock music, which tends to play root notes in bass lines. Fv4 will have high values when the same short phrase is simply repeated throughout the piece. Such repetetion is sometimes used in some types of music such as dance music. Fv5 and Fv6 were designed by referring to the pitch content features of Tzanetakis et al. [1].

2.4 Features of Pitch Motions

The following 10 features are used:

- Ft1: Mean of the numbers of pitch motions (the changes of the bass pitches) per unit time.
- Ft2: Percentage of each of the following kinds of pitch motions: chromatic, conjunct (i.e., either choromatic or diatonic), disjunct (i.e., leaps), and octave.
- Ft3: Percentage of each of the following kinds of successive pitch motions: conjunct+conjunct, conjunct+disjunct, disjunct+conjunct, and disjunct+disjunct.

Similarly to Fv1, Fv2, and Fv3, we designed Ft1 to distinguish jazz music, which tends to involve frequent pitch motions, from rock music, which tends to maintain the same note (usually the root note) within a chord. Ft2 and Ft3 were intended to distinguish walking bass lines from bass riffs involving octave motions used in electronic music.

3 ALGORITHM OF BASS-LINE FEATURE EXTRACTION

Our bass-line features and conventional low-level features are extracted through the following steps.

3.1 Extracting Pitch Trajectory of Bass line

Given an audio signal, the spectrogram is first calculated. The short-time Fourier transform shifted by 10 ms (441 points at 44.1-kHz sampling) with an 8,192-point Hamming window is used. The frequency range is then limited by using a bandpass filter to deemphasize non-bass-part features. The same filter setting as that used by Goto [7] is used.

After that, PreFEst-core [7], a multi-pitch analysis technique, is used. PreFEst-core calculates the relative predominance of a harmonic structure with every pitch in every frame. The most highly predominant pitch for each frame is basically extracted as the result of pitch estimation, but the second most highly predominant pitch is extracted if it is the same as the top one in the previous frame.

3.2 Extracting Bass-line Features

The 21 bass-line features defined in Section 2 are extracted from the pitch trajectory of the bass line. In addition, the same features are extracted from the note name trajectory (i.e., the octave is ignored) to avoid the influence of octave pitch-estimation errors; however, the percentage of octave pitch motions is excluded. The total number of bass-line features is 41.

3.3 Extracting Timbral Features

The timbral features used by Lu et al. [10] are used. The features are listed in Table 1. These features are extracted from the power spectrum in every frame and their temporal means and variances are then calculated.

3.4 Extracting Rhythmic Features

We designed rhythmic features by referring to previous studies, including that of Tzanetakis et al. [1]. For each 3-s window, auto-correlation is analyzed. The following features are then extracted:

Table 1. Timbral features used in our experiments

Intensity	Sum of intensities for all frequency bins
Sub-band intensity	Intensity of each sub-band (7 sub-bands were prepared)
Spectral centroid	Centroid of the short-time amplitude spectrum
Spectral rolloff	85th percentile of the spectral distribution
Spectral flux	2-norm distance of the frame-to-frame spectral amplitude difference)
Bandwidth	amplitude weighted average of the differences between the spectral components and the centroid
Sub-band peak	Average of the percent of the largest amplitude values in the spectrum of each sub-band
Sub-band valley	Average of the percent of the lowest amplitude values in the spectrum of each sub-band
Sub-band contrast	Difference between Peak and Valley in each sub-band

- Fr1: Ratio of the power of the highest peak to the total sum.
- Fr2: Ratio of the power of the second-highest peak to the total sum.
- Fr3: Ratio of Fr1 and Fr2.
- Fr4: Period of the first peak in BPM.
- Fr5: Period of the second peak in BPM.
- Fr6: Total sum of the power for all frames in the window.

3.5 Dimensionality Reduction

The dimensionality of the feature space is reduced to avoid the so-called curse of dimensionality. The dimensionality reduction is performed through the following three steps:

1. For every feature pair, if its correlation is higher than a threshold r, the feature that has a lower separation capacity is removed. The separation capacity is calculated as the ratio of the between-class variance to the within-class variance.
2. The s_1 features having the highest separation capacities are chosen.
3. The dimensionality is further reduced from s_1 to s_2 by using principal component analysis (PCA).

The parameters r, s_1, s_2 are determined experimentally.

4 APPLICATION TO GENRE CLASSIFICATION

In this section, we mention an application of our bass-line features to automatic genre classification. Automatic genre classification [1, 11] is a representative task in the content-based MIR field because genre labels can describe coarse characteristics of musical content despite their ambiguous definitions and boundaries. In fact, many researchers have attempted to perform automatic genre classification and entered their classification accuracies in competitions at the Music Information Retrieval Evaluation Exchange (MIREX). However, bass-line features were not used in previous audio-based genre classification studies. In this section, we show that our bass-line features are effective at improving the accuracy of automatic genre classification.

4.1 Experimental Conditions

We used an audio data set consisting of 300 musical pieces (50 for each genre) of six different genres: Pop/Rock, Metal/Punk, Electronic, Jazz/Blues, Classical, and World. This was taken from the data set distributed on the Web for the ISMIR 2004 Audio Description Contest[1]. To reduce computational costs, we used only a one-minute term for each piece. The one-minute term right after the first one minute was excerpted to avoid lead-ins, which sometimes do not contain bass lines.

Given an audio signal, the feature vector $\boldsymbol{x}$ consisting of the bass-line, timbral, and rhythmic features was extracted using the algorithm described in Section 3. The Mahalanobis distance of the feature vector $\boldsymbol{x}$ to the feature distribution of each genre c was calculated as follows:

$$D_c^2 = (\boldsymbol{x} - \boldsymbol{\mu}_c)^{\mathrm{t}} \Sigma_c^{-1} (\boldsymbol{x} - \boldsymbol{\mu}_c),$$

where t is the transposition operator, and $\boldsymbol{\mu}_c$ and Σ_c are the mean vector and covariance matrix, respectively, of the feature distribution for the genre c. Finally, the genre minimizing the Mahalanobis distance, that is, $\mathrm{argmin}_c D_c^2$, was determined as the classification result.

The experiments were conducted using the leave-one-out cross validation method for each of the with- and without-bass-line conditions. The parameters r, s_1, s_2 were determined as the best parameters found through preliminary experiments for each condition: we used $r = 0.5, s_1 = 7, s_2 = 2$ for the with-bass-line condition and $r = 0.65, s_1 = 8, s_2 = 2$ for the without-bass-line condition.

4.2 Experimental Results

Experimental results are listed in Table 2. The classification accuracy was improved from 54.3% to 62.7% on average. In particular, the result for Electronic was greatly improved, from 10% to 46%. The feature distribution for each genre is shown in Figure 1. Here, the second dimension in the with-bass-line condition contributes to the separation of the distributions for Electronic and other genres. The second dimension represents the pitch variability of bass lines.

The features selected through the dimensionality reduction process for the with- and without-bass-line conditions are listed in Table 3. In the with-bass-line condition, Fv1 (the number of different pitches that appear in at least one frame) and Ft3 (the ratio of successive pitch transition patterns) were selected instead of the temporal mean of the 2nd sub-band intensity and the temporal variance of the 4th sub-band contrast, which were selected in the without-bass-line condition. The temporal mean of the 2nd sub-band intensity was extracted from a low-pitch sub-band, so it can be

[1] http://ismir2004.ismir.net/genre_contest/index.htm

Table 2. Results of automatic genre classification

(a) without bass-line features Avg: **54.3%**

	P/R	M/R	El.	J/B	Cl.	Wo.	Acc.
Pop/Rock	**21**	13	3	7	2	4	42%
Metal/Punk	6	**42**	1	1	0	0	84%
Electric	11	8	**5**	12	9	5	10%
Jazz/Blues	0	0	0	**48**	0	2	98%
Classical	0	0	0	4	**31**	15	62%
World	3	4	0	7	20	**16**	32%

(b) with bass-line features Avg: **62.7%**

	P/R	M/R	El.	J/B	Cl.	Wo.	Acc.
Pop/Rock	**23**	10	8	5	0	4	46%
Metal/Punk	5	**42**	0	2	0	1	84%
Electric	10	1	**23**	6	3	7	46%
Jazz/Blues	5	1	1	**37**	1	5	74%
Classical	0	0	0	2	**45**	3	90%
World	7	1	6	10	8	**18**	36%

Vertical axis: ground truth, horizontal axis: classification results

Table 3. Features selected with dimensionality reduction.

Without bass-line features	With bass-line features
Mean of intensity	Mean of intensity
[Mean of 2nd sub-band intensity]	Mean of 5th sub-band contrast
Mean of 5th sub-band contrast	Fv1: # of pitches that appear
Var of 6th sub-band intensity	Var of 6th sub-band intensity
[Var of 4th sub-band contrast]	Var of 6th sub-band valley
Var of 6th sub-band valley	Mean of spectral flux
	Ft3: % of disjunct+conjunct motions

[ ] denotes features selected for the without-bass-line condition.
Underline denotes selected bass-line features.

regarded as an *implicit* bass-line feature. This result shows that our *explicit* bass-line features contribute to improve separation capacity much more than such an implicit bass-line feature.

One possible reason for genre classification errors is misestimation of bass-line pitches. The misestimation tended to occur at on-beat times when drum instruments sound. One possible solution to this problem is to use drum-sound reduction techniques [12, 13].

4.3 Examination of Ground Truth by Human Subjects

Another possible reason for genre classification errors lies in ambiguous definitions and boundaries of music genres. By conducting a listening test with human subjects, we discuss the appropriateness of the ground truth for pieces whose genres were or were not successfully identified.

We first chose one piece each at random from genre-identified pieces and mis-identified pieces for each genre; the total number of chosen pieces was 12. We then asked human subjects the most likely genre for each piece. The human subjects were 10 people who often listened to music in everyday life. The results are listed in Table 4. For Pop/Rock, Metal/Punk, Electronic, and Classical, at least half the subjects chose the same genres as our system, which were different from the official ground truth. This result implies that these pieces lie on the boundaries of the genres

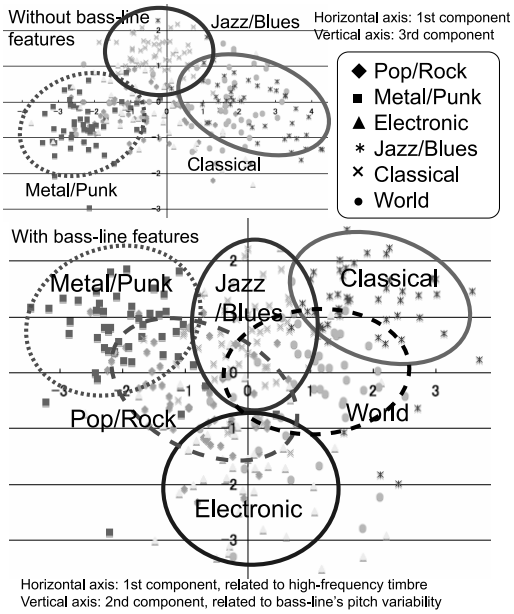


Figure 1. Feature distributions. While the distributions for Electronic and World spreaded out all over the feature space without the bass-line features, those gathered with the bass-line features.

Table 4. Results of listening test by human subjects

	P/R		M/P		El.		J/B		Cl.		Wo.	
	T	F	T	F	T	F	T	F	T	F	T	F
Pop/Rock	[7]	4	3	[**6**]	4	[**5**]		[ ]		2		1
Metal/Punk	3		[7]	3								
Electric		[**6**]		1	[4]	5		2				
Jazz/Blues					1		[10]	8				[2]
Classical									[10]	1		
World					1					[**7**]	[10]	7

T/F denotes the piece whose genre is correctly/incorrectly identified.
[ ] denotes the result of automatic genre classification.
Underline denotes the case that at least half the subjects chose the same genre as the system's output although the ground truth is different.

and that their genre classification is essentially difficult.

5 APPLICATION TO MUSIC ISLANDS

Music genres are useful concepts, but they are essentially ambiguous. Some pieces categorized into Rock/Pop could be very similar to Electronic, and others could be similar to Metal/Punk. We therefore need a mechanism for distinguishing pieces that have such different characteristics even though they are categorized into the same genre. One solution to this is to visualize music collections based on musical similarity. Because different listeners may focus on different musical aspects (e.g. melody, rhythm, timbre, and bass-line), it should be possible to adapt music similarity to such users' differences. We therefore choose to use the Music Islands technique developed by Pampalk [2] as a music collection visualizer and generate different views by switching weights given to the features.

Table 5. Settings of feature weighting

	Bass-line features	Timbral/rhythmic features
All Mix	1.0	1.0
Timbre&Rhythm	0.0	1.0
Bass Only	1.0	0.0

5.1 What are Music Islands

The key idea of Music Islands is to organize music collections on a map such that similar pieces are located close to each other. The structure is visualized using a metaphor of geographic maps. Each island represent a different style (genre) of music. This representation is obtained with an unsupervised clustering algorithm, the Self-organizing Map.

5.2 Switching Views of Music Islands

As Pampalk pointed out [2], there are various aspects of similarity. It is thus better to allow users to adjust the aspects that they are interested in exploring in their music collections. He therefore used three different aspects of features: periodicity histograms related to rhythmic characteristics, spectrum histograms related to timbre, and metadata specified manually by users. Different combinations of these aspects successfully generated different views, but there is room for improvement if features that are more clearly related to specific musical aspects are available.

In this paper, we introduce our bass-line features to generate Music Islands. By switching the weights given to the bass-line and other features, we produce music similarity measures and collection views that enhance different aspects of music.

5.3 Implementation

We implemented a system for visualizing a music collection using Matlab (Figure 2). This system generates three different views called *All Mix*, *Timbre&Rhythm*, and *Bass Only*. The settings of feature weights are listed in Table 5. The users are allowed to select one view based on their interests and to click any cell in a view to select and play back a musical piece. We used SOM Toolbox [14] and SDH Toolbox [15].

5.4 Experiments and Discussions

We conducted experiments on the generation of music collection maps to discuss the difference among three kinds of views. We used the 300 musical pieces (50 pieces per genre $\times$ 6 genres), which were the same as those used in the experiments described in the previous section. The map size was set to 14×14.

The results of generating music collection maps, shown in Figure 3, can be summarized as follows:

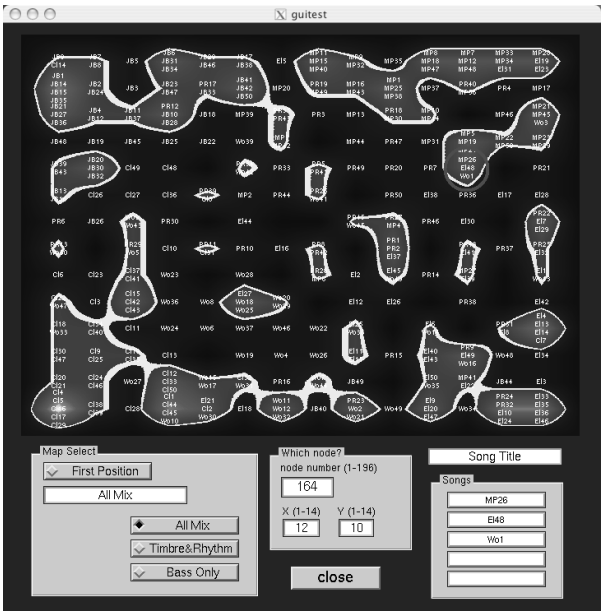
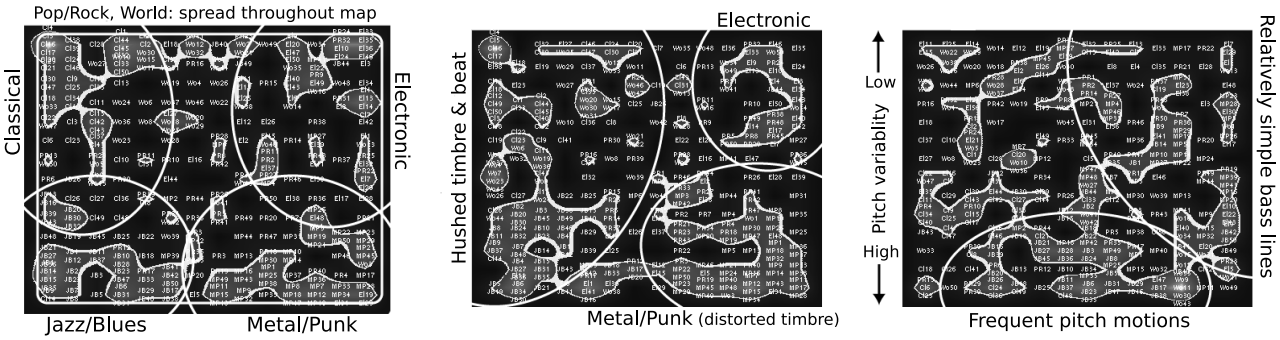


Figure 2. Prototype system of music collection visualizer.

- *All Mix*
  For Classical, Electronic, Jazz/Blues, and Metal/Punk, pieces in the same genre tended to be located close to each other. This is because these genres have comparatively clear characteristics in timbral, rhythmic, and bass-line aspects. Pop/Rock and World pieces, on the other hand, were spread throughout the map. This is because these genres cover a wide range of music, so their correspondence to acoustic characteristics is unclear. In fact, *rock* is sometimes treated as a super category of *metal* and *punk*. *Pop music* is widely used to classify any kinds of musical pieces that have a large potential audience and *world music* to classify any kinds of non-western music.
- *Timbre&Rhythm*
  The basic tendency was similar to that of *All Mix*. The major difference was that islands of Classical and Jazz/Blues were connected by a land passage. This is because music of both genres is played on similar instruments and has a comparatively weak beat. Moreover, Electronic pieces tended to spread much more throughout the map than *All Mix* ones. This result matches the improvement in genre classification.
- *Bass Only*
  Jazz/Blues pieces that had high pitch variability in their bass lines were located at the bottom of the map. On the other hand, pieces that have low pitch variability in their bass lines were located at the top of the map. Thus the map enhanced the characteristics of bass lines.

From these results, we consider that aspects enhanced in



(a) *All Mix* (b) *Timbre&Rhythm* (c) *Bass Only*

Figure 3. Music Islands with three different settings of feature weighting.

the similarity measure can be switched by changing feature weighting.

6 CONCLUSION

In this paper, we described our design of bass-line features to be extracted from audio signals and applied them to automatic genre classification and music collection visualization. Experimental results for automatic genre classification showed that the use of bass-line features improved classification accuracy from 54.3% to 62.7%. Experimental results for music collection visualization showed that we produced music collection views that could enhance different musical aspects by switching the weights to the bass-line and other features.

To achieve user-adaptive MIR, we need feature extraction techniques that can separately capture various musical aspects and feature integration techniques that are aware of users' preferences. If a user tends to focus on a specific musical aspect such as a melody, rhythm pattern, timbre, or bass line, an MIR system for this user should give higher weights to such an aspect than to other aspects. However, only a few attempts [16, 17] have been made to apply features that capture specific aspects to content-based MIR. In particular, no attempts have been made to apply bass-line features. In this paper, we described how we designed bass-line features and applied them to automatic genre classification and music collection visualization. The use of different feature weights achieved different collection views that can be switched by users according to their preferences.

Future work will include integrating our bass-line features with features capturing other musical aspects, for example, instrumentation [16], drum patterns [12], vocal timbre [17], and harmonic content [5].

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TOWARDS STRUCTURAL ALIGNMENT OF FOLK SONGS

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ABSTRACT

We describe an alignment-based similarity framework for folk song variation research. The framework makes use of phrase and meter information encoded in Humdrum scores. Local similarity measures are used to compute match scores, which are combined with gap scores to form increasingly larger alignments and higher-level similarity values. We discuss the effects of some similarity measures on the alignment of four groups of melodies that are variants of each other.

1 INTRODUCTION

In the process of oral transmission folk songs are reshaped in many different variants. Given a collection of tunes, recorded in a particular region or environment, folk song researchers try to reconstruct the genetic relation between folk songs. For this they study historical and musical relations of tunes to other tunes and to already established folk song prototypes.

It has often been claimed that their work could benefit from support by music information retrieval (MIR) similarity and alignment methods and systems. In practice however it turns out that existing systems do not work well enough out of the box [5]. Therefore the research context must be analyzed and existing methods must be adapted and non-trivially combined to deliver satisfying results.

1.1 Similarity and alignment

Similarity and alignment can be considered two sides of the same coin. In order to produce an automatic alignment we need a measure for the relatedness of musical units. Conversely, in order to compute the (local) similarity between two melodies we must know which parts of the melody should be compared.

Alignment can also be a prerequisite for higher-level similarity measures. In a previous paper we derived generalized queries from a group of folk song variants [3]. For a given group of musically related query melodies aligned by the user, we were able to retrieve melodies from a database that are good candidate members for this group.

Making a manual alignment is time-consuming and involves edit decisions, e.g. 'shall one *insert* a rest in one melody or *delete* a note in the other?'. When looking for good additional group members in a database, one should allow both options. However, keeping track of all options quickly becomes impracticable. In this paper we therefore look into automatic alignment of corresponding score positions and ways of controlling the alignment with basic similarity measures.

1.2 Overview and related work

In this paper we first discuss why automatic detection of (genetically related) folk song variants is very demanding and is a major research topic in its own. Next, to support research into similarity measures based on musically meaningful transformations, we develop a framework that helps to model the influence of local similarity measures on variation detection and alignment. Starting from the information encoded in our folk song collection, we motivate the use of available structural and metrical information within alignment directed similarity measures. Finally we compare automatically derived alignments with alignments annotated by an expert.

Generally, we follow a similar approach to Mongeau and Sankoff's [6], who tackled selected transformational aspects in a generic way. They set up a framework to handle pitch contour and rhythm in relation to an alignment-based dissimilarity (or quality) measure. They based their framework assumptions on musical

common sense and based their model parameter estimations on the discussion of examples.

We agree with them that musical time (as representated in common music notation) is a fundamental dimension in variant similarity. This distinguishes our approach from ones that deal with performance-oriented timing deviation problems. A shortcoming of Mongeau and Sankoff's global alignment algorithm is that it cannot handle aspects of musical form, such as repeats or reordering of parts. They are also criticized for sometimes ignoring barlines [2].

Contribution: In this paper we present an approach to tackle these shortcomings by proposing a phrase- and meter-based alignment framework. By using simple and out-of-the-box local similarity measures within the framework and studying the resulting alignments we prove that the framework is useful.

2 MUSICOLOGICAL MOTIVATION

For folk song researchers, an important question is to identify which songs or melodies are genetically related. A way for them to tackle this question is to order melodies into groups that share relevant musical features. We assume that in the process of incrementally building those melody groups, researchers construct a sort of mental alignment of related parts of the candidate melodies. The detection of relevant relationships between variants is affected by the perceived similarity of the material, the knowledge of common transformation phenomena and the discriminative power of the shared features with respect to the investigated corpus.

Researchers have identified transformations that range from a single note change to changes of global features like mood. A mood change can for example affect the tonality (major/minor), the number of notes (due to liveliness) and ambitus (due to excitation) [10].

2.1 Modeling musical similarity

To support folk song variation research, one could choose to model expert reasoning using rule systems. Such a system would consist of transformation rules and transformation sequences that taken together model the relation between melodic variants. A fundamental problem with this approach is that we are still a long way from sufficiently understanding music perception and cognition. Therefore it is impossible to fully formalize the necessary expert knowledge and model the rules of musicological discourse.

Also, it is difficult to find out which approaches to folk song variation are the most promising, because there is little scholarly documentation about music transformations. An exception is Wiora's catalog of transformation phenomena [10]. It describes musical features and related transformations and is a good inspirational source for models of similarity and human reasoning about it. But it lacks descriptions of the contexts in which certain transformations are permissible. It also does not provide the means to estimate the plausibility of one transformation chain as compared to another when explaining a certain variation. It is common in folk song research to reason about the relatedness of *specific songs* rather than to provide comprehensive models. However, what is needed for MIR is a comprehensive theory about this kind of reasoning.

We chose a different approach to model variant relationships, namely to investigate local similarity. This closely follows musicological practice. Often some *striking similarities* between some parts are considered sufficient evidence for a variant relationship. This seems to leave MIR with the task to design local similarity measures for variation research. However we also need to model what is *striking* and what are *relevant parts* and we must find ways to turn local similarity values into overall relevance estimates.

2.2 Structural, textual and metrical alignment

In this section we mention some relations between the parts of a song and between parts of variant songs that support or inhibit the detection of variants. By doing so, we motivate the structure-based alignment framework that we describe in the next sections. Our assumptions stem from general experience with folk songs and from dealing with a collection of Dutch folk songs that we currently help making searchable. [9]

Strophes: In songs, music is tightly bound to the lyrics. When two different songs have similar lyrics, we can often simply use textual similarity to relate musical parts with high confidence. However, we cannot rely on text alone: if textually different variants or different strophes are encoded, we need to make use of musical similarities. This should be unproblematic in principle, since different strophes of a song typically

share the same basic melody.

Phrases: Musical phrases in a folk song are typically related to the verse lines of a poem. When dealing with automatic alignment we must be aware that the level of phrase indication might differ: one phrase in song A might match with two shorter phrases in song B, or the same rhythm can be notated with different note values.

Accents: Word accents in a verse typically follow a common scheme, such as dactyl and trochee. These accents often correspond to the metrical accents of the notes, which can be found by looking at the barlines and measure signatures. We cannot always assume the same meter across variants, since different durations can turn a 2-foot verse scheme into a 3-accent melody. Also, extra syllables in the lyrics may make it necessary to insert notes.

Accent (beat position) and phrase position may also be important for musical similarity of variants: in our previous work we found that pitches on stronger beat positions tend to be more stable across variants than less accented notes. There also seems to be a higher agreement between variants at the beginning and end of a strophe or a phrase (cadence notes), while inner variation is higher [4]. We will study these claims further in future research.

3 A STRUCTURE-BASED ALIGNMENT FRAMEWORK

In this section we describe three components for structure-based folk song alignment: hierarchical segmentation, phrase-level alignment and strophe alignment. The general idea of the proposed framework is to use alignments and alignment scores of smaller musical parts in the alignment process on higher levels. The more part A from the first melody resembles part B from the second melody according to a similarity measure, the more preferable is it to align A with B rather than with another part of the second melody with a lower similarity. However, constraints in higher level alignments can overrule this preference.

3.1 Hierarchical song structure

In accordance with the analysis given in the previous section, we split songs hierarchically into smaller parts. We work on manually encoded musical scores in which meter information and phrases that correspond to the lyrics are indicated. For each song, at least one strophe is encoded. For some songs two or more strophes are separately encoded. We convert our encodings to Humdrum **kern format with the usual = bar line markers and special `!!new phrase` comments.

We use the *Humextra* [7] toolkit to access the relevant information in *Humdrum* `**kern` files (one file per *strophe*). In our data model, each strophe contains a sequence of *phrases*. We currently do not deal with repeat structures, since they are not encoded in our data, but written out. Each phrase is split into *bar segments* using the given bar lines.

A bar in turn is recursively split into smaller *beat segments* delimited by metrically strong beat positions. These are not encoded and need not coincide with note events, but are inferred from the bar lines and the measure signature.

To each structural unit we attach both the corresponding Humdrum fragment, which can be used to retrieve notes and extra information such as the syllables, and a beat range, which identifies the start and duration of the segment measured in beats.

When retrieving the notes for a structural unit, special care is taken to handle boundaries: incomplete bars can occur not only at the beginning or end of a strophe but also at the beginning or end of a phrase. A bar segment will only return those notes of the bar that are part of the parent phrase segment, and likewise for a beat segment.

3.2 Phrase level alignment

The user of the framework chooses an elementary similarity measure sim that is defined on either bar segments or beat segments. The framework computes a similarity value for any pair of segments (one segment from melody A, the second from melody B).

To combine segments of two phrases into a **phrase level** alignment, we use standard global string alignment techniques with match scores and deletion and insertion related gap scores [11]. We define the match score of two segments a and b as being equal to $sim(a, b)$. The scaling of gap scores with respect to sim is left to the user.

To cope with the common phenomenon of different durations of the upbeat bar and the last bar in the same phrase in two variants, we support different gap scores for *inner* and (left/right) *outer* gaps. Also, we could

use local instead of global alignment methods to look for motivic similarity instead of phrase similarity [6].

Future improvements will support *augmentation*, *diminution*, *fragmentation* and *consolidation* as described in [6] in combination with segment constraints. We will also look into inner-phrase form deviations, such repeats of a bar or beat segment. (See section 5.)

3.3 Strophe alignment

For the **strophe level** alignment the framework employs phrase alignments: from the alignment scores similarity values are calculated for all possible pairs of phrases (one phrase from melody 1, one from melody 2). Different phrase-level similarity values from alternative elementary similarity measures and from non-alignment similarity measures (e.g. on cadence tones) can be consolidated into one similarity value. Then a string alignment technique can be used again to find the best alignments of the phrase sequences based on these similarity values. This handles transformations from ABCD to ABD.

To assume sequential similarity and to use phrases only once on the strophe level would sometimes be misleading. Consider simple transformations from AABB to AAB or BBAA. Therefore the framework supports the creation of alignments where one strophe is fixed and each phrase p of it can be matched against any of the phrases q of the phrase set S variant strophe: $MatchScore(p, S) = max_{q \epsilon S}\{similarity(p, q)\}$

To cover cases where strophes of one song differ significantly, the framework simply performs all pairwise comparisons between all strophes from one variant with all strophes from the other variant.

4 EVALUATION

To study the usefulness of our framework, we compared alignments produced by framework-based models with manual alignments (annotations). One of the authors selected sets of similar phrases from a variant group and produced for each set a multiple alignment of their segments in a matrix format (one line per phrase, one column per set of corresponding segments). Segments are identified by their starting metrical position (e.g. 3.2 for bar 3, second part) and all segments in a bar must be of the same size.

From each multiple alignment annotation of N phrases we derived N(N-1)/2 pairwise alignments. We compared these to automatic alignments derived from specific framework setups. Each setup consists of:

A basic distance measure (*seed*) acting on segments defined by the expert annotation. The segments usually stem from the first subdivision of the bar (one half of a bar in 6/8, one third of a bar in 9/8 measure). Exception: a 4/4 variant in a 6/8 melody group is split into four segments per bar.

A normalization to turn the segment distance values into match scores between 0 and 1. We employ $e^{-distance}$ as the match score for this experiment.

Gap penalty scores for inner and outer gaps (between 0 and 1.5 for this experiment). Note: gap scores are subtracted and match scores are added in an alignment.

For each setup we generated a log file. For overall performance comparisons we produced summary fitness values per variant group and across all tested variant groups. For the fitness of a setup for a particular variant group (*group fitness*), we counted all pairwise alignments in which the automatic alignment has the same gap positions as the annotation and divided this number by N(N-1)/2. For the *overall fitness* of a setup, we took the average of the group fitnesses.

4.1 Results

Four (not necessarily representative) groups of variant phrases were manually aligned and used for evaluation. We only present a summary about the lessons learned from studying the log files [1], which contain annotations, links to the musical scores, alignment results, failure analysis information and summaries.

The overall performance of selected setups is shown in table 4.1 and discussed in the next sections.

4.2 Discussion of distance seeds

In this section we discuss the performance of increasingly complex elementary distance measures (seeds). Baselines are provided by **trivial** and **random**.

The trivial distance 0 turns into a match score of 1 to any pair of segments. As a consequence the actual alignment depends on the gap scores and the algorithm execution order only. In our test setup this always means that the algorithm always chooses left

Seed	IG	OG	G1	G2	G3	G4	A1	A2
trivial	0.0	0.0	33	50	20	10	28	34
random	0.5	0.3	50	16	30	20	29	32
random	1.5	1.0	83	50	20	10	40	51
beatPos	0.0	0.0	33	16	20	10	19	23
beatPos	0.5	0.3	33	50	20	10	28	34
beatPos	1.5	1.0	100	50	20	10	45	56
events	0.5	0.3	100	50	40	30	55	63
events	1.5	0.5	100	50	20	30	50	56
ptdAbs	0.5	0.3	66	83	50	30	57	66
ptdAbs	1.5	1.0	100	83	40	20	60	74
ptdRel	0.5	0.3	66	50	50	20	46	55
ptdRel	1.5	1.0	100	50	50	20	55	66

Table 1. Alignment fitness. IG/OG: inner/outer gap scores. G1-4: percentage of well-aligned phrases per group. A1: average of G1-4; A2: average of G1-3.

(outer) gaps to compensate for the beat segment count difference of the variants.

A random distance between 0 and 1 leads to a more even distribution of gaps. When outer gaps are cheaper, there is a preference for outer gaps. Interestingly in our examples *random* performs better than *trivial*, because the manual alignments contain more right than left outer gaps. As a consequence we should consider to lower the right outer gap penalty with respect to the left in future experiments.

To study the performance using phrase and meter information only, we defined the **beatPos** distance as the difference of the segment number relative to the previous barline. The second segment of a bar thus has distance 1 to the first segment. The algorithm should prefer to align barlines this way. Surprisingly it performed worse than *trivial*. However, we found that too many (relatively cheap) gaps were introduced to match as many segments as possible. We compensated this in another test run with gap penalties greater than 1 and achieved much higher fitness than *trivial*. In general there were only few examples where both phrases were supposed to have inner gaps at different positions.

The next distance measure **events** measures the difference of the number of notes in a segment. Tied notes that begin before the segment starts count for both the current and the preceding segment. The effect of this measure is that regions of same event density (related to tempo or rhythm) are preferred matches. *Events* performs overall better in the alignment than *beatPos*.

To take both onset and pitch into account at the same time, we used proportional transportation distance (PTD) [8]: we scaled the beat segment time into [0..1] and weighted the note events according to their scaled duration. As the ground distance we used $(4\Delta pitch^2 + \Delta onset^2)^{-2}$ with pitches measured as MIDI note numbers modulo 12. Our distance measure **ptdAbsolute** takes advantage of the fact that the given melodies are all in the same key. It performs best in comparison with the previous measures.

If we do not assume the same key, it does not make sense to employ absolute pitch, but instead one can compare only the contours. One approach to tackle this is **ptdRelative**, which takes the minimum over 12 *ptdAbsolute* distances. However, it performs much worse. The reason for this is that, given this distance measure, two segments that each contain a single note always have a distance 0. One should therefore apply this measure only on larger segments or model the tonal center in the state of the alignment process (see section 5).

4.3 Discussion of annotation groups

The four variant groups were chosen to display different kind of alignment problems.

The manual alignment of the group G1 (*Wat hoor ik hier...*) does not contain any inner gaps. There is little variation in absolute pitch, so *ptdAbsolute* achieves 100% the same alignments. The framework proves to be useful and handles different kind of measures (6/8 and 9/8) correctly.

Group G2 (*Ik ben er ...*) contains one inner gap. According to the annotation, the final notes *d,c,b* of one variant (*72564*) are diminished (*d,c* lasts one beat instead of two beats). Because the framework does not handle such transformations yet, this was annotated as a gap. For the similarity measures this gap is hard to find, probably because the neighboring segments provide good alternative matches and often a right outer gap is preferred. Lowering the gap penalties leads to the introduction of unnecessary extra gaps. However, *ptdAbsolute* with high gap penalties achieves 83% success and misses only one pair (*72564* and *72629*), because it matches *d* with *e*. The framework deals well with aligning 4/4 with 6/8 measures.

Group G3 (*Moeder ik kom ...*) contains a repeated segment. Variants that have no repeat are annotated with a gap at the first occurrence of the segment. However, there is no compelling reason why this gap cannot

be assigned to the second occurrence. This ambiguity accounts for many "failures" in the test logs.

Group G4 (*Heer Halewijn*) was chosen because of its complexity. Only when looking at the annotation for a while the chosen alignment becomes understandable. It is mainly based on tonal function of pitches and contains many inner gaps. For pairs of phrases, other alignments are plausible as well, but in the multiple alignment several small hints together make the given annotation convincing. Therefore there are only few correct automatic alignments. Interestingly, however, the algorithm manages to align a subgroup (*72256*, *74003*, and *74216*) without failure.

5 CONCLUSION

We have presented a structure-based alignment and similarity framework for folk song melodies represented as scores. We have done initial tests that show both the usefulness and limitations of our segmentation, alignment and evaluation approach. We see two continuations. First, we should use the framework to study similarity seeds that take the observed stability of beginnings and endings into account (see section 2.2).

Second, the alignment framework needs to be developed further into several directions. 1) We did not so far pay any attention to the relationship between the statistical properties of the distance measure, its normalization and the value of the gap penalties. 2) We should support the modeling of *states* and non-linear gap costs. 3) *Multiple alignment* strategies should be incorporated in order to relate more than two melodies. The need for this became apparent in the last alignment group. Multiple alignments are particularly needed for group queries [3]. Therefore we will not only evaluate the quality of the alignments but also the performance of melody retrieval using these alignments.

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A NEW MUSIC DATABASE DESCRIBING DEVIATION INFORMATION OF PERFORMANCE EXPRESSIONS

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ABSTRACT

We introduce the CrestMuse Performance Expression Database (CrestMusePEDB), a music database that describes music performance expression and is available for academic research. While music databases are being provided as MIR technologies continue to progress, few databases deal with performance expression. We constructed a music expression database, CrestMusePEDB. It may be utilized in the research fields of music informatics, music perception and cognition, and musicology. It will contain music expression information on virtuosis' expressive performances, including those of 3 to 10 players at a time, on about 100 pieces of classical Western music. The latest version of the database, CrestMusePEDB Ver. 2.0, is available. The paper gives an overview of CrestMusePEDB.

1 INTRODUCTION

Constructing music databases is one of the most important themes in the field of music studies. The importance of music databases has been recognized through the progress of music information retrieval technologies and benchmarks. Since the year 2000, some large-scale music databases have been created, and they have had a strong impact on the global research area [1, 2, 3].

Corpora of scores, audio recordings, and information on books of composers and musicians have been collected and used in the analysis of music styles, structures, and performance expressions. Meta-text information, such as the names of composers and musicians, has been attached to large-scale digital databases and been used in the analysis of music styles, structures, and performance expressions from the viewpoint of social filtering. In spite of there being many active music database projects, few projects have dealt with music performance expression, such as dynamics, tempo, and the progression of pitch. The performance expression plays an important role in formulating impressions of music. Providing a performance expression database, especially describing deviation information from neutral expression, can be used as a research in MIR fields.

In musicological analysis, some researchers construct a database of the transition data of pitch and loudness and then use the database through statistical processing. Widmer *et al*. analyzed deviation of tempi and dynamics of each beat from Horowitz's piano performances [4]. Sapp *et al*., working on the Mazurka Project [5], collected as many recordings of Chopin mazurka performances as possible in order to analyze deviations of tempo and dynamics by each beat in a similar manner to Widmer. Their researches are expected to provide the fundamental knowledge in the quantitative analysis of music style and performance expression. However, their archives are focused on the specified composer and player but ignored all other types of music. And they are not enough to be covered with the description of the deviation of duration and loudness of each note from score. Toyoda *et al*. have provided a performance expression deviation database which is aimed to describe those of individual note [6]. But their data format is limited to performances recorded by MIDI signals not audio signals. However, most of the analysis of classical Western music expression is determined by the researchers' acute musical sense; even less research has quantitatively analyzed music performance expression based on data [7, 8, 9, 10, 11]. One of the reasons for this absence of data is that this qualitative research has been based on the field methods of music performance education up until now.

MIDI files of performances are prolific on the Internet; however, very few of these are actually created by professional musicians, but rather by students and amateurs. Also, searching with the precise details on the performances is too difficult, although the information on each piece (e.g., the title, composer, genre, and the composition of the musical instruments) is rich.

A database, which describes outlines of changes in tempo, dynamics, the delicate control of each note, the deviation regarding starting time and duration to existing virtuoso performances in the form of acoustic signals as time-series data, can be used for new studies in the fields of music informatics, musicology and music education. To that end, we constructed a performance expression database, 'CrestMusePEDB,' that covers classical Western music, especially piano pieces performed by famous professionals.

2 CONSIDERATIONS IN THE DATABASE DESIGN

2.1 Ensuring the amount and quality of performances

A music database generally needs to contain high-quality performances. Goto *et al*. recorded performances by many professional musicians as part of their RWC music database [12]. Copyright restrictions pose difficulties for performance database creators. One method of avoiding such restrictions is to have the collectors record musical performances specifically for the database. Financially this is a onerous task and can compromise quality when extending a large database. For this problem, McEnnis *et al*. suggested an approach: collecting the parameters of audio features (covered in music information retrieval) in the real performances around the world and showing those reference information[3]. This approach is a rational way to avoid copyright problems and that the system can deal with raw data of the real world. We used the same approach.

2.2 Handling performance control data

The main information in a performance expression database pertains to instrument control. It has various kinds and levels of information. For keyboard instruments such as the piano, control data can be represented as the onset time, the offset time, the dynamics (velocity), and the depth of the damper pedal. The information on the instrument control deals with the feature quantity at the MIDI level. Describing them in a database requires having really useful information that can be extended in the future. Therefore, we constructed control information as XML-based data in CrestMusePEDB. We will extend and work out the details of these descriptions in accordance with the individual situations and technology.

Another problem which deals with music information is to identify the intensity of each note from acoustic signal. We especially need to be careful in describing the loudness of each note. To obtain the strict value of loudness of a note, we need to construct a sound model including the characteristics when recording the performance and architectural acoustic characteristics. However, getting that information from only the audio signal is quite difficult. Although some approaches using non-negative matrix factorization (NMF) are suggested, the problem has not been radically resolved yet in audio signal processing. In the view point of construction of a useful database, those information of the performance data should be collected as correctly by auditory as possible. Therefore, the velocity data is estimated approximately in CrestMusePEDB based on a specific digital audio source by well-trained experts using some support tools described in section 3.4.

2.3 Musical structure

A specific musical structure corresponds with a performance. CrestMusePEDB includes the data of the musical structures in an analysis of each performance, and it provides support to use the data in research on musicology, performance theories, and computational music systems.

The best solution is to interview a pianist and ask how his/her performance was achieved based on the musical structure. However, such an interview must be done immediately before and after the recording. The musical structure data should be collected by estimating the data from audio signals. Our approach involves interviewing as many players as possible to estimate the structures through discussion between a few musicologists and asking an expert pianist to play some performances with different musical structures. We aimed to get both the musical structure and the way it is expressed from the original recording

3 OVERVIEW OF CRESTMUSE PEDB

3.1 Music pieces

The focus of the database is on classical music from the Baroque period through the early twentieth century, including music by Bach, Mozart, Beethoven and Chopin. We chose around a hundred music pieces, including those referred to often by previous music studies in the past couple of decades. We also chose various existing performances by professional musicians; the database includes 3 to 10 performances for each musical score.

3.2 Contents of the database

The database consists of the following four kinds of component data. These kinds of data will sequentially be provided from SCR, DEV. The description of the database is based on XML files. CrestMusePEDB does not contain any acoustic signal data (WAV files, AIFF files, MP3 files.) except for PEDB-REC. Instead, it contains the catalogs of the performances from which expression data is extracted. The original audio sources can be purchased by the database users if necessary.

PEDB-SCR (score text information): The score data included in the database: files in MusicXML format and in SMF (standard MIDI file) format will be provided.

PEDB-IDX (audio performance credit): The catalogs of the performances from which expression data are extracted: album title, performer's name, song title, CD number, year of publication.

PEDB-DEV (performance deviation data): Tempo changes outline changes in dynamics, the delicate control of each note, deviation regarding starting time, duration, and dynamics extracted from expressive performances. Performances from 3 to 10 performers are analyzed a piece, and plural

deviation data analyzed by different sound sources are provided a performance. Each data is described in DeviationInstanceXML format (see below).

PEDB-STR (musical structure data): This contains the information on a musical structure data (hierarchical phrase structure and the top note of a phrase). The data is described in compliant MusicXML format. The structure data corresponds with a performance expression data in PEDB-DEV. But if plural adequate interpretations exist in a piece, the plural structure data is provided in the performance data.

PEDB-REC (original recordings): The audio performance data we will newly record based on PEDB-STR. The data will be useful to analyze performance expression from the view of music structure. It provides an audio signal and MIDI data that an expert pianist plays.

3.3 DeviationInstanceXML format

DeviationInstanceXML describes the deviation information from score information in a music performance as a subset of CrestMuseXML (CMX)[1] , which is the universal data format for music informatics. The latest version (0.32) deals with the control information of the piano playing, such as the onset time, offset time, and the velocity in MIDI of each note, tempo control, and damper pedal control. This control data is categorized into (1) common or multiple parts (non-partwise), (2) each part (partwise), and (3) each note (notewise) and is described as follows:

```
<deviation target="sample.xml" xmlns:
      xlink="http://www.w3.org/1999/xlink">
  <non-partwise>...</non-partwise> (1)
  <partwise>...</partwise> (2)
  <notewise>...</notewise> (3)
</deviation>
```

The details of this description are shown in the site, but here is the overview shown.

(1) <non-partwise> One of the common deviations of plural parts is tempo information. It is described as a combination of two XML tags: one is <tempo>, which describes the fundamental impression such as Andante and Allegro. It indicates the number of beats per minute (BPM). The other is <tempo-deviation>, which describes the micro tempo per beat; it is expressed as the ratio of the value of <tempo>.

(2) <partwise> The deviations of each part deal with the fundamental dynamics and damper pedal control into a <partwise> tag. The fundamental dynamics describe the global indication such as forte and piano. The pedal control is available to be described in the MusicXML format, but the control refers to the expression of the score and does not always correspond to the performance. The pedal description in DeviationInstanceXML handles the pedal control in an actual performance expression. Some people might think that the pedal control should be described as the common deviation of all the parts. For CrestMusePEDB, the pedal control is sufficient because the only instrument in all of the performances is the piano. However, the DeviationInstanceXML will be used to describe the control with plural instruments in the future. This issue is controversial, but here, the pedal control is described in <partwise>.

(3) <notewise> The deviation information of each note deals with the deviation of the onset time and offset time from the score and the deviation of individual dynamics (the velocity of each note key). The individual dynamics are for describing a micro-expression within a phrase to which the note belongs. The actual loudness of the note is computed by multiplying the individual dynamics by the fundamental dynamics at the score-time of the piece.

[1] http://www.crestmuse.jp/cmx/

3.4 Procedure for making DEV data

Transcribing an audio-signal performance into MIDI-level data is the core issue for constructing the CrestMusePEDB. Although we have improved the automation of the procedure, at present, an iterated listening process by plural experts with commercial audio editor and original alignment software possesses higher reliability. The data needs to be extracted from a recording as efficiently as possible. We now use an approach where four experts who have specialized in music including piano, composition and musicology manually identify the transcription/data in CrestMusePEDB using their ears and three support tools that we have implemented for the procedure.

Figure 1 illustrates the procedure for generating PEDB-DEV information.

Step 1: identify the attack time in the beat level of a piece while listening to the CD audio.

Step 2: estimate the approximation of the onset time, offset time, velocity of each note, and whether or not the damper pedal is used and whether or not it is fixed to the sound source.

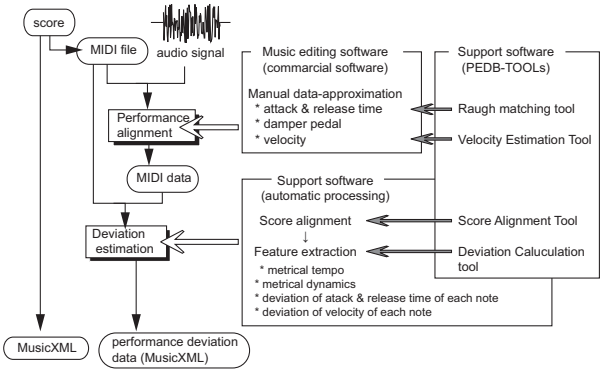

Figure 1. The procedure to make PEDB-DEV.

Step 3: refine the estimation with repeating step2 until a expert accepts and have it cross-checked by other experts. This step needs the most time of all the steps and takes at least five hours to a day a piece.

Step 4: match the expression data obtained through steps 1-3 with the normal expression data.

Step 5: calculate the time transition of tempi and dynamics, the deviation of the onset and offset time of each note, and the deviation of velocity from the fundamental dynamics (nominal expression) to the whole score.

The velocity estimation in step 2 and step 3 is necessary for resolving the problem with the loudness range when an audio signal differs because each audio CD was recorded using different musical instruments and in different environments. Here, we give priority to obtaining the value of the velocity for each note approximated to the listening data of the audio signal by experts who listen to the data of the audio signal. After justifying the range of the loudness for all of the audio signals, the experts repeat refining the approximated data with their headphones, which is always identical.

We have implemented three of support software to help with each step of the procedure.

Rough Matching Tool enables aligning the rough attack time of each note of a score corresponding with audio spectrogram, using Dynamic Programming.

Velocity Estimation Tool enables estimating the velocity of each note based on the original audio signal, the onset time of each note (which is roughly identified by the above tool), and an audio source for the sound of the MIDI data.

Score Alignment and Deviation Calculation Tool supports two procedures to extract the deviation information. The first one identifies all the notes of the score as those of the expressed performance. The second one calculates the deviation information (the onset time, the offset time, and the velocity) for each note. The deviation information is described in the DeviationInstanceXML format.

The rough matching tool and the velocity estimation tool do not have to function at a very high accuracy in order to be beneficial. The skill of experts is arguably higher than that of current systems. The former tool is used for step 1, and the latter is used when deciding the initial parameters in step 2. Meanwhile, the score alignment and deviation calculation tool seldom make errors. We determined the result data using these tools in step 4 and step 5. We will make these tools public so that a community of CrestMusePEDB users will grow.

3.5 Subset tools

The data description in the XML format has scalability and data integrity. Meanwhile, it is not so useful for researchers except for those in computer science. Therefore, we provide the following sub-tools for CrestMusePEDB so that it can be effectively utilized.

CrestMusePEDB player: a music player that inputs SCR and DEV data, then generates expressed performance data through the indicated audio source.

CrestMusePEDB time-line viewer: a deviation visualizer that inputs SCR and DEV data then shows each note and the data described in the DEV data to a piano-roll display.

CSV (Comma Separated Values) Converter: a plugin tool for the CrestMusePEDB time-line viewer that outputs selected deviation information in the CSV format.

4 PUBLISHING OF CRESTMUSE PEDB

CrestMusePEDB has been available since November 2007. The version 1.0 and 2.0 have been released on the Internet[2] and there are totally 60 performances included. Table 1 and 2 shows the list of the music pieces included in the database, and Table 3 shows a player list and a sound source corresponding with each piece.

Each DEV data is given a unique number (PerfID) and an identification ID to discriminate itself along with a form. Identification ID is given by the combination of ScoreID and DevID. For example, the database has three perfomance DEV data (Gould, Nakamura and Shimizu) of the 1st movement of Mozart's Piano Sonata K. 331. The IDs of the Gould's performance are described as "PEDB-DEV No. 22 (snt331-1-003-b)". The alphabet at the last of DevID means a sound source used for extracting deviation data from audio source; 'a' is B osendolfer PIANO/GIGA by Crypton Future Media Inc., 'b' is GPO-Concert-Steinway-ver2.sf2 (a free source from SF2midi.com) and 'c' is Yamaha's MOTIF XS. The performances using 'c' will be provided at the future version.

We have been preparing a listening contest for music performance rendering systems (Rencon) as part of another research project (`http://www.renconmusic.org/`). In the summer of 2008, the database is provided to the Rencon competition as the standard learning set for autonomous performance rendering systems that generate expressive performance. We have been already providing some sample data on CrestMusePEDB to researchers who plan to enter the Rencon contest.

5 DISCUSSION

5.1 Applied research area

CrestMusePEDB can be used in various research areas such as musicology and music education, not only MIR fields. Next, we describe some cases for application.

Musicology In musicology, many analyses of expressive performance style examine the presence of *legato*/*non-legato* phrasing [13]. Many of them depend on a subjective verdict by an analyst. As well as these verdicts, we suppose that the

[2] `http://www.crestmuse.jp/pedb/`

Table 1. CrestMusePEDB ver. 1.0 Performance Data (No. 1-39)

Composer (ComposerID) (amount of performances)						
	PerfID	ScoreID	Title			
			Player	DevID	CD index	Track.
J. S. Bach (1) (21 performances)						
	No. 1〜2	inv001	Invention No. 1 BWV 772			
			A. Schiff	001-a/b	POCL-5099	Tr.01
	No. 3	inv002	Invention No. 2 BWV 773			
			A. Schiff	001-b	POCL-5099	Tr.02
	No. 4	inv008	Invention No. 8 BWV 779			
			A. Schiff	001-b	POCL-5099	Tr.08
	No. 5	inv015	Invention No.15 BWV 786			
			A. Schiff	001-b	POCL-5099	Tr.15
	No. 6-11	wtc101-p	Das Wohltemperierte Klavier Vol. No.1 BWV 846 Prelude			
			S. Richter	002-a/b	BVCC-37139	Tr.01
			G. Gould	003-a/b	SRCR9496	Tr.02
			F. Gulda	004-a/b	416-113-2	Tr.01
	No. 12	wtc107-p	Das Wohltemperierte Klavier Vol. I No. 7 BWV 852 Prelude			
			S. Richter	002-b	BVCC-37139	Tr.13
	No. 13	wtc107-f	Das Wohltemperierte Klavier Vol. I No. 7 BWV 852 Fuga			
			S. Richter	002-b	BVCC-37139	Tr.14
	No. 14	wtc113-p	Das Wohltemperierte Klavier Vol. I No.13 BWV 858 Prelude			
			S. Richter	002-b	BVCC-37139	Tr.01
	No. 15	wtc113-f	Das Wohltemperierte Klavier Vol. I No.13 BWV 858 Fuga			
			S. Richter	002-b	BVCC-37140	Tr.02
	No. 16	wtc123-p	Das Wohltemperierte Klavier Vol. I No.23 BWV 868 Prelude			
			S. Richter	002-b	BVCC-37140	Tr.21
	No. 17	wtc123-f	Das Wohltemperierte Klavier Vol. I No.23 BWV 868 Fuga			
			S. Richter	002-b	BVCC-37140	Tr.22
	No. 18	wtc202-p	Das Wohltemperierte Klavier Vol. II No. 2 BWV 871 Prelude			
			S. Richter	002-b	BVCC-37141	Tr.03
	No. 19	wtc202-f	Das Wohltemperierte Klavier Vol. II No. 2 BWV 871 Fuga			
			S. Richter	002-b	BVCC-37141	Tr.04
	No. 20	wtc219-p	Das Wohltemperierte Klavier Vol. II No.19 BWV 888 Prelude			
			S. Richter	002-b	BVCC-37142	Tr.11
	No. 21	wtc219-f	Das Wohltemperierte Klavier Vol. II No.19 BWV 888 Fuga			
			S. Richter	002-b	BVCC-37142	Tr.12
W. A. Mozart (02) (8 performances)						
	No. 22-24	snt011-1	Piano Sonata No.11, K. 331, 1st Mov.			
			G. Gould	003-b	SRCR-9669	Tr.01
			H. Nakamura	005-b	AVCL-25130	Tr.07
			N. Shimizu	006-b	RWC-MDB-C-2001-M05	Tr.01
	No. 25	snt011-2	Piano Sonata No.11, K. 331, 2nd Mov.			
			H. Nakamura	005-b	AVCL-25130	Tr.08
	No. 26	snt011-3	Piano Sonata No.11, K. 331, 3rd Mov.			
			H. Nakamura	005-b	AVCL-25130	Tr.09
	No. 27	snt016-1	Piano Sonata No.16, K. 545, 1st Mov.			
			G. Gould	003-b	UCCG-5029	Tr.10
	No. 28	snt016-2	Piano Sonata No.16, K. 545, 2nd Mov.			
			G. Gould	003-b	UCCG-5029	Tr.11
	No. 29	snt016-3	Piano Sonata No.16, K. 545, 3rd Mov.			
			G. Gould	003-b	UCCG-5029	Tr.12
L. V. Beethoven (03) (5 performances)						
	No. 30	snt008-1	Piano Sonata No. 8, Op. 13, 1st Mov.			
			V. Ashkenazy	007-a	POCL-5005	Tr.07
	No. 31	snt014-2	Piano Sonata No.14, Op. 27-2, 2nd Mov.			
			B. Brendel	008-b	PHCP-21023	Tr.08
F. Chopin (04) (7 performances)						
	No. 32	pld001	Prelude Op. 28, No. 1			
			V. Ashkenazy	007-b	POCL-5064	Tr.01
	No. 33〜34	pld004	Prelude Op. 28, No. 4			
			V. Ashkenazy	007-b	POCL-5064	Tr.04
			M. Argerich	009-b	UCCG-5024	Tr.04
	No. 35	pld007	Prelude Op. 28, No. 7			
			V. Ashkenazy	007-b	POCL-5064	Tr.07
	No. 36	pld015	Prelude Op. 28, No. 15			
			V. Ashkenazy	007-b	POCL-5064	Tr.15
	No. 37	pld020	Prelude Op. 28, No. 20			
			V. Ashkenazy	007-b	POCL-5064	Tr.20
	No. 38	wlz007	Waltz Op. 64-2, No. 7			
			V. Ashkenazy	007-b	POCL-5024	Tr.07
R. Schuman (05) (1 performances)						
	No. 39	kdz007	Kinderszenen Op. 15, No. 7 "Träumerei"			
			V. Ashkenazy	007-b	POCL-5106	Tr.07

Table 2. CrestMusePEDB ver. 2.0 Performance Data (No. 40-60)

Composer (ComposerID) (amount of performances)						
	PerfID	ScoreID	Title			
W. A. Mozart (02) (6 performances)						
	No. 40〜42	snt011-1	Piano Sonata No. 11, K. 331, 1st Mov.			
			C. Eschenbach	010-a	UCCG-5029	Tr.07
			I. Haebler	011-a	COCQ-83691	Tr.07
			L. Kraus	012-a	SICC-487	Tr.04
	No. 43	snt001-1	Piano Sonata No. 1, K. 279, 1st Mov.			
			G. Gould	003-a	SRCR-9667	Tr.01
	No. 44	snt001-2	Piano Sonata No. 1, K. 279, 2nd Mov.			
			G. Gould	003-a	SRCR-9667	Tr.02
	No. 45	snt001-3	Piano Sonata No. 1, K. 279, 3rd Mov.			
			G. Gould	003-a	SRCR-9667	Tr.03
L. V. Beethoven (03) (6 performances)						
	No. 46	snt008-2	Piano Sonata No. 8, Op. 13, 2nd Mov.			
			V. Ashkenazy	007-a	POCL-5005	Tr.08
	No. 47	snt008-3	Piano Sonata No. 8, Op. 13, 3rd Mov.			
			V. Ashkenazy	007-a	POCL-5005	Tr.09
	No. 48	snt014-1	Piano Sonata No. 14, Op. 27-2, 1st Mov.			
			B. Brendel	008-a	438-862-2	Tr.09
	No. 49	snt014-2	Piano Sonata No. 14, Op. 27-2, 2nd Mov.			
			B. Brendel	008-b	438-862-2	Tr.10
	No. 50	snt014-3	Piano Sonata No. 14, Op. 27-2, 3rd Mov.			
			B. Brendel	008-a	438-862-2	Tr.11
	No. 51	snt017-1	Piano Sonata No. 17, Op. 31-2, 1st Mov.			
			M. Pollini	013-a	UCCG-7069	Tr.02
F. Chopin (04) (9 performances)						
	No. 52	wlz001	Waltz Op. 18, No. 1			
			V. Ashkenazy	007-a	POCL-5024	Tr.01
	No. 53	wlz003	Waltz Op. 34-2, No. 3			
			V. Ashkenazy	007-a	POCL-5024	Tr.03
	No. 54	wlz009	Waltz Op. 69-1, No. 9			
			V. Ashkenazy	007-a	POCL-5024	Tr.09
	No. 55	wlz010	Waltz Op. 69-2, No. 10			
			V. Ashkenazy	007-a	POCL-5024	Tr.10
	No. 56	etd003	Etude Op. 10, No. 3			
			V. Ashkenazy	007-a	POCL-5046	Tr.03
	No. 57	etd004	Etude Op. 10, No. 4			
			V. Ashkenazy	007-a	POCL-5046	Tr.04
	No. 58	etd023	Etude Op. 25-11, No. 23			
			V. Ashkenazy	007-a	POCL-5046	Tr.23
	No. 59	nct002	Nocturne Op. 9-2, No. 2			
			V. Ashkenazy	007-a	POCL-3880	Tr.02
	No. 60	nct010	Nocturne Op. 32-2, No. 10			
			V. Ashkenazy	007-a	POCL-3880	Tr.10

Table 3. Players' list included in ver. 1.0 and 2.0.

No.	Player's name	Included performances
001	Andras Schiff	5
002	Sviatoslav Richter	12
003	Glenn Gould	9
004	Friedrich Gulda	2
005	Hiroko Nakamura	3
006	Norio Shimizu	1
007	Vladimir Ashkenazy	19
008	Alfred Brendel	4
009	Martha Argerich	1
010	Christoph Eschenbach	1
011	Ingrid Haebler	1
012	Lili Kraus	1
013	Maurizio Pollini	1

analysts can work more objectively by referring to qualified performance expression data.

Music pedagogy in performance expression To quantify the information on performance expression clarifies its features and enables showing us the features objectively. CrestMusePEDB can be applied to a music-learning system in music education and at rehearsal sessions for players.

Supporting the performance expression design CrestMusePEDB will contain performance expression data of over 100 pieces of music. The database will likely be used as a data set for learning expressive performances in a rendering system and for referenced performances in a case-based music system. CrestMusePEDB will contain not only performance expression data but also musical structure information. Research on performance rendering and music analysis progress can be conducted by utilizing both the performance expression data and the musical structure information.

5.2 Future works

Handling damper pedal It can be impossibly difficult to extract detailed damper pedaling information, including

Table 4. A part of music pieces the future versions will include.

Composer	Titles
J. S. Bach	Sinfonia No. 3, 5, 8, 11
W. A. Mozart	Sonata No.8, K. 310
L. v. Beethoven	Sonata No. 9, 17, 19, 20, 23
F. Chopin	Mazurkas No. 5, 7, 13, 19, 23, 38
	Polonaise No. 3, 6
J. Brahms	Rhapsody Op. 79, No. 1, 2
	Rhapsody Op. 119-4
R. Schumann	Kinderszenen Op. 13, No. 1
F. Schubert	Moment Musicaux Op. 97, No. 3, 4
G. Fauré	Barcarolle No. 4, 6, 8, 12
C. Debussey	Suite Bergamasque (all)
	Deux arabesques (both)
	Préludes 1er livre VIII 'La fille aux cheveux de lin'
	Préludes1er livre X 'La cathédrale engloutie'
	Préludes2e livre X 'Canope'
	Children's Corner No. 1
E. Granados	12 danzas españolas Op. 37-5
I. Albéniz	Cantos de España Op. 232
S. Rakhmaninov	Prelude Op. 3-2
E. Satie	Trois Gymnopédies No. 1
G. Gershwin	Three Preludes No. 1
S. Prokofiev	Sonata No. 7 (1st & 3rd mov.)
	The Love for Three Oranges, 'March'
M. Ravel	Sonatine (1st mov.)
	Le Tombeau de Couperin I. Prelude

half pedaling, from audio signals. At present, we simply estimate the position as being possible if the player changes his pedaling, depending on the resonance in the audio signal and the pianist's experience for the performance method. Extracting and describing the information of pedal control should be discussed more in the future.

Handling audio (sound) sources In CrestMusePEDB, the velocity data is estimated approximately based on a certain digital audio source. In the future, we should construct an acoustic model of the physical characteristics of the piano and architectural acoustic characteristics. The database should include these description.

Formation of a research community The most meaningful purpose of constructing an open database is to enable a lot of research to be conducted using the database and for the global research area to be developed. At present, CrestMusePEDB mainly consists of the members of the CrestMuse Project. We are preparing some useful tools for writing down musical notation. We intend to create a far-reaching community for CrestMusePEDB.

6 CONCLUSION

We introduced CrestMusePEDB containing a description of the deviation of performance expression by many world virtuosos. This database is available to researchers around the world free of charge through the Internet after the submission procedure is completed. Information on obtaining the database is described at `http://www.crestmuse.jp/pedb/`. We believe that it can be useful for research in, but not limited to, music information retrieval, music appreciation, music structure analysis, music expression analysis, musical instrument analysis/identification, performance rendering systems, and music-related visualization. With this database, researchers can now use virtuosos' performances for each stage of finding problems, obtaining solutions, implementing these solutions, and for conducting evaluations and presentations.

In the future, more expressive performances and various meta-data (descriptions of contents) will need to be added to the database pieces through cooperation with several music databases on the Internet. Also, we are preparing some tools to facilitate using the database such as a data viewer and extracting the deviation data from audio signals so that world-wide users can add data to the database. Our goal for our database is for it to be widely used worldwide and to accelerate progress in this field of research.

7 ACKNOWLEDGEMENT

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BEAT TRACKING USING GROUP DELAY BASED ONSET DETECTION

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ABSTRACT

This paper introduces a novel approach to estimate onsets in musical signals based on the phase spectrum and specifically using the average of the group delay function. A frame-by-frame analysis of a music signal provides the evolution of group delay over time, referred to as phase slope function. Onsets are then detected simply by locating the positive zero-crossings of the phase slope function. The proposed approach is compared to an amplitude-based onset detection approach in the framework of a state-of-the-art system for beat tracking. On a data set of music with less percussive content, the beat tracking accuracy achieved by the system is improved by 82% when the suggested phase-based onset detection approach is used instead of the amplitude-based approach, while on a set of music with stronger percussive characteristics both onset detection approaches provide comparable results of accuracy.

1 INTRODUCTION

The task of estimating the times at which a human would tap his foot to a musical sound is known as beat tracking [1]. All state-of-the-art approaches ([1, 2, 3, 4]) for this task first conduct an onset detection. The output of the onset detection is a signal with a lower time resolution than the input signal, which has peaks at the time instances where a musical instrument in the input started playing a note. Usually, this onset signal is derived from the amplitude of the signal, as in [1, 2, 3]. Only in [4], phase information is considered, by computing the phase deviation between neighboring analysis frames. Several approaches of computing onsets from musical signals are compared in [5], resulting in the conclusion that in general the moments of big positive change in the amplitude spectrum of the signal provide the most preferable estimators for the onsets. This is because using information contained in the complex spectrum or in the phase changes might lead to similar onset estimations, but using only the amplitude spectrum is preferred for computational reasons [5]. A similar conclusion can be drawn from the results of the MIREX 2007 Audio Onset Detection contest [1] , where most systems use only the amplitude information. Considering the results on complex music mixtures, which are the signals of interest in beat tracking, the usage of phase deviation by some systems does not improve the onset estimation accuracy [6].

As can be seen in the results depicted in [7] (Table II), the state-of-the-art approaches for beat tracking decrease significantly in accuracy, when applied to folk music. These music signals contain weaker percussive content than music of rock or disco styles. This problem is of particular importance when dealing with traditional dances as well, as they are often played using string or wind instruments only [8]. Based on the results obtained in [7], it is necessary to improve beat tracking on music with little percussive content. While a decrease in the case of jazz and classical music can partly be attributed to the complex rhythmic structure, rhythmic structure of folk music is simpler, and thus the decrease in this forms of music may be attributed solely to the problem of detecting onsets.

In [9], the negative derivative of the unwrapped phase, *i.e.* the group delay, is used to determine instants of significant excitation in speech signals. This approach has been further developed and used for the detection of clicks of marine mammals in [10]. There, it has been shown that pulses can be reliably detected by using group delay, even when the pulses have been filtered by a minimum phase system. Motivated by these works, we suggest the use of the group delay function for onset estimation in musical signals, and then use this estimation as input to a beat tracker based on the state-of-the-art system presented in [2]. The goal of this paper is to provide an improved beat tracking performance on musical signals with simple rhythmic structure and little or no percussive content. The proposed approach to consider phase information is novel in Music Information Retrieval, as previous approaches computed a time derivative of phase [5], while the suggested approach makes use of group delay, which is a derivative of phase over *frequency*. The group delay function is computed in frame-by-frame analysis and its average is computed for each frame. This results in time-domain signal referred to as phase slope function [10]. Onsets are then simply estimated by detecting the positive zero-crossings of the phase slope function. Therefore, the suggested approach does not require the use of time-dependent (adaptive) energy-based thresholds as the ampli-

[1] http://www.music-ir.org/mirex/2007/index.php/Audio_Onset_Detection

tude and phase based approaches for detecting the onsets [5].

In Section 2, the characteristics of the group delay function for minimum phase signals are shortly reviewed to support our motivation. In Section 3 we present a method to compute an onset signal for music based on the phase slope function and how this information has been incorporated into the state-of-the-art system suggested by Klapuri et al. [2] for beat tracking. Section 4 shows results of the proposed approach on artificial and music signals comparing the suggested phase-based approach with a widely used amplitude-based approach. Section 5 concludes the paper and discusses future work.

2 BASIS FOR THE PROPOSED METHOD

Consider a delayed unit sample sequence $x[n] = \delta[n - n_0]$ and its Fourier Transform $X(\omega) = e^{-j\omega n_0}$. The group delay is defined as:

$$\tau(\omega) = -\frac{d\phi(\omega)}{d\omega} \tag{1}$$

so the group delay for the delayed unit sample sequence is $\tau(\omega) = n_0 \ \ \forall\omega$, since the phase spectrum of the signal is $\phi(\omega) = -\omega n_0$. The average over ω of $\tau(\omega)$ provides n_0 which corresponds to the negative of the slope of the phase spectrum for this specific signal and to the delay of the unit sample sequence. An example of a delayed unit sample sequence with $n_0 = 200$ samples as well as the associated group delay function are depicted in Fig.1(a) and (b), respectively. In the above example the Fourier Transform has been computed considering the center of analysis window to be at $n = 0$. When the window center is moved to the right (closer to the instant $n = n_0$), the slope of the phase spectrum is increased (the average of the group delay function is decreased) reflecting the distance between the center of the analysis window and the position of the impulse. When the center of the analysis window is at $n = n_0$ then the slope is zero. Continuing moving the analysis window to the right the slope will start to increase (while the average of the group delay will decreased). In this way, the slope of the phase spectrum is a function of n. Note that the location of the zero-crossing of this function will provide the position of the non-zero value of the unit sample sequence independently of the amplitude value of the impulse.

In general, the average value of the group delay is determined by the distance between the center of the analysis window and the delay of the unit sample sequence, even when it has been filtered by a minimum phase system [9]. The group delay function will still provide information about this delay value as well information about the poles of the minimum phase system. In Fig. 1(c),(d) the output of the minimum phase signal and the associated group delay are depicted. The slope function will have a similar behavior to this described earlier for the unit sample sequence.

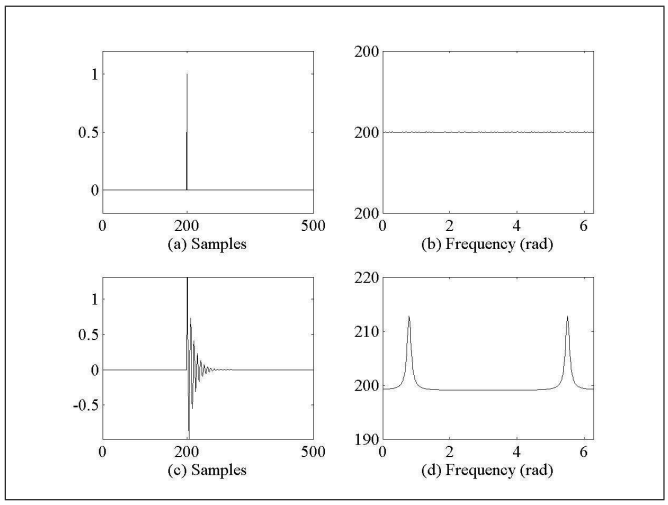

Figure 1. (a) A delayed by 200 samples unit sample sequence. (b) The group delay function of the signal in (a). (c) A minimum phase signal with an oscillation at $\pi/4$. (d) The group delay function of the signal in (c).

In Figure 2, the phase slope of a periodic sequence of minimum phase signals is depicted. The dash-dotted line depicts a phase slope resulting from a window shorter than the period of the signal, the dashed line results from an analysis using a longer window. The phase slope values have been assigned to the middle of the analysis window, the analysis step size was set to one sample. It can be seen that even in the case of low signal amplitude, the positive zero crossing coincides in each case with the beginning of the minimum phase signal. As a musical instrument could be considered as a causal and stable filter, that is driven by a minimum phase excitation, it can be assumed that an onset estimation using the positive zero crossings of the phase slope is a valid approach. To avoid the problems of unwrapping the phase spectrum of the signal for the computation of group delay, the slope of the phase function can be computed as [11]:

$$\tau(\omega) = \frac{X_R(\omega)Y_R(\omega) + X_I(\omega)Y_I(\omega)}{|X(\omega)|^2} \tag{2}$$

where

$$\begin{aligned} X(\omega) &= X_R(\omega) + jX_I(\omega) \\ Y(\omega) &= Y_R(\omega) + jY_I(\omega) \end{aligned}$$

are the Fourier Transforms of $x[n]$ and $nx[n]$, respectively. The phase slope is then computed as the negative of the average of the group delay function.

3 METHOD FOR BEAT TRACKING

3.1 Onset detection using group delay

The onset detection using group delay follows the concept explained in Section 2. The parameters of the onset detector

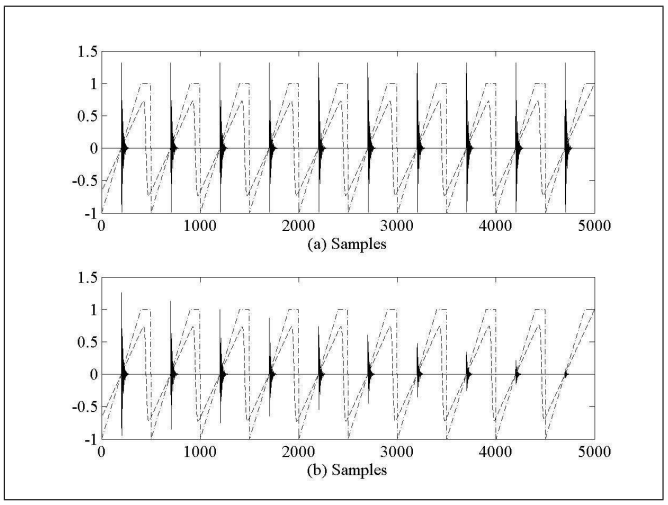

Figure 2. (a) A sequence of impulses of constant amplitude and the associated phase slope function using long (dashed line) and short (dash-dotted line) window (b) A sequence of impulses with linearly time varying amplitudes and the associated phase slope function using long (dashed line) and short (dash-dotted line) window.

have been evaluated on two development data sets: The first data set, referred to as D1, consists of periodic artificial signals like those depicted in Figure 2, with periods from 0.3s to 1s, which is related to the typical range for the tempo of musical pieces (60bpm-200bpm). This data set is also useful in evaluating the robustness of the suggested approach against additive noise. For this purpose, a Transient to Noise Ratio (TNR) is defined in the same way as the usual Signal to Noise Ratio (SNR):

$$TNR = 10 \log_{10} \frac{\frac{1}{L} \sum_{n=0}^{L-1} x^2(n)}{\sigma_u^2} \qquad (3)$$

where x denotes a signal of length L and σ_u^2 denotes the variance of the noise. The artificial signals have been mixed with white noise at different TNR. For each artificial signal a corresponding text file has been created containing the onset times of the impulses. The second development set, referred to as D2, is a data set of 28 song excerpts of 30 seconds length. This data set has been compiled and beat annotated by the authors. From these annotations, a unit sample sequence, $\mathbf{a}[n]$, may be obtained with pulses located at the annotated onset or beat time instance. In the same way, a unit sample sequence $\mathbf{y}[n]$ may be generated from the estimated onset times. The quality of an onset estimation was judged based on the function used in the MIREX 2006 Audio Beat Tracking contest [2] :

$$P_M = \frac{1}{S} \sum_{s=1}^{S} \frac{1}{N_P} \sum_{m=-W}^{W} \sum_{n=1}^{N} \mathbf{y}[n] \mathbf{a}_s[n-m] \qquad (4)$$

[2] http://www.music-ir.org/mirex2006/index.php/Audio_Beat_Tracking

where N is the length of the two pulse trains in samples, S is the number of different annotations per sound sample (equal to one for the development data), N_P is the maximum number of impulses in the two pulse trains, $N_P = \max(\sum \mathbf{y}[n], \sum \mathbf{a}_s[n])$, and W is equal to 20% of the average distance between the impulses in $\mathbf{a}_s[n]$ in samples. This function represents an estimator, of how much two pulse trains are correlated, accepting some inaccuracy regarding the placement of the onset estimation impulses. Note that for the development set containing music signals (*i.e.*, D2), no onset annotations exist but beat annotations.

The most crucial parameter in the click or onset detection using phase slope is the length of the analysis window. As it is indicated in [10], a large window is appropriate for detecting major sound events in an audio signal while shorter windows may be used in case additional sound events are needed to be detected. In this paper, the optimum length of the analysis window has been determined by trials and errors on the two (development) training data sets, D1 and D2. Figure 3 shows the phase slopes from a short excerpt of a music sample from D2 computed with three different analysis window lengths. The optimum analysis window was found to be 0.2s, thus slightly shorter than the shortest considered signal period (i.e., 0.3 s). A typical window like Hanning was used while the step size of the analysis was set to 5.8ms which corresponds to a sample rate, f_l, of 172 Hz for the onset signal, as suggested in [2].

Two further refinements of the approach as explained in Section 2 have been found to be necessary for music signals. The first is a zero crossing selection. In Figure 3 it can be observed that the shorter the analysis window the more positive zero-crossings are detected. It was observed that accepting only the zero crossings that are surrounded by large oscillations improves the accuracy on the development sets. Such oscillations can easily be detected by thresholding, as shown by the dotted lines in each sub plot of Figure 3. The positive threshold has been determined by the mean of the absolute values of the phase slope for a whole sample; the negative threshold is simply the negative of this value. A zero-crossing is selected if the minimum and the maximum amplitude of the phase slope function, before and after the zero-crossing, respectively, pass the corresponding thresholds. For example, the zero crossing at sample 800 in the middle plot was rejected, because it is not followed by a large maximum.

The second refinement is the usage of a multi band analysis. Dividing the spectrum into a number of bands has been shown to be meaningful for beat tracking [12, 13]. For this, the spectrum has been divided into four equally sized bands on logarithmic frequency scale. In each band, an onset detection using the phase slope method was performed. In order to get a single vector representation, the four band-wise onset signals, $\mathbf{y}_c[n], c = 1...4$, have been fused in the same

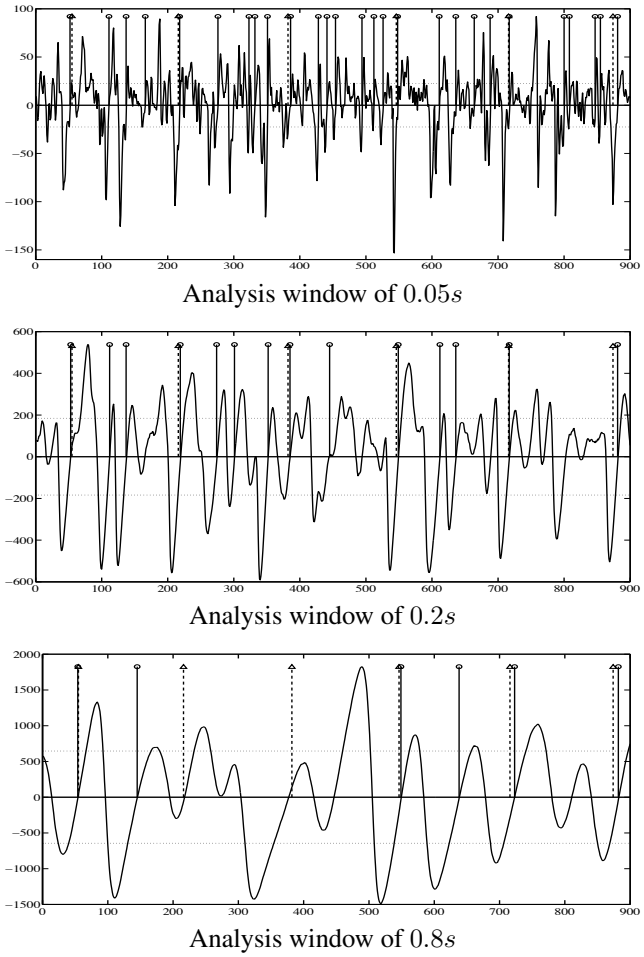

Figure 3. Influence of the analysis window length on the phase slope of a music sample from D2 (x-axis: samples, y-axis: phase slope amplitude, onsets: bold peaks with circle markers, annotation: dashed peaks with triangle markers, threshold for zero crossing selection: dotted lines)

way as in [2]:

$$\mathbf{y}[n] = \sum_{c=1}^{4}(6-c)\mathbf{y}_c[n] \tag{5}$$

giving more weight to lower bands. Indeed, the subband splitting as well as the fusion of bands have been found to improve the performance of the beat tracker as measured by (4).

3.2 Beat tracking

For the estimation of beat times from the band-wise onset signals an algorithm based on the method proposed by Klapuri et al. in [2] has been used. The algorithm had to be adapted to the type of onset signals that are obtained using the phase slope function. This modified beat tracker will be referred to as M-KLAP in the rest of the paper. Experiments for the development of M-KLAP have been conducted using the music data set, D2, described in Section 3.1.

3.2.1 Beat Period

For beat period estimation, Klapuri et al. suggest the computation of comb filter responses on each of the four bands separately, and summing afterwards. In M-KLAP, the band wise onset signals, $\mathbf{y}_c$, are simply summed using (5). Afterwards, the obtained onset vector is weighted with the sum of the spectral flux at each sample n:

$$\mathbf{y}_{flux}[n] = \mathbf{y}[n]\sum_{\omega} HWR(|X(\omega,n)| - |X(\omega,(n-1))|) \tag{6}$$

where HWR denotes a half wave rectification and $X(\omega, n)$ denotes the (short time) Fourier transform of the signal as used in the group delay computation in (2). Weighting the detected onsets in this way slightly but consistently improves performance.

The sample autocorrelation of the vector $\mathbf{y}_{flux}[n]$ is then computed in a rectangular window of $t_{win} = 8s$ length with a step size of one second. The maximum lag considered is $4s \times f_l$, which is equal to 688, since $f_l = 172Hz$. The centers of the analysis windows are positioned at times $[1s, 2s, ..., T_N]$, where $T_N = \lfloor N/f_l \rfloor$, zero padding has been applied. In the following, the beat periods β have been estimated using a *Hidden Markov Model*(HMM) as described in [2], where the beat period is referred to as tactus period. This results in a sequence of beat period estimations $\beta[k]$, with $k = 1...T_N/s$. The only change in the HMM is the use of flat priors for the beat periods, and that the beat periods do not depend on the simultaneous measure periods ((24) in [2]) in the Viterbi algorithm.

3.2.2 Beat Phase

In the phase estimation of the beat pulse ((27) in [2]), the computation of the likelihood of a phase $\Phi[k]$ in analysis frame k has been changed to

$$P(\hat{\mathbf{r}}_{\tilde{\mathbf{y}}\mathbf{y}}|\Phi[k]=l) = \sum_{c=1}^{4}(6-c)\sum_{n=0}^{8f_l}\tilde{\mathbf{y}}_k[n+l]\mathbf{y}_c[kf_L+n-4f_l] \tag{7}$$

where $\tilde{\mathbf{y}}_k$ is a reference pulse train of $t_{win}f_l + 1$ samples length, having an impulse at the middle position and a period equal to $\beta[k]$. Thus, just like in the estimation of the beat period, an eight second length window has been used. The weighted sum of the band wise correlations as computed in (7) is then used in an HMM framework as suggested in [2]. Again, incorporating spectral flux as in (6) has been tried, combined with an impulse selection instead of a Viterbi in order to get the final beat pulse. This would

have the advantage of avoiding the relatively long (8s) analysis window. However, results were slightly inferior to those achieved using the Viterbi algorithm. Because of this, this improvement was postponed for now. Note that the accuracy of measure and tatum periods [2] have not been evaluated, as the focus is the derivation of the beat information.

4 EXPERIMENTS

This Section compares the performance of the system as suggested by Klapuri et al.[2], denoted as KLAP, with the performance phase slope detected onsets as input to the modified beat tracker, which will be referred to as PS/M-KLAP. Two data sets of beat annotated pieces of music have been used for evaluation. The first has been used as a training set for the MIREX 2006 Audio Beat Tracking task [3], and consists of twenty 30 second excerpts from popular music songs. Each song has been beat annotated by several listeners, who were asked to tap the beat of the piece of music. In the following, this data set is referred to as T1. The second data set, (T2), consists of twenty 30 second excerpts from pieces of traditional Cretan music, downloaded from the Institute of Mediterranean Studies [4]. The beat for these pieces has been annotated by the first author. In contrast to T1, none of the songs contain percussive instruments, but only string instruments and vocals. It is worth to note, that none of the mentioned data sets have been used to find the optimal parameters of the system. For this purpose, only the development sets, D1 and D2, mentioned in Section 3, have been used.

As detailed in [2], the most appropriate estimator for the performance of a beat tracking system is the length of the longest continuous correct estimated section of the song, divided by the duration of the whole song. For example, for 30s duration of a song and 12s to be the longest continuously correct beat estimation duration, the accuracy is 40%. Furthermore, the beat estimation is judged as correct when its period is half or double the period of the annotation as well. A deviation of 0.175 times the annotated period length is tolerated. Note that a beat pulse train with the same period as the annotation pulse train is considered as incorrect whenever it has a half period offset (*off-beat*). Accuracies measured with this method will be referred to as A_{cont}. For convenience, also the accuracies as computed by (4) will be shown, denoted as A_{mir}, in order to be able to compare with scores achieved at the MIREX contest.

4.1 Proof of concept

In this Section, the KLAP and PS/M-KLAP beat trackers are applied to D1, the development set containing artificial signals. For each TNR level, the accuracies of the two beat tracking systems have been computed using (4) for all the signal periods in D1 (0.3s to 1s). Then the mean values and the standard errors have been computed for each TNR level. The mean accuracy values along with their corresponding standard errors, shown as error bars, are depicted in Figure 4. Without the addition of noise both approaches estimate a beat pulse train that is perfectly correlated with the position of the impulses in the signal. When the TNR decreases, the presented approach PS/M-KLAP is persistently more accurate. This proofs the hypothesis, that using the proposed approach, beat tracking will be more robust against noise which is important if an audio recording is noise corrupted. Also the presence of noise makes some of the possible percussion components found in music to soften. Based on the above results we expect the proposed approach to be also appropriate for musical signals without strong percussive components.

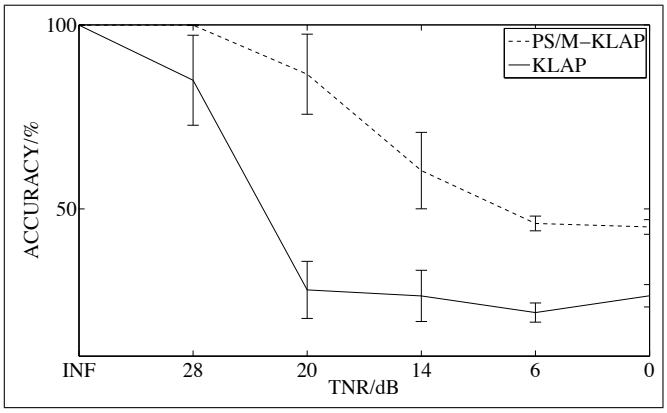

Figure 4. Accuracy of the beat tracking using the proposed method (PS/M-KLAP) and the algorithm of [2] (KLAP), on artificial signals of varying TNR

4.2 Results

The accuracies of the beat trackers applied to the music data sets T1 and T2 are depicted in Tables 1 and 2 for the accuracy measures A_{cont} and A_{mir}, respectively. On T1, the KLAP beat tracker is superior. The advantage of using the phase slope in the proposed way is distinct on T2. Here, the improvement compared to the state-of-the-art approach is 82%. This shows that using the proposed method, beat tracking in a signal with weak or no percussive content can be improved, while approaches using the amplitude information clearly fail on this data. Since the difference between the KLAP and PS/M-KLAP system, doesn't solely rely on the onset detection approach (there are modifications in beat tracking approach as well), it may be assumed that the differences in accuracy cannot be solely attributed to the onset detection method. To check this we decided to provide as input to the M-KLAP system the standard input of the KLAP

[3] http://www.music-ir.org/mirex/2006/index.php/Audio_Beat_Tracking
[4] http://gaia.ims.forth.gr/portal/

	PS/M-KLAP	KLAP
T1	54.3(0.058)	58.4(0.063)
T2	48.0(0.171)	26.3(0.122)

Table 1. Accuracies A_{cont} of the beat tracking on the two data sets, mean value/%(variance)

	PS/M-KLAP	KLAP
T1	45.0(0.028)	50.0(0.026)
T2	39.3(0.071)	21.4(0.085)

Table 2. Accuracies A_{mir} of the beat tracking on the two data sets, mean value/%(variance)

system, *i.e.* that derived from spectral flux [2]. For T1 and T2 data sets, the obtained results are $48\%/43.1\%$ and $22\%/27.8\%$, respectively for A_{cont}/A_{mir} measures. This shows the importance of the phase slope function in onset detection and in the context of beat-tracking. The slightly lower performance of PS/M-KLAP in T1 as compared to the standard system (KLAP) may be attributed to the implementation of beat tracking and we expect to further improve that part in the near future.

5 CONCLUSIONS

In this paper a new method to detect onsets using the average of the group delay was introduced and evaluated in a beat tracking framework. Advantages are the immediate detection of the onsets by locating the positive zero crossings of the phase slope, and the robustness for signals with little percussive content as shown in the experiments. The next steps to improve the method are a more efficient implementation of the phase slope computation and refinements of the beat tracker. Also, evaluation on different data sets of folk music will be performed.

6 ACKNOWLEDGEMENTS

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Structured Polyphonic Patterns

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ABSTRACT

This paper presents a new approach to polyphonic music retrieval, based on a structured pattern representation. Polyphonic patterns are formed by joining and layering pattern components into sequences and simultaneities. Pattern components are conjunctions of features which encode event properties or relations with other events. Relations between events that overlap in time but are not simultaneous are supported, enabling patterns to express many of the temporal relations encountered in polyphonic music. The approach also provides a mechanism for defining new features. It is illustrated and evaluated by querying for three musicological patterns in a corpus of 185 chorale harmonizations by J.S.Bach.

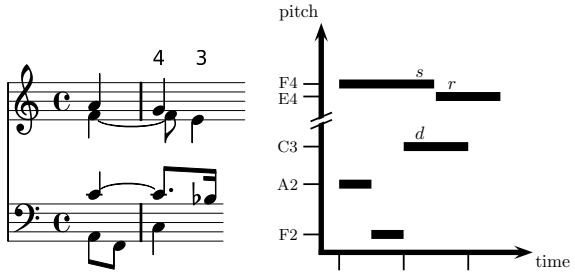

Figure 1. A 4-3 suspension between bass and alto voices in bars 16-17 of Bach's chorale BWV 283 (left) and a piano-roll representation of the alto and bass voices (right).

1 INTRODUCTION

1.1 Motivation

The concept of pattern in music is very important for diverse computational problems: database query and indexing; statistical modeling of musical style; and computer assisted musical analysis. Recently, there have been efforts to extend monophonic patterns to the general polyphonic case. Though there is agreement on the expressive requirements of monophonic patterns (e.g. transposition invariance), a consensus is yet to emerge in the polyphonic case. It is clear that patterns must be able to represent sequences of simultaneous events (simple first species counterpoint) and note against note counterpoint (second, third species). In addition, it is important that patterns represent relations between events that are overlapping but not simultaneous (fourth, fifth species).

Formalizing the knowledge found in treatises on counterpoint is a promising way to propose key features of polyphonic music, which can then be reused in other applications such as machine learning. This paper introduces structured polyphonic patterns ($\mathcal{SPP}$) and shows how they can be used to define polyphonic features and patterns for the musicological notions of parallel fifth, suspension, and cadential voice leading.

The concept of a *suspension* in counterpoint provides a useful case study for the expressive power required for polyphonic patterns. In a suspension (see Figure 1, left), a dissonance formed on a strong beat is resolved by step to a consonance. The pattern presented here is restricted to a simpler "proto-suspension" pattern for ease of exposition. The temporal relations and consonance and dissonance are captured, but features that refer to strong and weak beats, stepwise resolution, and relative duration of preparation and resolution notes are omitted.

Figure 1 (right) shows the notes involved in the suspension in piano-roll notation. The pattern involves three notes: the suspended note s; the note d introducing a dissonance; and the resolution note r (respectively F4, C3 and E4). A suspension pattern must precisely represent the temporal relations between these three notes. In particular, the pattern must represent that d starts while s is unfolding and that r starts while d is unfolding. At the same time, the pattern must represent harmonic pitch class intervals and also group these into classes of consonant and dissonant intervals. In addition, the suspension might occur between any two voices in a multiple voice texture, hence the pattern must capture voice combinations.

The Humdrum toolkit [5] supports polyphonic pattern representation and retrieval. However, the development of a polyphonic pattern can be a prohibitively complex task. To represent the suspension, the encoding of the source needs to be first transformed to a form where a single regular expression can test for the "starts while" temporal relation and

bass	tenor	alto	sop.
8AA	[4c	[4f	4a
8FF	.	.	.
=	=	=	=
4C	8.c]	8f]	4g
.	.	4e	.

bass	alto	hint	pc	cons
AA	f	-20	8	T
FF	(f)	-24	0	T
C	(f)	-17	5	F
(C)	e	-16	4	T

```
[a-g A-G]+[- # n]*[^)]*[(]   [a-g A-G]+[- # n]*.*F$
[a-g A-G]+[- # n]*[)]   [^(]*[a-g A-G]+[- # n]*.*T$
```

Figure 2. Hudrum/kern representation of the 4-3 suspension of Figure 1 before (top left) and after (top right) preprocessing. Simplified suspension pattern in Humdrum (bottom).

the occurrence of a consonance. Consider, for example, the last line of the top left of Figure 2. The E4 in the alto voice starts while the C3 in the bass voice is unfolding, introducing a consonant interval. The null token ".", used to indicate that C3 is still unfolding, must be replaced by the duplicated C3 (to allow harmonic interval computation), but also placed in round brackets to indicate that it is a continuation and does not represent the onset of a new event (this can be done with the Humdrum "ditto" command). Several additional preprocessing steps are necessary, for example: ties must be removed (otherwise the second component of a tie will not appear as an unfolding event); slurs must be removed (as slurs are also denoted by round brackets); and lines containing only null markers "." must be removed (otherwise, a suspension could span three lines instead of two).

Once preprocessing steps are executed, the musical content at the top left of Figure 2 is transformed into a form ready for querying, shown at the top right of Figure 2, which includes all additional columns needed in the pattern matching phase. These additional columns can be computed using Humdrum commands such as "hint", "pc", and "recode".

The pattern matching itself is executed by searching for two successive lines that respectively satisfy the regular expressions shown at bottom of Figure 2. Note that this will only find instances in which the d note appears in the first column (the bass voice in Figure 2). To capture every suspension in Humdrum, one must iterate the preprocessing steps, extracting every two voice combination, and testing for both ordering of those two voices. This must be programmed at a scripting language level above Humdrum.

2 METHODS

As the example above illustrated, the expression of even a simple polyphonic pattern is difficult in Humdrum. A central aim of this research is a pattern description and matching algorithm that is a more usable and efficient alternative to Humdrum. This section describes $\mathcal{SPP}$, a polyphonic pattern language inspired by algebraic representations of music [1, 4] and music knowledge representation methods [2].

2.1 Pattern components

Pattern components in $\mathcal{SPP}$ are represented using *feature sets*. A feature f has a *feature name* τ, a *feature value* v, and optionally a *voice* γ:

Definition 1 $\quad f \quad ::= \quad \tau : v \quad | \quad \tau(\gamma) : v$

A specific feature name is taken from a set of feature names (e.g. `pitch`). A feature value is either a value (e.g. 60) or a value variable. A voice is either a voice name or a voice variable. A feature of the form $\tau : v$ can be primitive (e.g. $\texttt{pitch} : 60$) or can encode a relation with an event occurring in the same voice (e.g. melodic interval $\texttt{mi} : -1$). A feature of the form $\tau(\gamma) : v$ is used to encode a relation with an event occurring in some other voice; for example, the feature $\texttt{vi}(\texttt{alto}) : -5$ encodes a vertical interval of -5 with some event in the alto voice.

A *pattern component* α has a feature set and a voice:

Definition 2 $\quad \alpha \quad ::= \quad \{f, \ldots, f\}_\gamma$

Example *E1* below illustrates a component containing two value variables (in upper case) and one voice variable (in lower case):

$$\{\texttt{duration} : D,\ \texttt{pitch} : P\}_y \qquad \textit{(E1)}$$

An *event* is a component with no variables, whose feature set uses at least the following feature names: `onset`, `duration` and `pitch` (omitted if the event is a rest). For example, the following event encodes the r note of Figure 1 (using MIDI pitch numbers and MIDI ticks at a resolution of 24 ticks per quarter note):

$$\{\texttt{onset} : 36,\ \texttt{duration} : 24,\ \texttt{pitch} : 64\}_{\texttt{alto}} \qquad \textit{(E2)}$$

A component α *matches* an event if, after assignments of the variables of α to values, the feature set of α is a subset of the feature set of the event and their voices are equal. For example, the component *E1* matches the event *E2* with the variable assignments $D \mapsto 24$, $P \mapsto 64$ and $y \mapsto \texttt{alto}$.

2.2 Pattern construction

Patterns in $\mathcal{SPP}$ are formed by joining components sequentially using the ";" operator and by layering components vertically using the "$=\!=$" operator. In addition, a component

may be modified by the "$-$" operator. Formally, the set of $\mathcal{SPP}$ patterns is defined inductively as follows:

Definition 3

$$\begin{array}{rcl} \phi & ::= & \alpha \\ & | & -\alpha \\ & | & \phi \,;\, \phi \\ & | & \dfrac{\phi}{\phi} \end{array}$$

To illustrate $\mathcal{SPP}$ semantics, the "proto-suspension" pattern will now be developed in multiple steps. First, the following pattern captures the sequence formed by the s and r notes in Figure 1:

$$\{\texttt{pitch} : 65\}_{\texttt{alto}} \;\; ; \;\; \{\texttt{pitch} : 64\}_{\texttt{alto}} \qquad \textit{(E3)}$$

This construction enforces a single temporal relation: the event matching the left component must directly precede the event matching the right component (i.e. the event matching the left component must end exactly as the event matching the right component starts). To capture both the alto and bass voice of Figure 1 (in particular to capture the d note of the suspension), two layers are joined as follows:

$$\frac{\{\texttt{pitch} : 65\}_{\texttt{alto}}}{\{\}_{\texttt{bass}}} \;\; ; \;\; \frac{\{\texttt{pitch} : 64\}_{\texttt{alto}}}{\{\texttt{pitch} : 48\}_{\texttt{bass}}} \qquad \textit{(E4)}$$

This construction enforces four temporal relations: i) the event matching the top left component must directly precede the event matching the top right component; ii) the event matching the bottom left component must directly precede the event matching the bottom right component; iii) the events matching the top and bottom right components must start together; and iv) the events matching the top and bottom left components must start together. Consequently, example *E4* does not correctly capture the temporal relations of a suspension, and the "$-$" operator is used to extend the pattern as follows:

$$\frac{-\{\texttt{pitch} : 65\}_{\texttt{alto}}}{-\{\}_{\texttt{bass}}} \;\; ; \;\; \frac{\{\texttt{pitch} : 64\}_{\texttt{alto}}}{-\{\texttt{pitch} : 48\}_{\texttt{bass}}} \qquad \textit{(E5)}$$

This construction also enforces four temporal relations. The first two are unchanged. As a modified component is layered with a non-modified component (*E5*, right), the third condition becomes: iii) the event matching the top right component must start while the event matching the bottom right component is unfolding. As two modified component are layered (*E5*, left), the fourth condition becomes: iv) the events matching the top and bottom left components must overlap. Note how the "$-$" does not represent a tie between the two bass components. The construction correctly captures the temporal relations between the s, d and r notes in the suspension example of Figure 1. Example *E5* can now be generalized to account for transposition invariance. This is done by replacing the concrete pitch features by more abstract vertical interval features:

$$\frac{\boxed{s}}{\dfrac{-\{\}_{\texttt{alto}}}{-\{\}_{\texttt{bass}}}} \;\; ; \;\; \frac{\boxed{r}}{\dfrac{\{\texttt{vi}(\texttt{bass}) : -16\}_{\texttt{alto}}}{-\{\texttt{vi}(\texttt{alto}) : 17\}_{\texttt{bass}}}} \qquad \textit{(E6)}$$

$$\boxed{d}$$

The feature $\texttt{vi}(\texttt{alto}) : 17$ specifies that the d note forms a vertical interval of 17 semitones with some overlapping note in the alto voice. This correctly represents the vertical interval between the d note and the s note. However, it lacks precision as it could also represent an interval between the d note and the r note. The feature $\texttt{vi}(\texttt{bass}) : -16$ also lacks precision as it specifies that the r note forms a vertical interval of -16 with any overlapping note in the bass voice. Therefore, the vertical interval feature must be specialized to the the "start while" temporal context:

$$\frac{-\{\}_{\texttt{alto}}}{-\{\}_{\texttt{bass}}} \;\; ; \;\; \frac{\{\texttt{sw\_vi}(\texttt{bass}) : -16\}_{\texttt{alto}}}{-\{\texttt{sw\_vi}(\texttt{alto}) : 17\}_{\texttt{bass}}} \qquad \textit{(E7)}$$

The feature $\texttt{sw\_vi}(\texttt{alto}) : 17$ is restricted to vertical intervals formed when the d note starts while some note in the alto voice unfolds. Therefore, it unambiguously represents the interval between the d note and the s note. Similarly, the feature $\texttt{sw\_vi}(\texttt{bass}) : -16$ unambiguously represents the vertical interval between the r note and the d note. Further generalization is achieved by replacing vertical intervals with classes of consonant and dissonant intervals:

$$\frac{-\{\}_{\texttt{alto}}}{-\{\}_{\texttt{bass}}} \;\; ; \;\; \frac{\{\texttt{sw\_cons}(\texttt{bass}) : \texttt{t}\}_{\texttt{alto}}}{-\{\texttt{sw\_dis}(\texttt{alto}) : \texttt{t}\}_{\texttt{bass}}} \qquad \textit{(E8)}$$

The `sw_cons`/`sw_dis` features are similar to the `sw_vi` feature, except that the absolute value of the vertical interval is tested for inclusion (respectively exclusion), modulo twelve, in the following set : $\{0, 3, 4, 7, 8, 9\}$. Finally, voice variables are used to capture suspensions between any two voices, resulting in the final "proto-suspension" pattern *P1*:

$$\frac{-\{\}_{x}}{-\{\}_{y}} \;\; ; \;\; \frac{\{\texttt{sw\_cons}(y) : \texttt{t}\}_{x}}{-\{\texttt{sw\_dis}(x) : \texttt{t}\}_{y}} \qquad \textit{(P1)}$$

In any instance, all three occurrences of the voice variable x (respectively y) must be assigned to the same voice name (i.e. the scope of variables is the whole pattern).

2.3 Defining polyphonic features

A strength of $\mathcal{SPP}$ is that the polyphonic features used in the previous section are given precise formulations in terms of feature definition rules. For example, the `sw_vi` feature is defined as follows:

where	**add**	
$\frac{\{\texttt{pitch} : P\}^*_x}{=\{\texttt{pitch} : Q\}_y}$	$\texttt{sw\_vi}(y) : Q - P$	*(R1)*

The **where** part uses a pattern to indicate that the `sw_vi` feature is formed wherever a note in voice x starts while a note in voice y is unfolding. The value variable P (respectively Q) is used to capture the pitch of the note in voice x (respectively y). The **add** part defines the feature to add. It is evaluated after the variables of the **where** part have been assigned. The resulting feature is added to the event matching the distinguished component in the **where** part (indicated with an asterisk in rule *R1*).

The same mechanism can also accommodate horizontal features, such as the melodic interval:

where	**add**	
$\{\texttt{pitch} : Q\}_x \; ; \{\texttt{pitch} : P\}^*_x$	$\texttt{mi} : P - Q$	*(R2)*

Applying the rules *R1* and *R2* to the source excerpt of Figure 1 would add the following features to the r note:

$$\begin{array}{l} \texttt{mi} : -1 \\ \texttt{sw\_vi}(\texttt{bass}) : -16 \end{array} \qquad \textit{(E9)}$$

Matching the **where** part of a feature definition rule is done with the same algorithm used for $\mathcal{SPP}$ pattern matching.

2.4 Pattern matching algorithm

The matching algorithm proceeds in two phases. In the first phase, the corpus is scanned and every component of the pattern is given an instance list (i.e. a list of matching events and corresponding variable assignments). In the second phase, the pattern is recursively analyzed and instance lists are joined, either according to a ";" operator or a "$=\!=$" operator. Joins that do not respect $\mathcal{SPP}$ semantics or result in inconsistent variable assignments are simply discarded. The joining of two instance lists is done efficiently using data structures that take advantage of the fact that pairs of instances resulting in valid joins are always neighbors in the time dimension.

Figure 3. Examples of the two most frequent kinds of suspension found in the chorales: 2-3 suspension in BWV 272 bar 1 (left), and 4-3 suspension in BWV 262 bar 6 (middle). In addition, a 7-(5)-6 suspension in BWV 328 bars 32-33 (right).

3 RESULTS

An $\mathcal{SPP}$ parser and matching algorithm has been implemented in Ocaml and tested with 3 polyphonic patterns on a corpus of 185 chorale harmonizations by J.S.Bach. The corpus, originally encoded in the Humdrum format, was retrieved from www.kernscores.net [9] and saturated with the features described in Table 1.

3.1 Suspension

A total of 1177 instances of the suspension pattern *P1* developed in Section 2 were found in the corpus. Some instances of the suspension pattern are shown in Figure 3. The first two instances are examples of two of the four most frequent kinds of suspension found: 2-3 (509 instances), 4-3 (278 instances), 7-6 (128 instances) and 4-5 (98 instances). Note how the d note (the note introducing a dissonance) can be either in the upper voice (Figure 3, left) or lower voice (Figure 3, middle): components layered with the "$=\!=$" operator may freely match any voice permutation.

An interesting instance of the pattern is illustrated at the right of Figure 3. In this example, a 7-6 suspension (A4-G4 in soprano) is delayed by an intermediate F4 in the soprano voice, forming a (consonant) interval of a fifth which matches the consonance captured by the `sw_cons` feature found in the top right component of *P1*. If desired, this type of match could be easily removed by introducing a feature representing stepwise motion in the same component.

3.2 Parallel fifth pattern

The second pattern selected for illustration is the parallel fifth, a voice leading structure avoided by Renaissance and tonal composers. In the corpus, 5 instances were discovered. Figure 4 presents two instances. The remaining instances can be found in BWV 263 bar 6, BWV 301 bar 3, and BWV 361 bar 12. The results here are identical (on a smaller data set) to those reported by Fitsioris and Conklin [3] who use

Feature name	Range	Description
`pc`	$\{0,\ldots,11\}$	Modulo twelve of `pitch` (pitch class)
`pc_r`	$\mathbb{B}$	True if the last `pc` is repeated
`mi`	$\mathbb{Z}$	Melodic interval (pitch difference with previous note)
`mi_m`	$\{0,\ldots,11\}$	Modulo twelve of `mi` (pitch class interval)
`vi`(x)	$\mathbb{Z}$	Vertical interval with an overlapping note in voice x
`st_vi`(x)	$\mathbb{Z}$	Vertical interval with a note in voice x that starts together with the current note
`sw_vi`(x)	$\mathbb{Z}$	Vertical interval with a note in voice x that is unfolding while the current note starts
`et_vi`(x)	$\mathbb{Z}$	Vertical interval with a note in voice x that ends together with the current note
`vi_r`(x)	$\mathbb{B}$	True if the last vertical interval with voice x is repeated
`et_vi_am`(x)	$\{0,\ldots,11\}$	Modulo twelve of the absolute value of `et_vi`
`sw_cons`(x)	$\mathbb{B}$	True if `sw_vi`(x) corresponds to a consonant interval
`sw_dis`(x)	$\mathbb{B}$	True if `sw_vi`(x) corresponds to a dissonant interval

Table 1. List of user-defined features used in this paper (top: horizontal features; bottom: vertical features). The ranges $\mathbb{Z}$ and $\mathbb{B}$ respectively refer to integers and booleans.

Figure 4. Selected instances of the parallel fifth pattern: BWV 323 bar 8 (left) and BWV 355 bar 15 (right).

Figure 5. Instance of the cadential pattern in the bass and tenor voices: BWV 293 bar 8

a Prolog encoding of the parallel fifth pattern and provide a musicological interpretation of all instances.

The $\mathcal{SPP}$ pattern of the parallel fifth was carefully designed to avoid false positives (e.g. antiparallel fifths, cases separated by rests):

$$\{\texttt{et\_vi\_am}(y):7\}_x \;;\; \{\texttt{vi\_r}(y):\texttt{t},\ \texttt{pc\_r}:\texttt{f}\}_x \qquad \textit{(P2)}$$

The first component of *P2* captures a vertical interval of a perfect fifth (7 semitones modulo twelve, hence compound fifths are also captured). The second component ensures that the fifth is repeated ($\texttt{vi\_r}(y) : \texttt{t}$ feature) and that there is melodic motion ($\texttt{pc\_r} : \texttt{f}$ feature). The $\texttt{vi\_r}(y) : \texttt{t}$ feature uses a voice variable to ensure that the successive fifths occur between the same two voices (the current voice x and some other voice y).

3.3 Cadential melodic interval pattern

The final pattern reported in this paper was initially reported in the context of the discovery of vertical patterns in the Bach chorales [1]. It was observed that a particular structure of two simultaneous pitch class intervals (2-5 in the bass and 11-8 in a higher voice; Figure 5) occurs at many cadences. Interestingly, in these cadences the leading tone falls to the fifth scale degree rather than rising to the tonic. Here, a two-voice fragment of that pattern is encoded as follows:

$$\frac{\{\}_x}{\{\}_{\texttt{bass}}} \;;\; \frac{\{\texttt{mi\_m}:11\}_x}{\{\texttt{mi\_m}:2\}_{\texttt{bass}}} \;;\; \frac{\{\texttt{mi\_m}:8\}_x}{\{\texttt{mi\_m}:5\}_{\texttt{bass}}} \qquad \textit{(P3)}$$

The query on the corpus returned a total of 70 instances. In most (64) of the instances, the first chord is a ii^6_5. The first instance of Figure 6 illustrates such a case. It contains the pattern in the bass and alto lines, with extensive melodic and rhythmic elaboration occurring in the surrounding tenor and soprano voices. In the remaining 6 cases a IV or IV^7 is formed on the first chord. Figure 6 (right) illustrates such a IV^7 case. Note that these latter cases could easily be excluded by adding the feature $\texttt{st\_vi}(y) : 9$ to the component at the bottom left of *P3*. This would capture the vertical interval of a major sixth (9 semitones) between the bass note and some note in another voice, which has to occur in a ii^6_5 chord.

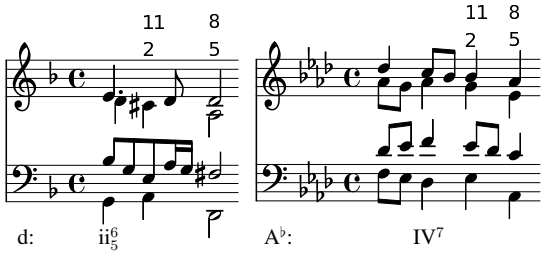

Figure 6. Selected instances of the two-voice cadential pattern: BWV 297 bar 11 (left) and BWV 354 bar 6 (right).

4 DISCUSSION

The paper introduced $\mathcal{SPP}$, a structured polyphonic pattern language and matching algorithm. An important feature of $\mathcal{SPP}$ is that voicing is handled in a general way and that voice permutations are explored automatically. It also provides a simple feature definition mechanism to offer a great deal of flexibility.

Most existing approaches to polyphonic pattern representation lack the expressiveness to accurately capture the patterns discussed in this paper. For example, vertical patterns [1] can only match polyphonic sources that have been expanded and sliced to yield a homophonic texture. A point set pattern representation [7, 8] can only encode a class of suspensions for which the duration ratios and the vertical intervals are always the same (capturing every suspension would require a set of patterns, the size of which can grow quickly as many different ratios and vertical intervals are likely to be found in the source). Techniques that rely on approximate matching to a source fragment [6] can confuse simultaneous notes with notes that overlap without being simultaneous. This can result in a significant loss of precision when retrieving patterns such as the suspension in which events overlap but are not simultaneous. In comparison with Humdrum, $\mathcal{SPP}$ shifts the kind of understanding required from operational (understanding the processing steps required to search for a pattern) to denotational (understanding the meaning of particular features and patterns). This shift offers many advantages, such as the ease to specialize a pattern, and a greater confidence in the precision of the query. Finally, $\mathcal{SPP}$ is much easier to extend for a researcher, as adding a new operator can be done without significantly modifying the pattern matching algorithm.

In the future, the formal properties of the language will be explored, e.g. the possibility of reasoning about pattern equivalence. Also, the expressiveness of $\mathcal{SPP}$ will be formally investigated (including a comparison with Humdrum). Further evaluation of the approach will be achieved by encoding more patterns (e.g. appoggiatura, chained suspensions, neighbor tones and passing tones, cross relation) and analyzing the result of matching in wider more diverse corpora, including corpora of piano music where voices can appear or disappear (this paper used a corpus that had an unchanging four-voice texture). The resulting catalog of polyphonic patterns might be used as global piece features for machine learning. Finally, the topic of pattern discovery based on the $\mathcal{SPP}$ language will be explored.

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STREAMCATCHER: INTEGRATED VISUALIZATION OF MUSIC CLIPS AND ONLINE AUDIO STREAMS

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ABSTRACT

We propose a content-based approach to explorative visualization of online audio streams (e.g., web radio streams). The visualization space is defined by prototypical instances of musical concepts taken from personal music collections. Our system shows the relation of prototypes to each other and generates an animated visualization that places representations of audio streams in the vicinity of their most similar prototypes. Both computation of music similarity and visualization are formulated for online real time performance. A software implementation of these ideas is presented and evaluated.

1 INTRODUCTION

A massive amount of music is already available on the web in the form of internet radio streams. At the time of writing this paper, the "radio-locator" website [1] lists more than 2500 entries in its directory of free internet radio stations, a number that is likely to increase rapidly in the future. Because it is infeasible for users to keep an overview of available radio streams, software that recommends possibly interesting radio streams would be desirable.

When talking about music, humans usually use examples of similar music to describe new and unknown songs or artists. For example, one might characterize *Nirvana* as a combination of the *Pixies'* and *Sonic Youth's* guitar sound, the *Beatles'* sense for catchy melody lines, and the heritage of American folk-rock music in the style of *Neil Young*. So, instead of resorting to predefined taxonomies for music categorization (e.g., by *Genre*), an alternative is to present a "query by example" based user interface directly to the user.

To support the user in finding new music he or she might like in audio streams, we propose a simple interactive visualization approach that incorporates the user's musical vocabulary into the definition of semantic spaces for music similarity judgement. The user defines her own semantic space by supplying, coloring and labeling an arbitrary number of reference songs from her own music collection. In this way, she can define her personal musical concepts via examples she is familiar with.

In our application, songs from personal music collections are imported in an incremental, user-feedback based manner: (1) A new sound clip (concept prototype) is loaded into the system and it is compared to already loaded clips with a content-based music similarity measure, (2) the software shows the user what it thinks the new music is similar to and puts a preliminary label on it, (3) the user refines the software suggestion by correcting the label, (4) go back to (1). When the user decides that the concepts he or she is interested in are sufficiently well represented, unknown music from internet radio streams can be compared to the sound prototypes (using the same similarity metric), and a visualization is generated by means of a multidimensional scaling [4] technique. The visualization also provides direct interaction facilities that allow the playback of streams and clips to interactively explore the audio streams audio-visually.

Additionally to mapping acoustic similarity to proximity information in the 2D visualization, color is used to encode the user's view of what the music sounds like. Colors can be assigned to songs upon loading them into the system by the user. This idea is motivated by the neurologically based phenomenon of *synesthesia* [1, 14], in which stimulation of one sense automatically triggers experiences in another sense.

2 RELATED WORK

Berenzweig et al. [3] construct an *Anchor Space* by training N classifiers to prototypical instances of concepts, classifying unknown music, and mapping posterior probabilities to the individual dimensions of vectors in an N-dimensional space. Distances in this space have been shown to better reflect the user's notion of music similarity.

Lidy and Rauber [7] calculated features derived from long-term observations of (terrestrial) radio stations and used a SOM to generate a 2D visualization of the types of music

[1] http://www.radio-locator.com/

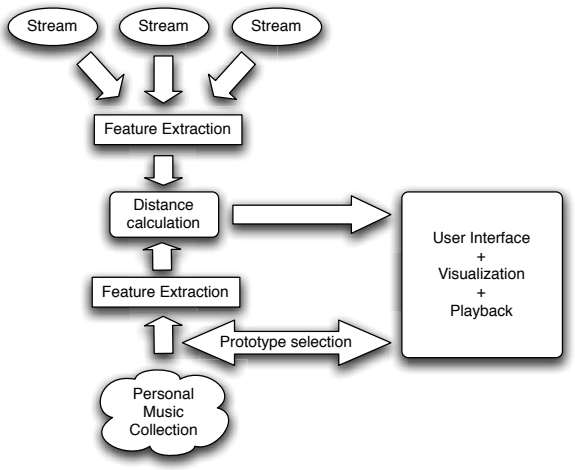

Figure 1. System components

played in different radio stations. However, they do not perform online analysis of radio streams.

Tzanetakis [13] extracted features from audio streams in real time and displayed genre membership of the streams in a real time visualization. By reducing dimensionality of the feature space with a Principal Component Analysis, he also mapped audio signals from high-dimensional timbre-similarity space into a three-dimensional visualization space. To the authors' knowledge, this was the first published implementation of online/real time audio signal classification and similarity analysis.

Lamere [6] uses a technique based on acoustic similarity and multidimensional scaling to visualize and explore music collections in 3D space. This work is related to our work in that it also shows the disparity between acoustic and high-level similarity.

Lübbers [9] derives prototypical songs by hierarchically clustering music collections, and proposes a multi-modal user interface for interactively exploring the cluster centers' neighborhood.

Recently, some work in the field of Human-Computer Interaction has been published (see also [1, 14]) that gives rise to the presumption that organizing music by color is quite intuitive. We also used this idea in our application to identify music that belongs to a common abstract concept, which could be a genre, a particular kind of instrumentation, a certain guitar sound, and so on.

3 SYSTEM OVERVIEW

To evaluate our approach, we implemented an application prototype that performs online (a) feature extraction, (b) similarity calculation, and (c) visualization. Figure 1 sketches the main components of the application.

The feature extraction subsystems are responsible for extracting feature frames from offline clips and online streams of audio data. The distribution of these feature frames is then modeled as a single Gaussian with full covariance [10] per clip/stream.

Central to our framework is the similarity calculation component that calculates distance matrices holding clip-to-clip and stream-to-clip similarities by calculating the symmetric Kullback-Leibler (KL) divergence [5] between each pair of Gaussians.

Distances between clips are then projected to a 2D visualization space with a Multidimensional Scaling [4, 2] algorithm, whereas streams are linearly embedded into a local subspace spanned by the 3 nearest neighbors of a stream model in feature space.

3.1 Implementation notes

The application was implemented in C++ using the Open-Source QT [2] toolkit , and we use OpenGL [3] for the visualizations. All signal processing and statistical computing was implemented within the *FLOWer* framework, a portable and efficient C++ library for data flow-oriented media processing, which is being developed by the first author of this paper.

4 FEATURE EXTRACTION & SIMILARITY CALCULATION

In order to be able to extract features from offline files as well as from live streams, the system contains two feature extraction pipelines. In both pipelines, Mel Frequency Cepstral Coefficients (MFCCs) [8] are computed for each frame of audio data. The main difference between online and offline analysis is the way statistics (means and covariances) are calculated. While in the offline case, the distribution of MFCC's in an entire audio file is estimated, the online scenario requires more diligence; since the distribution of features in an internet radio stream is not likely to stay constant, we only take the most recent feature frames into account. For performance reasons, we decided to use a recursive estimator for means and covariances instead of a sliding window approach.

The input to the feature extraction stage is either an MP3 file or an MP3 stream (which can be transmitted over an HTTP connection). This data is decoded to PCM at a sample rate of 44.1kHz, converted to mono, and sliced into frames of size 2048 samples with 50% overlap. Then, the frames are multiplied with a Hamming window, the magnitude spectrum and MFCCs are calculated, and a statistical model is derived from the data.

[2] http://www.trolltech.com/products/qt
[3] http://www.opengl.org

4.1 Offline processing

In the offline case, the calculation the mean vector μ and the covariance matrix Σ in the stream processing framework is straighforward. Let $\mathbf{X}$ be a vector of random variables, and $\mathbf{x}_i$ a concrete instantiation (sample) of $\mathbf{X}$ (in our case an MFCC vector from frame i).
Since

$$\Sigma = E(\mathbf{X}\mathbf{X}^\top) - \mu\mu^\top \tag{1}$$

and

$$\mu = \frac{1}{n}\Sigma\mathbf{x_i} \tag{2}$$

$$E(\mathbf{X}\mathbf{X}^\top) = \frac{1}{n}\Sigma\mathbf{x_i}\mathbf{x_i}^\top \tag{3}$$

all that is needed is one accumulation vector and one accumulation matrix for the sums and the sums of products of observations, respectively.

4.2 Online processing

Since in an online scenario the complete sequence of feature frames is not available beforehand, we use a recursive estimate of mean and covariance to parameterize the distribution of timbral features (we preferred a recursive estimator to windowed statistics for performance reasons).
The recursive estimate of the mean vector μ is calculated as:

$$\mu_n = (1-\alpha)\mu_{n-1} + \alpha\mathbf{x_n} \tag{4}$$

The covariance matrix Σ of a multidimensional vector of random variables $\mathbf{X}$ can be written as

$$\Sigma = E(\mathbf{X}\mathbf{X}^\top) - \mu\mu^\top \tag{5}$$

$E(\mathbf{X}\mathbf{X}^\top)$ can be estimated recursively as

$$E(\mathbf{X}\mathbf{X}^\top) = R_n = (1-\alpha)R_{n-1} + \alpha\mathbf{x}\mathbf{x}^\top \tag{6}$$

and

$$\Sigma_n = R_n - \mu\mu^\top \tag{7}$$

which can be refactored to

$$\begin{aligned}\Sigma_n &= (1-\alpha)R_{n-1} + \alpha\mathbf{x}\mathbf{x}^\top - \mu\mu^\top \\ &= (1-\alpha)[\underbrace{R_{n-1} - \mu_{n-1}\mu_{n-1}^\top}_{\Sigma_{n-1}} + \\ &\ldots\quad \alpha\underbrace{(\mathbf{x}_n\mathbf{x}_n^\top - \mu_{n-1}\mu_{n-1}^\top - \mathbf{x}_n\mu_{n-1}^\top + \mathbf{x}_n^\top\mu_{n-1})}_{(\mathbf{x}_n-\mu_{n-1})(\mathbf{x}_n-\mu_{n-1})^\top}]\end{aligned}$$

Thus,

$$\Sigma_n = (1-\alpha)[\Sigma_{n-1} + \alpha(\mathbf{x}_n - \mu_{n-1})(\mathbf{x}_n - \mu_{n-1})^\top] \tag{8}$$

Small values of the parameter α assign small weights to present values, thus the sequence of estimates is smoother with smaller α. In experiments, we found $\alpha = 0.001$ to be a reasonable setting.

For the implementation of the recursive estimation algorithms only one vector and one matrix accumulator are necessary. This is a clear advantage over windowed estimation approaches, where all measurements that fall inside the window must be memorized.

4.3 Distance calculation

Clip-to-clip distance is calculated upon loading of a new clip by storing the Kullback-Leibler (KL) divergences between each clip pair to a distance matrix. Clip-to-stream distances must be recalculated continuously because the distribution of stream features can change at any time.

4.4 Evaluation of the incremental similarity measure

In figure 2, we have plotted the similarities of a set of music clips to a music stream. We used the 4 reference clips listed in table 1.

The 4 plots correspond to the 4 clips, each plot shows the similarity of the current stream content to the reference clip (x axis is in audio frame numbers, 1 frame corresponds to the FFT hop size of 1024 samples). We calculate the similarity of the stream at time t to the clip C_i as the reciprocal value of the KL divergence between the stream model and the clip model, and we normalize the sum of clip-to-stream similarities over all clips at time t to one.

Artist/Composer	Title
Absolute Beginner	Rock On
Metallica	The Four Horsemen
Al Green	Listen
Heitor Villa-Lobos	Etude n. 1 en mi mineur

Table 1. Reference clips used for the evaluation

Artist/Composer	Title	Frames
A Tribe Called Quest	Buggin Out	0–9388
Megadeth	This was my Life	9389–18776
Al Green	Listen	18777–25150
Heitor Villa-Lobos	Etude n.2 en la majeur	25151–29715
Male	Speech	29716–34539

Table 2. Stream used for the evaluation

The similarity measure successfully identifies music from the same genre, style, or artist ("A Tribe Called Quest" have the highest similarity with the "Absolute Beginners" clip, "Megadeth" are most similar to "Metallica", and the clip-to-stream self-similarity for Al Green's "Listen" stays at ~ 0.8 almost over the entire duration of the song). However, the system tends to confuse Rap vocals with pure speech although it should emit low similarity scores with respect to all reference clips (last part of the stream versus the "Absolute Beginner" song) - this observation seems to be related to the fact that simple timbral similarity based on MFCC statistics wrongly classifies "Rap" and "Speech" as similar. Another shortcoming can be observed using the Absolute Beginner's and Al Green's songs as examples - both yield high similarity scores in the beginning of the Al Green tune, which seems to be related to the use of strong percussion in both pieces.

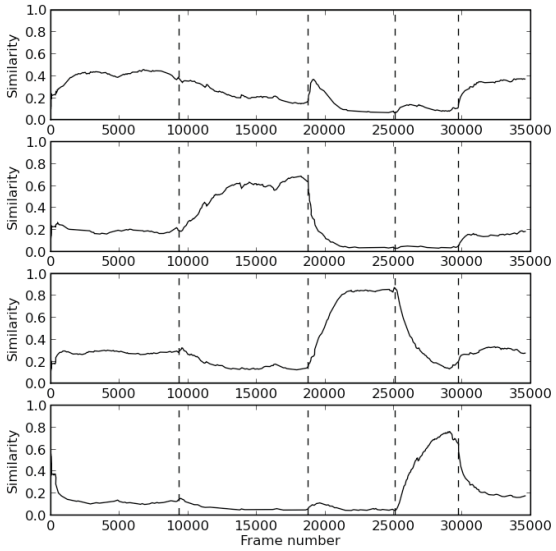

Figure 2. Online stream similarity to offline reference clips

5 GRAPHICAL USER INTERFACE

Figure 4 shows a screenshot of the running *StreamCatcher* application. The GUI of the application currently consists of a visualization that allows direct manipulation (zooming, panning, rotating, playback of clips and streams) of the visualized objects via a pointing device like a mouse or a touch screen.

By pressing the "Load Clip" button, a new sound clip is loaded into the application. After analyzing and k-NN-classifying the clip, a label (consisting of a textual description and a color) is suggested by the software in a dialog window. Now the user has the possibility to accept the suggestion, to change the label to another – already existing – label or create a new label. Furthermore, it is always possible to alter the labeling of the clips later on. If the "Open Stream" button is pressed, the user can load an MP3 stream, which is subsequently visualized with the algorithm described below.

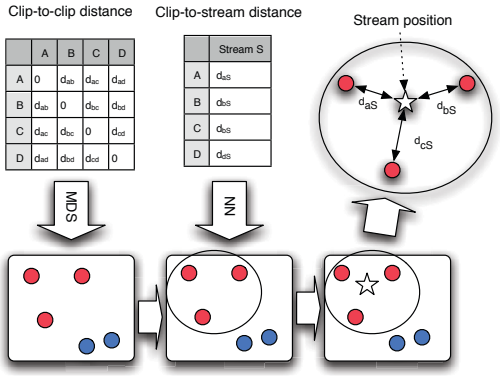

Figure 3. Placement of online streams in visualization space

5.1 Visualization algorithm

It is well known that the symmetric KL divergence between Gaussians does not satisfy the requirements to a proper distance measure [12]. Therefore, we use a visualization technique that seeks to preserve the KL divergence values in a low-dimensional visualization space and relaxes this demand if that is not possible. A classic distance-preserving data mining/visualization technique is Multidimensional Scaling [4], which aims at placing data points in low dimensional visualization space while approximating the distances in feature space as closely as possible. Multidimensional scaling can be implemented as a spring model [2, 11], a physically inspired model of spring-connected nodes aiming at finding a node placement that minimizes the cumulative deflection from the springs' resting states. By mapping distances in feature space to spring lengths, the spring model solves the MDS problem approximately.

We chose to use a spring model variant of MDS because of its ease of implementation and because of the attractive visualizations that can be generated by constantly drawing the gradual relaxation of the system. By using a gradient-descent based solution algorithm (see algorithm 1), a placement of the nodes that minimizes the overall stress (the deflection from the springs' resting state) is constructed.

While the overall stress in the model (S_{cum}) is larger than a threshold (S_{thresh}), the algorithm loops over all nodes and determines for each node a force acting upon the node by calculating a weighted sum of unit vectors pointing from the current node to all other nodes. The weights can be positive or negative, depending on the difference between high- and low-dimensional distances. Then, a velocity vector is

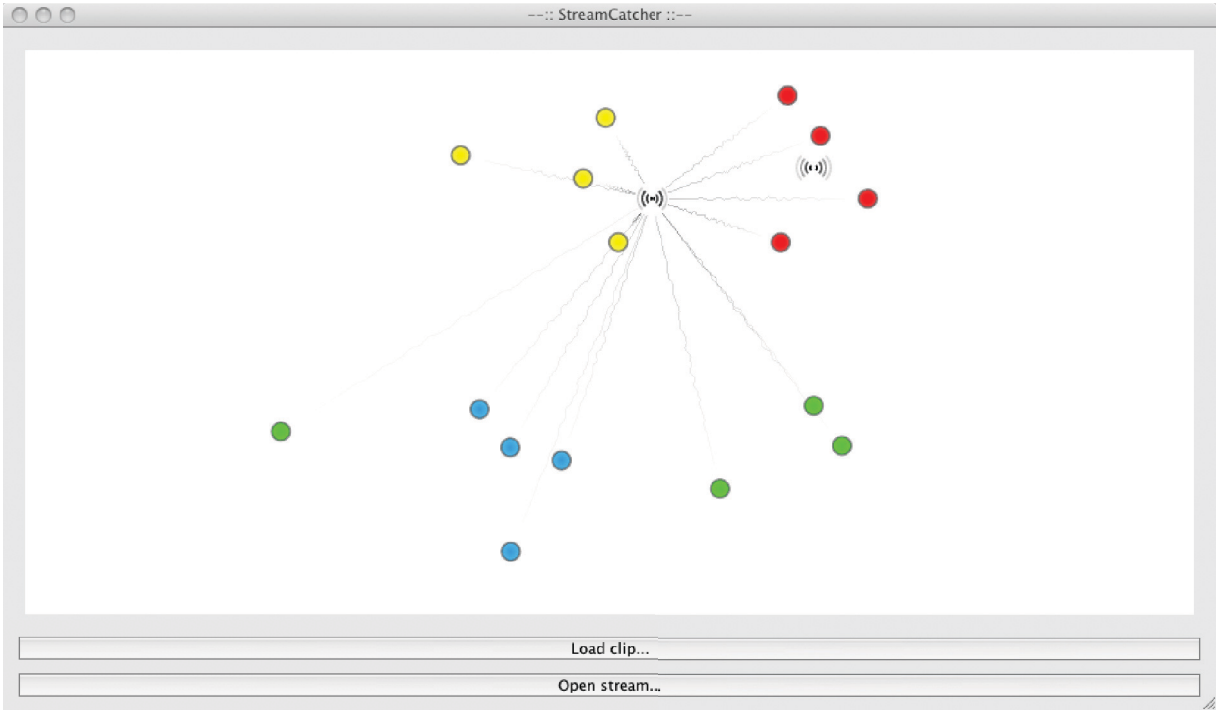

Figure 4. Screenshot showing the running application

Algorithm 1 Spring model layout algorithm

repeat
  $S_{cum} = 0$
  for $C_a \in \mathcal{C}$ **do**
    $\overrightarrow{f} = \overrightarrow{0}$
    for $C_b \in \mathbf{C}$ **do**
      $\overrightarrow{u} \leftarrow unit(p(C_a) - p(C_b))$
      $S \leftarrow d_{high}(C_a, C_b) - d_{low}(C_a, C_b)$
      $\overrightarrow{f} \leftarrow \overrightarrow{f} + \overrightarrow{u} * S$
      $S_{cum} \leftarrow S_{cum} + |S|$
    end for
    $v(C_A) \leftarrow v(C_A) + \overrightarrow{f}$
    $p(C_a) \leftarrow p(C_a) + v(C_A)$
    $v(C_A) \leftarrow v(C_A) * dampingFactor$
  end for
until $S_{cum} < S_{thresh}$

S_{cum} : Cumulative stress in the model
$\mathcal{C}$: Set of clip nodes
$unit$: Function returning a unit vector
p : Vector-valued function returning the 2D position of a clip node
d_{high} : Distance of clips in feature space
d_{low} : Distance of clips nodes in visualization space
v : Vector-valued function returning the current velocity of a clip node
$\overrightarrow{f}$: Force acting upon a node

updated for the current node by adding the force vector to its current value. Finally, the node's position is updated by moving it into the direction of its velocity, and the node's velocity is multiplied with a damping factor.

We use spring model MDS to place the static clips in a 2D visualization. Upon loading of a new clip, an MDS procedure is executed until the cumulative stress goes below a threshold value. The placement of the dynamic streams is determined by calculating the distances to the 3 nearest neighbors in the feature space and using the reciprocal value of those distances to calculate a convex combination of the positions of the nearest neighbors that have been placed by the MDS algorithm (see figure 3). Additionally, the true distances to the anchor clips are visualized by drawing the audio waveform of the stream's audio signal between the stream's position and the clips' positions, and by modulating the opacity of this waveform with the feature space distance. The smaller the feature space distance, the more opaque is the drawing of the waveform.

In the screenshot (figure 4), example clips for four different styles/genres of music are loaded (Heavy Metal: Red, HipHop: Yellow, Jazz Guitar: Green, Classical Guitar: Blue). The green outlier, which is closer to the Classic Guitar cluster than to Jazz Guitar, is a solo guitar piece by Pat Metheny, which makes the placement quite reasonable. The two streams (the loudspeaker-like symbols emanating the waveforms) play Heavy Rock music and Soul/Funk. The Rock stream is placed near the Heavy Metal-cluster, whereas the Soul stream

is placed close to the HipHop clusters, which characterizes the music quite well.

6 CONCLUSIONS & FUTURE WORK

We have presented an application prototype for similarity analysis and visualization of audio files and online audio streams in a common framework. Music clips from personal music collections are used as prototypical instances of musical concepts that define a visualization space in which online streams are embedded. For the online calculation of music similarity from streams we derived a similarity measure which is based on incrementally updated statistical models. The system comprises a user interface that supports the user in identifying prototypical examples of high level musical concepts. Acoustic similarity is mapped to proximity data in a 2D visualization, which in turn is derived from a high dimensional timbre similarity space by means of multidimensional scaling. Sound prototypes can be labeled with user-definable colors as well as textual labels.

The system successfully identifies streams that sound similar to prototypical clips. Thus, it can be used to get a rough overview of audio stream content, e.g., of internet radio stations. By exploring the suggestions of the software, the user may find music he or she likes faster than by just blindly trying radio stations.

We are currently working on improving the underlying distance measure and on special cases like speech-music discrimination, which could be a very useful feature for this application.

7 ACKNOWLEDGEMENTS

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EVALUATING AND VISUALIZING EFFECTIVENESS OF STYLE EMULATION IN MUSICAL ACCOMPANIMENT

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ABSTRACT

We propose general quantitative methods for evaluating and visualizing the results of machine-generated style-specific accompaniment. The evaluation of automated accompaniment systems, and the degree to which they emulate a style, has been based primarily on subjective opinion. To quantify style similarity between machine-generated and original accompaniments, we propose two types of measures: one based on transformations in the neo-Riemannian chord space, and another based on the distribution of melody-chord intervals. The first set of experiments demonstrate the methods on an automatic style-specific accompaniment (ASSA) system. They test the effect of training data choice on style emulation effectiveness, and challenge the assumption that more data is better. The second set of experiments compare the output of the ASSA system with those of a rule-based system, and random chord generator. While the examples focus primarily on machine emulation of Pop/Rock accompaniment, the methods generalize to music of other genres.

1 INTRODUCTION

Automatic generation of music in a specific style has been a focal topic in computational music research since the early days of computing. Many researchers have designed systems that emulate music styles in compositions. The evaluation of the musical output, the degree to which it achieves its goal of emulating a particular style remains a challenge. Evaluation is often in the form of subjective opinion. It is our goal to fill this gap in quantitative evaluation of style emulation in automatic accompaniment systems.

In [1], Meyer states that "style is a replication of patterning, ..., that results from a series of choices made within some set of constraints." Accompaniment can be considered the outcome of a series of choices over possible chords under certain contextual constraints, such as melody, phrase, and key. For example, in four-part harmonization, the composition must follow the counterpoint and voice-leading rules in music theory. In contrast, modeling of the accompaniment style in Pop/Rock tends to be vague and difficult because, according to Stephenson [2], "in rock, ..., melody is allowed to interact with harmony more freely," and the process becomes more complex because, as Everett states [3], "many different tonal systems are now practiced by the same artist, on the same album."

In this paper we propose methods to visualize, measure, and quantify the quality of an accompaniment generated with the goal of emulating an original accompaniment, the "ground truth". These measures are designed to evaluate automatic style-specific accompaniment (ASSA) systems. We examine the style as captured by the musical relations between melody and harmony, and between adjacent chords. We develop six metrics based on these two types of measures.

Using these quantitative methods, we design experiments to explore training set selection strategies that further the ASSA system's style emulation goals. For machine learning tasks, it is often the case that more training data guarantee better results. For ASSA goals, more training songs may not necessarily improve the output quality if the training set is not consistent with the desired style. The first set of experiment explores the factors impacting style emulation success.

A second set of experiments compare the degree of style-specificity between the best ASSA systems, a rule-based harmonization system, and a random chord generator. We conduct the experiments on five well known Pop/Rock albums by Green Day, Keane, and Radiohead, providing detailed statistics and case studies for the resulting accompaniments. The findings we report generalize to genres outside of Rock music, and to some extent to simulation of style in music in general.

The paper is organized as follows: Section 2 describes related automatic accompaniment systems, with and without style requirements, and their evaluations. It concludes with a brief description of an ASSA system, which forms the basis of much of our evaluations. We present the evaluation methods in Section 3. Intra-system comparisons are shown in Sections 4, and inter-system results in 5, followed by the conclusions.

2 RELATED WORK

Baroque style four-part harmonization has been a popular accompaniment generation application since the earliest days of computing. Recent examples include [4, 5]. Many rules govern these Chorale-type harmonizations, which can be applied in their synthesis and evaluation. Sixteenth century compositions in the style of Palestrina have also be emulated using Markov models [6]. Such compositions are similarly and strictly circumscribed by a host of rules, which can be used in the evaluation of their correctness [7]. Temperley and Sleator proposed a set of preference rules to harmonizing melodies in the Western classical style [8]; these rules are implemented in their Harmonic Analyzer [9].

In the popular realm, the i-Ring system [10] generates an accompaniment for any eight-measure melody. The accompaniment is based on state transition probabilities calculated from a training set of 150 songs. For evaluation, 10 participants were asked to rate the accompaniment as Good, Medium, or Bad.

More recently, MySong [11] uses Hidden Markov Models, training on 298 popular songs in genres including Pop, Rock, R&B, Jazz, and Country Music. Users can choose between two style-related modes: 'Jazz,' which uses less common triads, or 'Happy,' which selects more major-mode chord transitions. 26 sample accompaniments by MySong were evaluated subjectively by 30 volunteer musicians.

While the output of the above systems fit the melody, because the systems learn from general training sets, the style captured by the output lacks specificity. Thus, these systems do not address the emulation of specific accompaniment styles, as embodied in a distinctive song or a particular band's output. Evaluations take the form of subjective opinion, and lack an objective or quantitative component.

In [12], Chuan & Chew proposed an ASSA system that can generate style-specific accompaniment to a melody given only a few examples. The system consists of a chord tone determination and a chord progression module. The chord tone determination module applies machine learning to determine which notes in a melody are likely chord tones based on the specified style. The system then constructs possible chord progressions using neo-Riemannian transforms, representing them in a tree structure, and selects the highest probability path based on the learned probabilities in a Markov chain. This system was rated informally by subjectively judgement via a Turing test.

3 QUANTIFYING ACCOMPANIMENT DISTANCE

This section first presents ways to quantify and visualize distance between two chords, and metrics to evaluate distance between two different accompaniments to the same melody. The examples are generated by the ASSA system of [12].

3.1 Visualization

Two visualization models are described as follows. The first considers musical distance between two chords based on neo-Riemannian transforms, the second addresses the difference between chord choices in relation to the melody.

3.1.1 Neo-Riemannian Distance in Chord Space

Music theorists have used neo-Riemannian transformations to analyze harmonic diversity in Pop/Rock music [13]. In the neo-Riemannian chord space, chords are connected by three types of neo-Riemannian operations (NROs), P (parallel), L (leading-tone exchange), and R (relative).

Figure 1 shows the three types of NROs in chord space, where chords are represented in Roman numerals. Vertical neighbors are connected by P/R operations, and horizontal neighbors have Dominant/Subdominant (D/S) relationships.

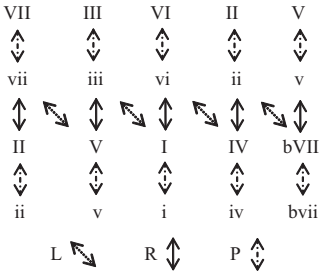

Figure 1. Neo-Riemannian transforms in chord space.

To quantify the distance between two chords – for example, to determine how well a generated chord compares to the original – we can compute the shortest distance between the two chords in the neo-Riemannian space. Two chords connected by an NRO share two tones, with the third being one or two half steps apart. The fewer NROs between the chords, the closer they are perceptually.

Chord Space Distance-Time Plot: To visualize the distance between a machine-generated and an original accompaniment, we can graph this distance in chord space over time. Figure 2 shows one such graph for comparing the generated and original accompaniments.

The x-axis marks the chord count; the yz-axes represent chord space. The plot charts a given accompaniment's distance from the baseline (original), according to chord space distance. If the two accompaniments are identical, the graph would be a horizontal line defined by $y = z = 0$. The advantage of this visualization is that we can see the quality of each generated chord by observing its neo-Riemannian distance from the original, and we can observe their chord relation by examining the direction of the deviation.

Chord Map Distribution: The second visualization employs the key map idea proposed in [14]. A chord map (Fig-

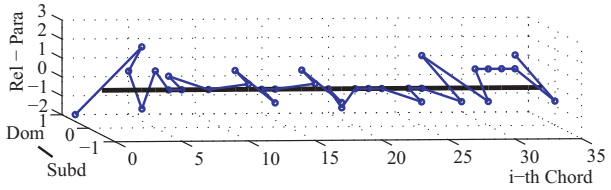

Figure 2. Chord space distance-time plot for Keane's *Somewhere Only We Know* (K1) trained on *Bedshaped* (K11) in the album *Hopes and Fears*.

ure 3) presents a nine-cell grid in which each cell presents a closely related chord with respect to the center chord. Apart from D/S and P/R, the nine cells include the I (Identity) and the combination operations: DR, SR, DP, and SP.

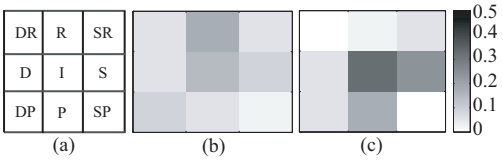

Figure 3. Chord map distributions: (a) the nine close chords; (b) result for K1 trained on K5 (*She Has No Time*); and, (c) result for K1 trained on K11

We count the number of chords in these nine categories, with respect to the original, in a generated accompaniment, and divide the number by the total number of chords. Two resulting grayscale plots are shown in Figures 3 (b) and (c). The plots show that the accompaniment for K1 trained on K11 is markedly better than that trained on K5.

3.1.2 Melody-Chord Interval Distribution

Another way to compare two accompaniments is by examining the relation between the chords and the melody they harmonize, to capture the degree of consonance-dissonance.

Suppose a subsequence of melodic notes, with pitch-duration values $\{(a_i, d_i), i = 1, \ldots, m\}$, is accompanied by a chord $\mathbf{x}$ containing the pitches $\{x_1, \ldots, x_n\}$. Assume that the a_i's and the x_j's have been normalized by the key of the melody so that 0 represents the tonic. The weight of a particular interval, k, between the melody pitches, $\mathbf{a}$, and the chord, $\mathbf{x}$, is defined as follows:

$$W_{\mathbf{a},\mathbf{x}}(k) = \sum_{\{i,j:(a_i - x_j) mod 12 = k\}} d_i, \qquad (1)$$

where $k = 0, \ldots, 11$. For example, the interval weight vector for a melodic note C, with duration d, and a C major triad is $[d, 0, 0, 0, 0, d, 0, 0, d, 0, 0, 0]$. We then normalize the interval weight vector to obtain the interval distribution:

$$Dist_{\mathbf{a},\mathbf{x}}(k) = \frac{W_{\mathbf{a},\mathbf{x}}(k)}{\sum_{i=0}^{11} W_{\mathbf{a},\mathbf{x}}(i)}. \qquad (2)$$

To compare two accompaniments, we take the difference between their melody-chord distributions. Suppose we have a sequence of melodic fragments $\{\mathbf{a}_1, \ldots, \mathbf{a}_T\}$, their original accompanying chords, $\{\mathbf{x}_1, \ldots, \mathbf{x}_T\}$, and chords in the generated accompaniment, $\{\mathbf{y}_1, \ldots, \mathbf{y}_T\}$. We calculate the difference between the distributions as follows:

$$\Delta_i = Dist_{a_i,y_i} - Dist_{a_i,x_i}, \text{ where } i = 1, \ldots, T. \qquad (3)$$

Melody-Chord Interval Distribution Difference: Figure 4 shows a visualization of the difference between two melody-chord interval distributions: the original accompaniment to the melody of K1, and one by the system trained on K11. The three-dimensional map of interval (half-step) distribution over time is shown as a two-dimensional grayscale plot. The horizontal axis shows the chord count, while the vertical axis the number of half steps.

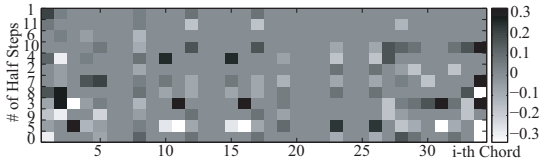

Figure 4. Ordered melody-chord interval distribution difference between the original K1 accompaniment and one trained on K11.

A predominance of darker gray indicates that the generated accompaniment contains extra types of half steps not in the original, while lighter colors mean that the generated chords lack certain types of half steps.

For visual coherence, we order the y-axis by the distribution of the original accompaniment, $Dist_{\mathbf{a},\mathbf{x}}$, in descending order, up the vertical axis. In this way, we can expect a generated accompaniment to be more stylistically distant from the original (having more frequent rare intervals) when more darker cells appear near the top of the graph, and less stylistically consistent (having fewer of the popular intervals) if more lighter cells are present near the bottom.

3.2 Metrics

We propose six metrics, three percentages and three distance metrics, to quantitatively assess a generated accompaniment, as shown in Table 1.

The first metric, *correct rate*, calculates the percentage of melodic notes that are correctly classified as chord tones and non-chord tones in the generated accompaniment, where chord and non-chord tones are given by examining the original accompaniment. This measure pertains to the ASSA system in [12], and to other systems that classify melody notes into chord and non-chord tones.

Table 1. Metric values' names, meanings and ranges

name	meaning	max	min
correct rate	% correct chord/non-chord tone classification	100	0
same chords	% of generated chords identical with original	100	0
chords-in-grid	% of generated chords in chord map	100	0
NR distance	ave NR distance bet generated/original chords	7	0
HS distance	MSE between interval (half-step) dist	2	0
wHS dist	weighted MSE bet interval distributions	2	0

The second metric, *same chords*, records the percentage of chords generated that are identical to the original. Note that this is also the value in the center cell of the chord maps shown in Figure 3. The third metric, *chords in grid*, gives the percentage of chords generated that are closely related to the original on the chord map; this value can be computed by summing all values in the cells of the chord map.

The *NR distance* metric shows the average shortest neo-Riemannian distance between the generated chord and the original, i.e., the average minimum neo-Riemannian distance between each data-point and the center line in the chord space distance-time plot shown in Figure 2. Note that the maximum NR distance between any two chords is 7.

The last two metrics are generated from the melody-chord interval distribution described in Section 3.1.2. The *HS distance* is the mean squared difference between the interval (half-step) distributions of the generated and the original accompaniments:

$$HS = \frac{1}{T}\sum_{i=1}^{T}\sum_{k=1}^{11}\Delta_i(k)^2. \quad (4)$$

The worst case occurs when only one type of half step dominates the generated and original accompaniments' distributions, but the types are different, causing a distance of 2.

The final metric *wHS distance* is like the HS distance, except that Δ_i is multiplied by the inverse of the half-step distribution value for the original accompaniment. The weighted HS distance increases the weight for infrequent half steps, reflecting the fact that rare, dissonant intervals can significantly impact the perceived accompaniment style.

Let $Dist_{\mathbf{a},\mathbf{x}}$ be the half-step distribution of the original accompaniment for melody fragment $\mathbf{a}$. The weighting factor for each half step, $k = 0, \ldots, 11$, can be calculated as:

$$w(k) = \begin{cases} 1/Dist_{\mathbf{a},\mathbf{x}}(k), & \text{if } Dist_{\mathbf{a},\mathbf{x}}(k) > 0 \\ 0.001, & \text{if } Dist_{\mathbf{a},\mathbf{x}}(k) = 0 \end{cases} \quad (5)$$

The weighted HS distance is then calculated as:

$$wHS = \frac{1}{T}\sum_{i=1}^{T}\sum_{k=1}^{11} w(k) \times \Delta_i(k)^2, \quad (6)$$

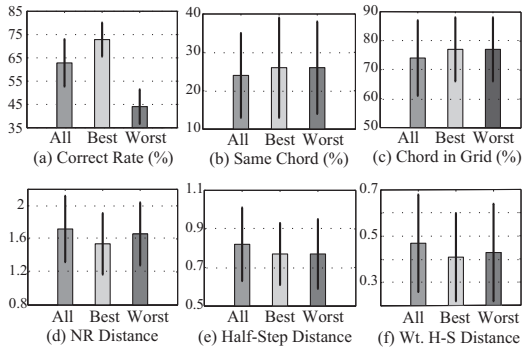

Figure 5. Summary statistics for generated accompaniments over five albums

4 INTRA-SYSTEM COMPARISONS

This section describes experiments involving only the ASSA system of [12]. Recall that this system has a chord tone determination (CTD) module, and a chord progression module. We examine the general quality of the results generated by the system, focussing on the impact of training data on the quality of the machine-generated accompaniment, so as to improve the choice of training songs.

The data set consists of songs from the five albums: Green Day's *Dookie* and *American Idiots*, Keane's *Hopes and Fears*, and Radiohead's *Pablo Honey* and *Hail to the Thief*. We use commercial lead sheets for the ground truth.

4.1 One-Song v.s. All-Except-One Training Policy

To test whether more data is indeed better in ASSA, these experiments compare two training song choices. The first selects one song from an album for training, and tests the model on another song in the same album. The second uses all except one song for training, and tests the model on the held out song.

Figure 5 shows the statistics. "Best" and "Worst" report the performance by the model that is trained on a single-song in the same album, and that has the highest and lowest correct CTD rates, respectively. "All" refers to results where all other songs in the same album form the training set.

Note that except for the correct CTD rate, Figure 5(a), the statistics for Worst-CTD consistently outperform those for All. The results indicate that more data is not always better, and that the use of neo-Riemannian transforms offer a robust way to generate chord progressions, even when chord tone information is poor.

4.2 Correct Chord Tones vs. Best Chord Overlap

To test the ASSA system's sensitivity to different parameters, we compare the Best-CTD performance in the previous section with two other training data selection policies.

The first uses the three songs in the same album with the best CTD correct rates; the second uses the three songs that share the most common chords (CC) with, and having the least extra chords over, the test song.

Figure 6 shows the summary statistics for these tests. We observe that the CC training set achieves the highest *same chord* and *chords in grid* percentages, and reports the lowest half-step distance metrics.

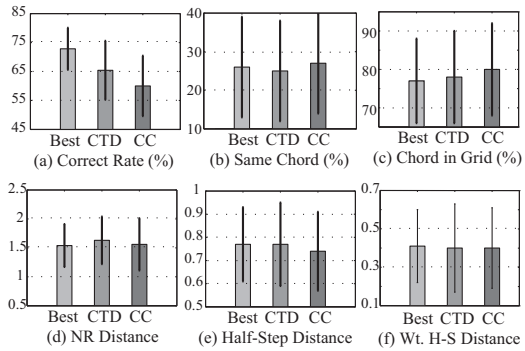

Figure 6. Summary statistics for accompaniments generated using Best, CTD and Common Chord training data sets

4.3 Case Study: Visualization of Details of Two Results

In this section we examine two accompaniments by visualizing their chord space distance-time plots, and their melody-chord interval distribution difference graphs. We compare the output of systems trained on the CC set (the best training set), and on all other songs in the same album (the worst training set.) Figure 7 shows the first 12 bars of the melody and the original accompaniment on Keane's *Your Eyes Open*, and the chords generated by ASSA with the CC and All training sets.

Figure 8 shows the neo-Riemannian distance plots over time. Observe that the accompaniment generated by All contains larger deviations from the original than that by the CC training set. Figure 9 shows the melody-chord interval distribution visualizations of the two generated accompaniments. Note that Figure 9(b) has more dark cells near the top than Figure 9(a). These darker cells indicate that the accompaniment generated by the All training set contains more melody-chord intervals that are rare in the original.

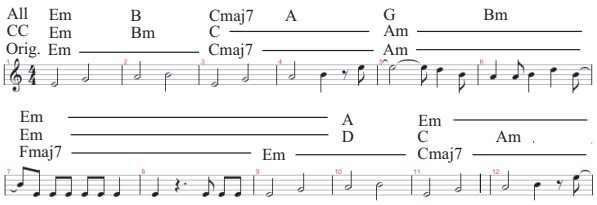

Figure 7. The first 12 bars of Keane's *Open Your Eyes*

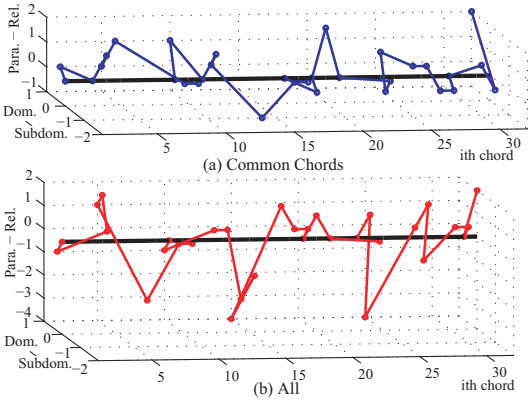

Figure 8. Chord space distance-time plot for Keane's *Open Your Eyes*

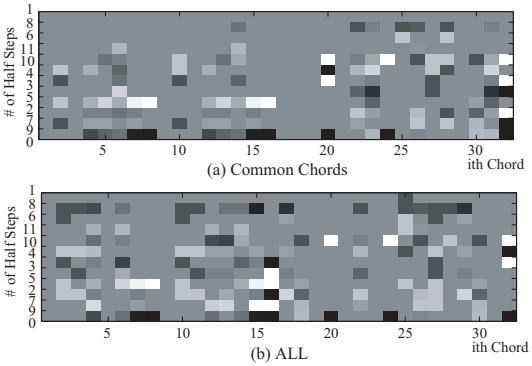

Figure 9. Interval distribution distance graphs for Keane's *Open Your Eyes*

5 INTER-SYSTEM COMPARISONS

In this section we compare the ASSA system [12] with two rule-based harmonization systems, Temperley and Sleator's (T-S) Harmonic Analyzer [8], and a random harmonizer with only one constraint.

The Harmonic Analyzer [9] generates the root of the chords for harmonization without indicating their modes (major or minor). For comparison, we interpret the chords as being the common ones as described in [15]. For example, when G is reported by the Harmonic Analyzer in a song in C major, we assign a G major instead of a G minor chord.

We further design a simple random chord selector as the base case for the comparisons. In order to construct a reasonable accompaniment system, we add one constraint to the random chord generator: an accompaniment chord must contain at least one of the melody notes it harmonizes, and are randomly assigned if the bar has no melody note.

Figure 10 shows summary statistics comparing the ASSA with the Best-CTD and the CC training sets, the T-S Harmonic Analyzer, and the random chord selector (Rand). The figure shows that (a) the ASSA and T-S systems report sim-

ilar neo-Riemannian distances, (b) and (c) T-S generates shorter half-step distances, and (e) achieves a higher percentage of chords in grid; however, (d) ASSA with both training sets outperforms T-S on same chord percentage.

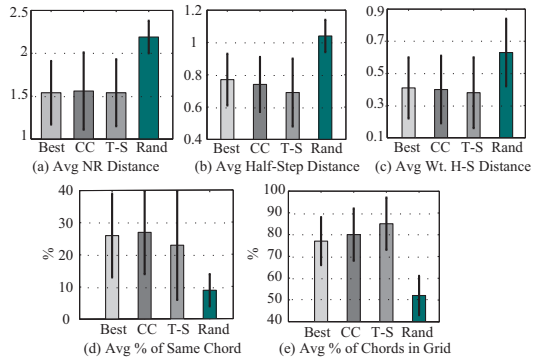

Figure 10. Overall statistics over systems.

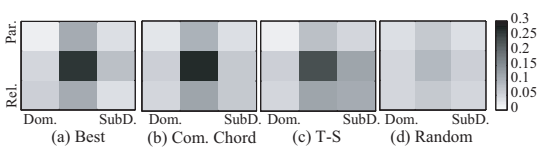

Figure 11. Chord map distributions of systems.

Figure 11 shows the chord map distributions for the four systems. (a) and (b) show that more chords generated by the ASSA system match the original exactly. They also show that ASSA tends to generate chords that are parallel or relative to the original. In (c), we observe that the T-S system generates almost equal numbers of chords in R, S, and SR. In (d), the chords generated by the random chord selector are relatively evenly distributed among the close chords.

6 CONCLUSIONS

We have proposed quantitative methods for evaluating and visualizing machine-generated style-specific accompaniments. Using these methods, we showed that a training set with more chords in common with original leads to better style emulation. In the inter-system comparisons, we showed that the ASSA system produces more chords identical to those in the original song than the T-S system.

7 ACKNOWLEDGEMENTS

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MUSIC STRUCTURE ANALYSIS USING A PROBABILISTIC FITNESS MEASURE AND AN INTEGRATED MUSICOLOGICAL MODEL

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ABSTRACT

This paper presents a system for recovering the sectional form of a musical piece: segmentation and labelling of musical parts such as chorus or verse. The system uses three types of acoustic features: mel-frequency cepstral coefficients, chroma, and rhythmogram. An analysed piece is first subdivided into a large amount of potential segments. The distance between each two segments is then calculated and the value is transformed to a probability that the two segments are occurrences of a same musical part. Different features are combined in the probability space and are used to define a fitness measure for a candidate structure description. Musicological knowledge of the temporal dependencies between the parts is integrated into the fitness measure. A novel search algorithm is presented for finding the description that maximises the fitness measure. The system is evaluated with a data set of 557 manually annotated popular music pieces. The results suggest that integrating the musicological model to the fitness measure leads to a more reliable labelling of the parts than performing the labelling as a post-processing step.

1 INTRODUCTION

Western popular music pieces tend to follow a sectional form where the piece can be thought to be constructed of smaller parts (e.g., verse or chorus) which may be repeated during the piece, often with slight variations. The analysis of music structure is the process of recovering a description of this kind from audio input.

Automatic music structure analysis enables several applications, including a structure-aware music player [4] and music summarisation or thumbnailing [3, 12, 9] (see [8] for a discussion of further applications). A popular aim has been to locate a representative clip of the piece, such as the chorus, but there are also systems which aim to describe the structure of the whole piece, for example [2, 7].

In the proposed method (block diagram illustrated in Figure 1), the audio content is described using three sets of features: mel-frequency cepstral coefficients (MFCCs), chroma, and rhythmogram, to address different perceptual dimensions of music. [1] The use of multiple features is motivated by [1], which suggests that changes in timbre and rhythm are important cues for detecting structural boundaries in addition to repetitions. A large set of candidate segmentations of the piece is first constructed. All non-overlapping segment pairs are then compared based on the features and two different distance measures: one based on the average value of a feature over a segment and the other matching the two temporal sequences of features in the segments. Utilising the distance values we calculate the probability of the two segments to be occurrences of same musical part (i.e., repeats). The values are used as terms in a fitness measure which is used to rank different candidates for the piece structure description.

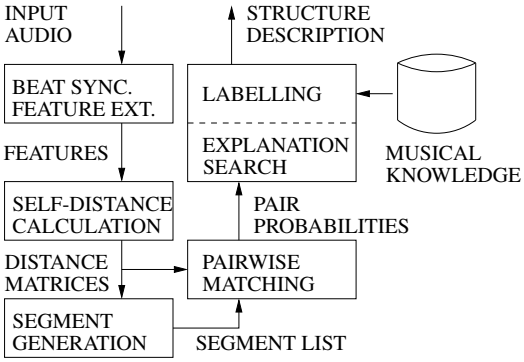

Figure 1. Analysis system block diagram.

The found structural description consists of subdivision of the piece into segments and of forming *groups* of segments that are occurrences of the same musical part. To make the description more informative, the segment groups are automatically named using musical part labels, such as "verse" or "chorus". The labelling is done by utilising a musicological model, in practice an N-gram model for musical parts estimated from a manually annotated data set. Two different strategies for the labelling are evaluated: either per-

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[1] A similar set of three features have been used earlier in [5], but the features were considered individually instead using a combination of them.

forming it as a post-processing step after the segments and their grouping has been decided, or by integrating it into the fitness measure for the descriptions. The latter method allows assigning the labels already while searching for the description, and enable utilising the potentially useful musical information of the part N-grams in the search.

The main original contributions of this paper concern the last two blocks in Figure 1, the pairwise matching of segments, the search algorithm, and musicological model for naming the found parts.

The performance of the proposed system is evaluated using a data set of 557 popular music pieces with manually annotated structure information. Different configurations of the overall system are studied in order to determine potential points for improvement.

2 PROPOSED METHOD

Different parts of the proposed method are described now in more detail.

2.1 Feature Extraction

The system uses three sets of acoustic features, all of them with two different time scales to provide the necessary information for further analysis. The extraction starts by estimating the locations of beat (tactus) pulses using the method from [6]. The pulse periods are halved by inserting an extra pulse between the estimated location to alleviate problems due to possible π-phase errors. The pulses are used to create a beat-synchronised time grid for making the features less sensitive to tempo variations.

MFCCs are used to describe the timbral content of the signal. They are calculated using 42-band filter bank energies which are decorrelated with discrete cosine transform. The lowest coefficient is discarded, and the following 12 coefficients are used as the feature. The MFCCs are calculated with 92.9 ms frames with 50% overlap.

The harmonic content is described with chroma, which is calculated using the method described in [14]. First, the saliences of fundamental frequencies in the range 80–640 Hz are estimated. The linear frequency scale is transformed to a musical one by retaining only the largest salience value in each semitone range. The chroma vector of length 12 is obtained by summing the octave equivalence classes. The frame division is the same as with MFCCs.

The rhythmic content is described with *rhythmogram* proposed in [5], but replacing the original perceptual spectral flux front-end with an onset accent signal obtained as a by-product of the beat estimation. After removing the mean value from the accent signal, it is divided into several seconds long, overlapping frames with spacing corresponding to the 46.4 ms frame hop on MFCCs and chroma calculation. From each frame, autocorrelation is calculated and the values below the lag of 2 seconds are retained. The values are normalised to produce value 1 on lag 0.

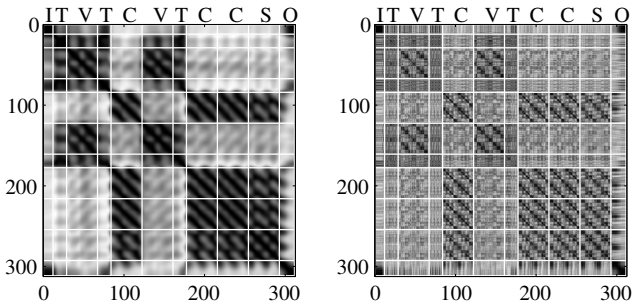

Figure 2. Examples of SDMs calculated using chroma features with two different time-scale parameters. The SDM on the left was calculated with a low cut-off frequency making different parts resemble blocks, and the SDM on right was calculated with a high cut-off making -45° stripes visible. All axes show time in seconds and a darker pixel value denotes lower distance. The overlaid grid illustrates the annotated part borders. The part labels are indicated as: intro (I), theme (T), verse (V), chorus (C), solo (S), outro (O).

Beat-synchronised feature vectors are obtained by calculating the average value of a feature between each two beat pulses. From each feature, two versions focusing on different time scales are produced. With MFCCs and chroma, this is done by low-pass filtering the feature vector series over time with second order Butterworth IIR filters. The filter cut-off frequency ω is determined by $\omega = 1/\tau$, where τ is the time scale parameter. The used values of τ for MFCCs are 8 for finer time scale and 64 for coarse time scale, for chroma the values are 0.5 and 64. The filtering is done twice along time, first forward and then backwards to double the magnitude response of the filters and to cancel out phase distortions. The rhythmogram feature is not filtered, but instead the length of the autocorrelation window is adjusted to a length which corresponds to 4 beat frames for the shorter time scale and 32 beat frames for the longer one. Each feature is normalised to zero mean and unity variance over the whole piece. The feature vector in beat frame k is denoted by $\mathbf{f_k}$. As the remaining operations are similar for all features and time scales, the same notation is used for all feature vectors. The above-mentioned time scale parameter values were selected based on the results from [10].

2.2 Self-distance Matrices

Using the extracted feature vector series, self-distance matrices (SDMs) are calculated. Each element in the SDM denotes the distance between feature vectors in frames k and l, and is calculated with the cosine distance measure $D_{k,l} = d(\mathbf{f_k}, \mathbf{f_l})$. Figure 2 illustrates the SDMs calculated for an example pieces using the chroma features.

2.3 Formal Model of Structure Description

The method searches for a *description* E for the structure of a piece S, which is represented as a sequence of *beat frames* $S = c_1, c_2, \ldots, c_K$, K being the length of the piece in beats. The sequence S can be divided into M subsequences of one or more consecutive frames $S = s_1, s_2, \ldots, s_M$, where each subsequence is a *segment* $s_m = c_k, c_{k+1}, \ldots, c_{k+L-1}$. A segment represents an individual occurrence of a musical part. For each segment s_m, there is also associated information about the *group* g_m where it belongs to. The combination of segment and group information is denoted by $(s, g)_m$. All possible segmentations and groupings of the piece form a set $\mathbb{S} = \{(s, g)_1, (s, g)_2, \ldots, (s, g)_Z\}$, where Z is the total number of all segments and groupings. The set of segments with the same group assignment represents all occurrences of a musical part (for example a chorus). A structure description $E \subset \mathbb{S}$ consists of the division of the entire piece into non-overlapping segments and of the grouping of the segments.

2.4 Border Candidates

Because the number of possible segmentations is very high, a set of potential segment border candidates is generated to narrow the search space. Not all of the candidates have to be used in the final segmentation, but the final segment borders have to be selected from the set. The main requirement for the border candidate generation method is to be able to detect as many of the true borders as possible while keeping the total amount of the borders still reasonable.

The border candidates are generated using the novelty detection method from [3]. A Gaussian tapered checkerboard kernel is correlated along the main diagonal of the SDMs to produce novelty vectors. The vectors from different features are summed and peaks in the resulting vector are searched using a median based dynamic thresholding.

2.5 Segment Distance Measures

All segments between all pairs of border candidates are generated. These segments, when replicated with all possible groupings, form the set $\mathbb{S}$, from which the final description is a subset. The fitness measure for the structure description operates on probabilities that two segments in the piece are occurrences of the same musical part. The probability is obtained by matching the features of the segments.

The matched two segments s_m and s_n define a submatrix $\tilde{D}_{[m,n]}$ of the SDM. Two distance measures for the segment pair are defined using this submatrix: *stripe* and *block* distances. The stripe distance $d_S(s_m, s_n)$ is calculated by finding the minimum cost path through $\tilde{D}_{[m,n]}$ with dynamic programming. No transition cost is applied, but instead the total path cost is the sum of the elements along the path. The local path constraint forces the path to take one step in one or both directions at a time. The distance measure is obtained by normalising the total path cost with the maximum of the two submatrix dimensions. The block distance $d_B(s_m, s_n)$ is calculated as the average element value in the submatrix $\tilde{D}_{[m,n]}$.

2.6 Probability Mapping

Given the acoustic distance $d(s_m, s_n)$ [2] between two segments s_m and s_n, it is possible to define the probability $p(s_m, s_n)$ that the segments belong to the same group (are occurrences of the same part) using a sigmoidal mapping

$$p(s_m, s_n) = p(g_m = g_n | d(s_m, s_n) = \delta) \quad (1)$$

$$= \frac{1}{1 + e^{A\delta + B}}, \quad (2)$$

where δ is the observed distance value. The sigmoid parameters A and B are determined using two-class logistic regression with Levenberg-Marquardt algorithm [13].

The probabilities obtained from the mapping of all six distance values are combined with geometric mean. Heuristic restrictions on the segment groupings can be enforced by adjusting the pairwise probabilities. An example of such restriction is to set the segment pair probability to zero if the segment lengths differ too much (ratio 6/5 was used as the limit in the evaluations). The aggregated probability value after adjustments is denoted by $\hat{p}(s_m, s_n)$.

2.7 Fitness of a Description

A probabilistic fitness measure for different structure description candidates is defined using the segment pair probabilities in the description E by

$$P(E) = \sum_{m=1}^{M} \sum_{n=1}^{M} A(s_m, s_n) L(s_m, s_n), \quad (3)$$

where

$$L(s_m, s_n) = \begin{cases} \log(\hat{p}(s_m, s_n)), & \text{if } g_m = g_n \\ \log(1 - \hat{p}(s_m, s_n)), & \text{if } g_m \neq g_n \end{cases}. \quad (4)$$

The weighting factor $A(s_m, s_n)$ is the number of elements in the submatrix $\tilde{D}_{[m,n]}$ defined by the two segments. It is used to enable comparing descriptions with different number of segments, and its intuitive motivation is "the need to cover the whole area of the SDM".

The main concepts related to the calculation are illustrated in Figure 3. The short ticks in top and left side of the figure are the locations of chunk borders. The description claims that the piece consists of five segments (A_1, A_2, B_1, B_2, C) of varying lengths, and that segments A_1 and

[2] Similar definition applies to all three feature sets and two distance measures.

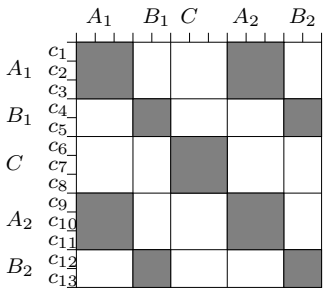

Figure 3. Illustration of the concepts regarding the calculation of the overall probability of a description. See text for details.

A_2 belong to same group as well as B_1 and B_2. There is a probability value assigned to each of the 25 submatrices, denoting the acoustic probability that the segments defining the submatrix belong to the same group. When evaluating Eq. (4), the upper alternative is used with the shaded submatrices, whereas the lower alternative is used for all the other submatrices.

The task to solve is to find the description E maximising the total log-probability of Eq. (3), given the acoustic information embedded to the pairwise probabilities

$$E_{OPT} = \underset{\mathrm{E}}{\operatorname{argmax}} \{P(E)\}. \tag{5}$$

2.8 Segment Labelling

The found description E_{OPT} defines a segmentation of the piece to musical sections and grouping of the segments which are occurrences of the same part. However, the groups have no musically meaningful labels. From the application point of view, the knowledge of the "musical role" of each part would be valuable. Musical pieces tend to follow certain forms when considering the sequences formed by the part names, e.g., "intro, verse, chorus, verse, chorus, chorus". These sequential dependencies are here modelled with N-grams of length 3. The 12 most often occurring musical part labels cover 90% of all part occurrences in the data set and are retained; the other labels are replaced with an artificial label "MISC". The estimated trigrams are smoothed using Witten-Bell smoothing [15]. The musicological information can be used in a post-processing stage to label the segments, or integrated into the fitness measure.

2.8.1 Labelling as Post-processing

In post-process labelling, the description found using Eqs. (3)–(4) is handled as a sequence of parts. Each of the groups in the description is mapped on trial with a unique musical part label in the trained vocabulary. The probability over the resulting label sequence is evaluated by calculating the cumulative Markov probability over the sequence and the most probable labelling is chosen. The post-processing labelling method is described in more detail in [11].

2.8.2 Integrated Labelling Model

As earlier experiments have shown the N-gram model to be informative in labelling the structural descriptions, an attempt is made to utilise this information already in the search of the descriptions. This is done by modifying the fitness measure of Eq. (3) to include a term containing the probabilities p_N from the N-grams:

$$P(E) = \sum_{m=1}^{M}\sum_{n=1}^{M} A(s_m, s_n) L(s_m, s_n) \tag{6}$$
$$+ \frac{w}{M-1} \sum_{o=1}^{M} \log\left(p_N(g_o|g_{1:(o-1)})\right) \sum_{m=1}^{M}\sum_{n=1}^{M} A(s_m, s_n),$$

where w is the relative weight given for the labelling model. [3] In effect, the additional term is the average part label transition log-probability weighted with the total area of the SDM. The labelling model likelihoods are normalised with the number of transitions $(M-1)$ to ensure that descriptions with different number of parts would have equal weight for the musicological model.

3 SEARCH ALGORITHM

Given all segments, segment pairs, and the probabilities that a pair of segments are from the same group, the search algorithm attempts for find the description E_{OPT} maximising the total fitness of Eq. (3) or Eq. (6). This section describes a novel search algorithm *Bubble token passing* (BTP) solving the optimisation task. An exhaustive search over all descriptions as is possible, but it is computationally too heavy for practical applications even with admissible search bounding. The exhaustive search was used to verify the operation of the greedy BTP algorithm presented next.

The proposed BTP can be seen as a variant of the N-best token passing (TP). The states in the algorithm are formed by the set $\mathbb{S}$ of all segments and all possible group assignments to each segment. In other words, the problem is finding the best path in a directed acyclic graph where the vertices are the segments with group information and there is an edge from a segment to the segments that start from the same border the previous ends. Because of the form of the total fitness measure of Eq. (3) or Eq. (6), the edge weights depend on the earlier vertices occupied by the path. Therefore, more efficient dynamic programming algorithms can not be used.

In the conventional TP, tokens are propagated between states time synchronously. Tokens record the travelled path and the associated path probabilities. At each time instant, the states take the best contained token, copy it and propagate it to all connecting states updating the token path and

[3] Values in the range 0.02–0.3 were noted to be most suitable in experiments with a subset of the evaluation data.

probability. When tokens are arriving to a state, they are sorted and only the best one is retained to be propagated at the next iteration. In N-best TP the N best tokens are propagated. [16]

The BTP operates as follows: an initial token is set to a start state, it is replicated and propagated to all connecting states updating the token information. The states store β best tokens that have arrived and propagate α tokens at the next iteration. If $\alpha < \beta$, the tokens not propagated remain in the state and they are considered for propagation in the next iteration. After some iterations, tokens start arriving to the end state. They contain the found paths (structural descriptions) and the related probabilities (fitness measures). As iterations are continued, more tokens "bubble" through the states to the end state and more description alternatives are found. It is likely that the found paths initially improve, but after a while the improving stops. The iterations can be stopped at any time, e.g., if the solution has converged or there are no more tokens in the system.

The proposed algorithm contains some beneficial properties mostly based on the two parameters α and β. As the tokens are propagated in a best-first order, the algorithm finds some reasonable solution fast and then continues improving it. The search "beam width" can be controlled with α, while β controls the the overall greediness. If the two are equal the algorithm is approximately the N-best TP. If β is set to infinity and the search is run until all tokens are out of the system, the search is exhaustive and guaranteed to find the global optimum. In experiment it was found to be sufficient to propagate a moderate amount ($\alpha = 5 - 50$) of tokens at a time while retaining larger amount of them ($\beta = 50 - 500$). With these values, the BTP was noted to find the same result as the exhaustive search in almost all of the test cases with considerably less computational cost. The exact values depend on the number of possible states and their connectivity.

When the description labelling done as a post-processing step, the search can be optimised by occupying different groups in order which eliminates much of the search space. [4]

4 RESULTS

The proposed algorithm is evaluated using 557 manually annotated pieces. The evaluation metrics are calculated from different evaluation aspects. The effect of the segmentation and border candidate generation is evaluated. In addition to them, both alternatives for the musical part labelling are evaluated. By comparing these two results, it is possible to test if the musicological knowledge is able to provide useful information for deciding which description is the best.

[4] With the number of border candidates used in the evaluations (32), and possible labels (13), there are approximately $3.1 \cdot 10^{35}$ different descriptions E. With post-processing labelling the amount is reduced to $1.1 \cdot 10^{26}$.

4.1 Data

The evaluation data set consists of 557 Western popular music pieces. [5] The pieces were selected to provide a representative sample of the music played on mainstream radio. The pieces are mainly from the pop/rock genre, but also some pieces from jazz, blues and schlager are present. For each piece, the sectional form was annotated by hand by segmenting and labelling the musical parts. The annotations were made by two research assistants with some musical background. The simulations were run using 10-fold cross-validation scheme. At each iteration, the training subset was used to train the labelling model N-grams and the distance to probability mapping function parameters. The presented results are averaged over all folds.

4.2 Evaluation Metrics

Two different metrics are used in the evaluations: frame pairwise grouping F-measure, and total frames labelled correctly. The first measure has been used in evaluation of a structure analysis algorithm in [7]. It considers all beat-frame pairs and whether or not the frames in the pair are assigned correctly in the same group. The recall rate R_r is calculated as the ratio of correctly found frame pairs with the frames assigned to the same group to the number of all segments pair from the same group. The precision rate R_p is the ratio of correctly found frame pairs to the claimed frame pairs. The F-measure is calculated from these two as $F = 2R_pR_r/(R_p + R_r)$.

The second used evaluation metric is motivated by the desired final output of the system: the result should be a segmentation to musical parts and the labelling of each segment with the appropriate musical part name. The measure is the amount of the piece assigned with the correct musical part name.

4.3 Evaluation Results

The evaluation results are given in Table 1. Results are shown for six different system configurations. First, the effect of segment boundary candidate generation (see Sec. 2.4) is studied by considering three different cases:

- "full": Fully automatic system which generates the segment border candidates using the novelty vector. The system has to decide at each candidate border should is be included, group the created segments, and label the groups. (This configuration uses all features except rhythmogram stripes.)
- "bord": The border candidate set is taken from the annotated segmentation points. Otherwise the same as above. (Uses MFCC and chroma stripes.)

[5] Full list of the pieces is available at <http://www.cs.tut.fi/sgn/arg/paulus/TUTstructure07_files.html>.

System	F (%)	R_p (%)	R_r (%)	label hit (%)
full w/ LM	61.4	64.2	63.9	39.1
full post-LM	61.7	65.9	63.1	36.1
bord w/ LM	77.0	80.5	77.8	46.8
bord post-LM	77.9	81.8	78.2	45.4
segs w/ LM	86.1	96.0	80.6	49.3
segs post-LM	86.2	95.6	81.1	46.5

Table 1. Evaluation results for six different system configurations and four evaluation measure. See text for details.

- "segs": The segmentation is taken from the ground truth. The system has to group the different occurrences of a part together and label the groups. (Uses MFCC and chroma stripes.)

Secondly, the strategies of using the musicological model are: "w/ LM" when then labelling is done during the description search using Eq. (6), and "post-LM" when the description search is done with Eq. (3) and the labelling is done as post-processing.

The results indicate that the method for generating the border candidates is currently a bottle-neck for the system, and should be considered in future work. The integrated usage of musicological model in the fitness measure does not seem to have a big effect on the result when looking at the F-measure (the difference is statistically insignificant, the level of $p < 0.05$ would require F-measure difference of at least 1.5 %-units). From the point of view of the label hit measure, the integrated labelling improves the results slightly, but the improvement is statistically significant ($p < 0.05$) only in the results obtained from fully automatic system. Comparing the results for "full" with [7] where the same F-measure was used, the performance is very similar, although it should be noted that differring data sets were used.

5 CONCLUSIONS

A method for analysing the sectional form of a music piece was presented. The method uses a probabilistic fitness measure for comparing different structural descriptions. This allows the focus of the development work to be concentrated on the definition of the fitness measure and its terms which is typically more intuitive and conceptually simpler than algorithm development. A musicological model of the sequential dependencies of musical parts was integrated to the proposed fitness measure. The integration improves the performance of the system when measuring the amount of time where the piece is assigned with the correct musical part label. A novel algorithm was presented for searching a description which maximises the defined fitness measure. The algorithm can be controlled with two intuitive parameters and its average- and worst-case performance is considerably better than that of an exhaustive search. Future work will concentrate on improving the method that generates segment border candidates since it seems to be a bottleneck of the current system.

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AUTOMATIC IDENTIFICATION OF SIMULTANEOUS SINGERS IN DUET RECORDINGS

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ABSTRACT

The problem of identifying singers in music recordings has received considerable attention with the explosive growth of the Internet and digital media. Although a number of studies on automatic singer identification from acoustic features have been reported, most systems to date, however, reliably establish the identity of singers in solo recordings only. The research presented in this paper attempts to automatically identify singers in music recordings that contain overlapping singing voices. Two approaches to overlapping singer identification are proposed and evaluated. Results obtained demonstrate the feasibility of the systems.

1. INTRODUCTION

In music recordings, the singing voice usually catches the listener's attention better than other musical attributes such as instrumentation or tonality. The singer's information, therefore, is essential to people for organizing, browsing, and retrieving music recordings. Most people use singer's voice as a primary cue for identifying songs, and performing such a task is almost effortless. However, building a robust automatic singer identification system is a difficult problem for machine learning. One of the challenges lies in training the system to discriminate among the different sources of sounds in music recordings, which may include background vocal, instrumental accompaniment, background noise, and simultaneous, or overlapping singings.

Although studies on automatic singer identification from acoustic features have been reported, most systems to date, however, reliably establish the identity of singers from recordings of solo performances only [1][2][3]. Tsai *et al*., in [4], investigated automatic detection and tracking of multiple singers in music recordings. However, the study only considered singing by multiple singers who performed in a non-overlapping matter. Other works related to the problem of singer identification include speech overlapping [5][6] in multi-speakers environments and voice separation from music accompaniment [7][8]. The research presented here attempts to automatically identify singers in music recordings that contain both simultaneous and non-simultaneous singings. We refer to this problem as overlapping singer identification (OSID).

2. APPLICATIONS

OSID can be applied in a number of areas. For example, a successful OSID system can be used as an automatic tool to locate, identify, and index singers in music recordings, thus reducing, if not replacing, human documentation efforts. OSID, moreover, can be applied in the context of karaoke. A personalized Karaoke system has been developed by Hua et al. [9] to create background visual content using home video and/or photo collections. As an extension to the current Karaoke system, OSID can be used to identify and index singers in their recorded karaoke songs, and using intelligent transformation and transition effects, the singing portions can be aligned with the respective singer's home video and/or photo collections to create a seamless personalize music video. In addition, OSID can be applied in the area of copyright protection and enforcement. Content description such as singer names, among many other metadata, is fundamental to the content and rights managements of digital music. OSID can be applied to generate singer information automatically and be integrated into the existing protection technologies to enhance the current copyright protection solutions.

3. PROBLEM FORMULATION

Since it is difficult to consider all application scenarios in an initial development stage, we began this research by identifying the important factors that influence the effectiveness of OSID. We then defined the scope of this preliminary study. The factors we identified include the following:

i) **Multiplicity.** Depending on the number of singers, a music recording may be classified as a pure instrumental, solo, duet, trio, band, or choral performance. In general, the complexity of an OSID

problem grows as the number of singers in a music recording increases. This study focuses on vocal duets, i.e., the OSID system determines the identity of a singer or singers who sang in a given music recording.

ii) **Overlapping duration percentage.** Although two singers perform a duet, they may not always be singing simultaneously. Therefore, an excerpt from a music recording can be a) an instrument-only segment, b) a solo-singing segment, or c) an overlapping-singing segment. To simplify the problem, the test recordings used in this study are either b) or c), with the overlapping duration percentages of 0% or 100%, respectively.

iii) **Overlapping energy ratio.** As in many bands, one or more musicians in addition to the lead singer often sing background vocal while they play their instruments. The audio signal energy of the background singer(s), therefore, may be very low compared to that of the lead singer. In such a case, identifying the background singers would be more difficult. For this preliminary study, no test recordings have background singers, and the singers in our test recordings all sing with roughly equal signal energies.

iv) **Tune/lyrics variations.** Multiple singers performing simultaneously may sing in a) exactly the same tune and lyrics, b) exactly the same tune but different lyrics, c) different tunes but exactly the same lyrics, or d) different tunes and different lyrics. We consider only cases a) and d) for this study.

v) **Background accompaniment.** A majority of popular music contains background accompaniment that inextricably intertwines singers' voice signals with a loud, non-stationary background music signal. During this initial stage of the development, we do not attempt to solve the problem of background interference but only deal with vocal music that has no background accompaniments.

vi) **Open-set/close-set.** The OSID problem at hand is a close-set classification problem, which identifies the singer(s) among a set of candidate singers. This study does not discuss the problem of open-set classification, which determines whether the singer(s) identified is/are among the candidate singers performed in a set of test recordings.

vii) **Audio quality.** Although most test recordings are taken from high quality sources such as CDs, many often undergo signal degradation due to audio filtering, encoding/decoding, or noise corruption. A successful OSID system should be robust against various signal distortions. This study, however, places this issue aside because audio quality is an inevitable problem for most music classification research. The music data we use for this study are all of high-quality audio recorded on PC.

4. DATA

Since no public corpus of music recordings meets the specific constraints of the OSID problem defined here, a small database of test recordings was created. The database contains vocal recordings by ten male amateur singers, aged between 20 and 35. Every singer was asked to perform 30 passages of Mandarin pop songs with a Karaoke machine. The duration of each passage ranges from 13 to 20 seconds.

The database was divided into two subsets, one for training the OSID system and the other for evaluating the system. The training subset consists of the first 15 passages, while the evaluation subset consists of the remaining 15 passages. Passages of the pop songs were recorded in a quiet room. The Karaoke accompaniments were output to a headset, and thus not recorded. All the passages were recorded at 22.05 kHz, 16 bits, in mono PCM wave.

Test recoringdgs of duets were then obtained by mixing the wave files sung by a pair of singers. Two sets of recordings (i.e., for training and evaluation), sung by 45 (C_2^{10}=45) different pairs of singers, were created. One set included 675 ($C_2^{10} \times 15$ =675) recordings of duets sung in exactly the same tune and with the same lyrics; the other set included 4,725 ($C_2^{10} \times C_2^{15}$ =4,725) recordings of duets sung in different tunes and with different lyrics.

To facilitate the discussions in the following sections, thereafter, recordings of duets sung in exactly the same tune and with the same lyrics are referred to as "STSL duet recordings." Similarly, recordings of duets sung in different tunes and with different lyrics are referred to as "DTDL duet recordings."

5. METHODOLOGY

Following the most popular paradigm of stochastic pattern recognition, we propose two approaches to OSID system design.

5.1. Two-Stage OSID System

Figure 1 shows the block diagram of our two-stage OSID system. The system first consists of a "Solo/Duet Recognition" component. If a solo singing is recognized, the problem becomes the conventional single singer identification. If a duet singing is recognized, a "Duet Singer Identification" component handles the case. Each of the components in this system is a Gaussian mixture classifier [9].

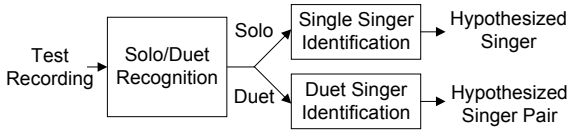

Figure 1. Two-stage OSID system.

5.1.1. Solo/Duet Recognition

Figure 2 shows the block diagram of the "Solo/Duet Recognition" component. The component is divided into two phases: training and testing. During the training phase, two Gaussian Mixture Models (GMMs), λ^s and λ^d, are created, where λ^s represents the acoustic pattern of a solo singing passage while λ^d represents the acoustic pattern of a duet singing passage. Combinations of Gaussian densities generate a variety of acoustic classes, which, in turn, reflect certain vocal tract configurations. The GMMs provide good approximations of arbitrarily shaped densities of spectrum over a long span of time [9]. Parameters of a GMM consist of means, covariances, and mixture weights. λ^s is generated from all solo singing passages and λ^d is generated from all duet singing passages. Then prior to Gaussian mixture modeling, singing waveforms are converted into Mel-scale frequency cepstral coefficients (MFCCs). In the testing phase, an unknown test recording is converted into MFCCs and then tested for λ^s and λ^d. The results are based on likelihood probabilities, $\Pr(\mathbf{X}|\lambda^s)$ and $\Pr(\mathbf{X}|\lambda^d)$, where the recording is hypothesized as a duet singing passage (or a solo singing passage) if $\log\Pr(\mathbf{X}|\lambda^d)$–$\log\Pr(\mathbf{X}|\lambda^s)$ is larger (or smaller) than a pre-set threshold η.

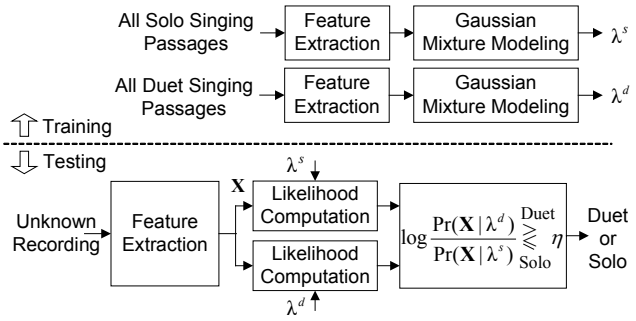

Figure 2. Solo/Duet recognition.

5.1.2. Single Singer Identification

Figure 3 shows the "Single Singer Identification" component. If there are N different candidates singers, then N GMMs, λ_1, λ_2, …, λ_N, are created to represent the acoustic patterns of their singings. When an unknown recording is received at the system input, the component calculates and decides in favor of singer I^* when the condition in Eq. (1) is satisfied:

$$I^* = \arg\max_{1\le i\le N} \Pr(\mathbf{X}|\lambda_i) \quad (1)$$

5.1.3. Duet Singer Identification

The "Duet Singer Identification" component is similar to the "Single Singer Identification" component. The only difference between the two is that the GMMs of solo singers are replaced with the GMMs of pairs of singers. However, generating the GMMs of pairs of singers is not as straightforward as generating the GMMs of solo singers, because it may be impractical to collect singing data from every possible combination of pairs of singers. Hence, two approaches were taken to sidestep the collection of real simultaneous singing data. The first approach uses direct waveform mixing, which is shown in Figure 4.

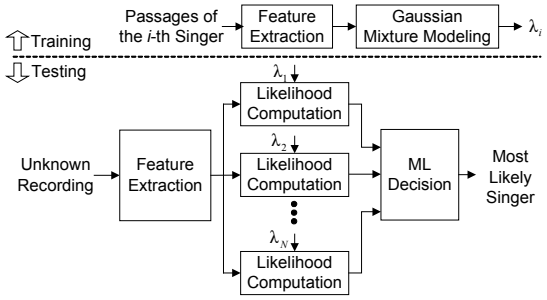

Figure 3. The "Single Singer Identification" component.

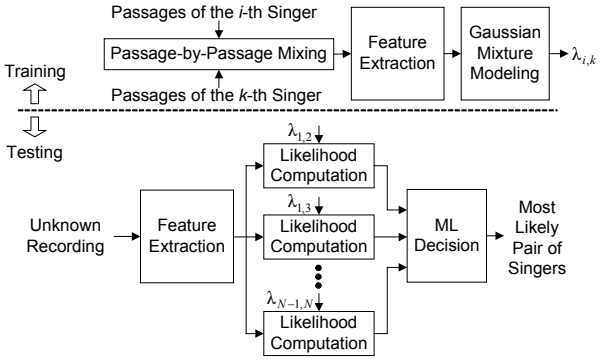

Figure 4. The "Duet Singer Identification" component using direct waveform mixing.

In the training phase of this system, audio waveforms from every pairs of singers are mixed, based on roughly equal energies, to simulate real duet singings. The resulting waveforms are then converted into MFCCs. For each pair of singers, a GMM is built using these features. Hence, for a population of N candidate singers, a total of $C_2^N = N!\,/\,[2!(N-2)!]$ singer-pair GMMs $\lambda_{i,j}$, $i \ne j$, $1 \le i, j \le N$ are created. In the testing phase, an unknown audio recording is converted into MFCCs and then tested for each of the C_2^N GMMs. The system then determines the most-likely singer pair (I^*, J^*) performed in the recording based on the maximum likelihood decision rule:

$$(I^*, J^*) = \arg\max_{1\le i,j\le N, i\ne j} \Pr(\mathbf{X}|\lambda_{i,j}). \quad (2)$$

One shortcoming of the direct waveform mixing approach is that the training process can become very cumbersome if the number of candidate singers is large or

if a new singer needs to be added. As an alternative to this problem, a second approach based on Parallel Model Combination (PMC) technique [10] is used, as shown in Figure 5. Given a set of N solo singer GMMs, each GMM is used to generate C_2^N singer-pair GMMs. Since duet singing signals overlap in the time/frequency domain while the GMMs are in the cepstral/quefrency domain, the parameters of the GMMs need to be converted to the linear spectral/frequency domain before they can be added.

In addition, since two K-mixture GMMs would result in a large $K \times K$-mixture GMM, UBM-MAP [11] is used to control the size of the resulting GMM K-mixture. The basic strategy of UBM-MAP is to generate a universal GMM using all solo singers' data, and then adapt the universal GMM to each solo singer GMM based on maximum a posterior (MAP) estimation. Since all of the solo singer GMMs are adapted from the universal GMM, the mixtures of the GMMs are aligned. Thus, we do not need to consider the combination of the k-th Gaussian of one GMM with the ℓ-th Gaussian of another GMM, where $k \neq \ell$, but we only need to consider the case when $k = \ell$. For a pair of singers i and j, the combined mean and covariance of the k-th mixture is computed by

$$\boldsymbol{\mu}_{i,j}^{k} = \mathbf{D}\left\{\log\{\exp(\mathbf{D}^{-1}\boldsymbol{\mu}_i^{k}) + \exp(\mathbf{D}^{-1}\boldsymbol{\mu}_j^{k})\}\right\}, \quad (3)$$

$$\boldsymbol{\Sigma}_{i,j}^{k} = \mathbf{D}\left\{\log\{\exp[\mathbf{D}^{-1}\boldsymbol{\Sigma}_i^{k}(\mathbf{D}^{-1})'] + \exp[\mathbf{D}^{-1}\boldsymbol{\Sigma}_j^{k}(\mathbf{D}^{-1})']\}\right\}, \quad (4)$$

where $\boldsymbol{\mu}_i^{k}$ and $\boldsymbol{\Sigma}_i^{k}$ are the mean vector and covariance matrix of GMMs λ_i, respectively; $\mathbf{D}$ represents the discrete cosine transform matrix; prime ($'$) denotes the transpose.

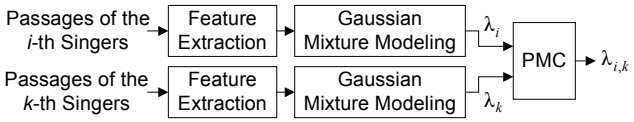

Figure 5. The training phase of "Duet Singer Identification" component based on parallel model combination.

5.2. Single-Stage OSID System

As an alternative approach, we present a second system that combines the three components in the Two-Stage OSID system. The system unifies the three components into a Single-Stage system, eliminating the stage of first determining if a test recording is a solo or duet singing performance. This is done by using the N single-singer GMMs from the Single-Singer Identification and the C_2^N singer-pair GMMs from the Duet-Singer Identification to build a unified classifier with $(N + C_2^N)$ GMMs. In the testing phase, an unknown recording is converted into MFCCs and then tested for each of the $(N + C_2^N)$ GMMs. Then, if we denote each single-singer GMM λ_i as $\lambda_{i,i}$, $1 \leq i \leq N$, the system should decide in favor of singer(s) (I^*, J^*) if the condition in Eq. (5) is satisfied.

$$(I^*, J^*) = \arg\max_{1 \leq i,j \leq N} \Pr(\mathbf{X} \mid \lambda_{i,j}), \quad (5)$$

Note that if $I^* = J^*$, then the recording is hypothesized to be performed by a single singer I^*.

6. EXPERIMENTS AND RESULTS

6.1. Solo/Duet Recognition Experiments

The first experiment conducted examined the validity of the solo/duet recognition component. There were 150 solo test recordings, 675 STSL duet test recordings, and 4,725 DTDL duet test recordings, with a total of 5,550 test recordings. The recognition accuracy was measured by

$$\frac{\#\text{Correctly} - \text{recognized Recordings}}{\#\text{Testing Recordings}} \times 100\%.$$

Table 1 shows the recognition results with respect to the different numbers of Gaussian mixtures in λ^s and λ^d. We can see that most of the recordings were correctly recognized.

No. of Mixtures	Accuracy
16	96.1%
32	94.2%
64	95.2%

(a) Recognition accuracy

	Actual	
Classified	Solo	Duet
Solo	99.3%	4.6%
Duet	0.7%	95.4%

(b) Confusion matrix of the 16-mixture case

Table 1. Solo/duet recognition results.

6.2. Single-Singer Identification Experiments

For the purpose of comparison, experiments of the conventional SID for solo recordings were also conducted. The identification accuracy was measured by

$$\frac{\#\text{Correctly} - \text{identified Recordings}}{\#\text{Testing Recordings}} \times 100\%.$$

Table 2 shows the results of singer identification in 150 recordings sung by 10 different singers. As the singer population was small, the result obtained was almost perfect.

No. of Mixtures	SID Accuracy
16	96.7%
32	98.7%
64	100.0%

Table 2. Results of singer identification for solo recordings.

6.3. Duet-Singer Identification Experiments

Then the feasibility of OSID in duet recordings was examined. In these experiments, test data consisted of 675 + 4,725 duet singing wave files, i.e., no solo recordings were considered. Here the performances of the direct waveform mixing and the PMC methods of Duet-Singer Identification component were evaluated.

Depending on the context of application, the performance of OSID is evaluated differently. This study considers two types of OSID accuracy. The first one takes into account the number of *singer pairs* identified correctly. Specifically,

$$\text{Acc.1 (in \%)} = \frac{\#\text{Correctly} - \text{identified Singer Pairs}}{\#\text{Testing Recordings}} \times 100\%.$$

The second one takes into account the number of *singers* identified correctly. Specifically,

$$\text{Acc. 2 (in \%)} = \frac{\#\text{Correctly} - \text{identified Singers}}{\#\text{Testing Singers}} \times 100\%.$$

For example, if a recording contains simultaneous singings by two performers, s_1 and s_2, and the identified singers are s_1 and s_4, then `#Correctly-identified Singer Pairs` = 0 and `#Correctly-identified Singers` = 1. Consequently, Acc. 2 is always higher than Acc. 1.

Tables 3 shows the OSID result obtained with direct waveform mixing methods. Here, the OSID results for four cases are presented: i) both training and testing data consist of STSL duet recordings; ii) training data consist of STSL duet recordings, while testing data consist of DTDL duet recordings; iii) training data consist of DTDL duet recordings, while testing data consist of STSL duet recordings; iv) both training and testing data consist of DTDL duet recordings.

It can be seen from Table 3 that OSID using STSL duet recordings for training always outperformed than those that using DTDL duet recordings for training. Similarly, the performance of OSID using STSL duet recordings for testing was always better than those that using DTDL duet recordings for testing. When both training and testing data consist of STSL duet recordings, we obtained the best OSID performance, showing that 85.0% of singer pairs or 92.5% of singers in the testing data can be correctly identified.

Tables 4 shows the OSID result obtained with the PMC method. In this experiment, since no duet singing is required in the training process, we considered two cases: i) testing data consist of STSL duet recordings; ii) testing data consist of DTDL duet recordings. It can be observed in Table 4, that similar to the results in Table 3, the performance of OSID was always better when STSL duet recordings were used for testing. Comparing Table 4 with Table 3 (a) and (b), it can also be found that the direct waveform mixing method was superior to the PMC method when STSL duet recordings were used for testing. However, the PMC method performed better than the direct waveform mixing method when DTDL duet recordings were used for testing. This indicates that the PMC method is not only better at scaling up the singer population, but it is also better at generalizing the singer identification problem than the direct waveform mixing method.

No. of Mixtures	Acc. 1	Acc. 2
16	80.7%	90.1%
32	84.3%	92.1%
64	85.0%	92.5%

(a) Both training and testing data consist of STSL duet recordings

No. of Mixtures	Acc. 1	Acc. 2
16	67.9%	83.7%
32	69.8%	84.8%
64	73.6%	86. 7%

(b) Training data consist of STSL duet recordings, while testing data consist of DTDL duet recordings

No. of Mixtures	Acc. 1	Acc. 2
16	77.3%	88.7%
32	78.4%	89.3%
64	80.7%	90.4%

(c) Training data consist of DTDL duet recordings, while testing data are STSL duet recordings

No. of Mixtures	Acc. 1	Acc. 2
16	52.3%	75.8%
32	47.1%	73.4%
64	43.6%	71.6%

(d) Both training and testing data consist of DTDL duet recordings

Table 3. Results of identifying duet recordings based on direct waveform mixing method.

No. of Mixtures	Acc. 1	Acc. 2
16	75.1%	87.1%
32	75.1%	87.3%
64	78.1%	88.7%

(a) Testing data consist of STSL duet recordings

No. of Mixtures	Acc. 1	Acc. 2
16	71.1%	85.0%
32	69.9%	85.0%
64	75.3%	87.6%

(b) Testing data consist of DTDL duet recordings

Table 4. Results of identifying duet recordings based on PMC method.

6.4. Singer Identification Experiments: Solo and Duet Recordings

Lastly, for a more realistic case, a test recording, which may be either a solo singing or duet singing, was tested. There were 150 solo test recordings, 675 STSL duet test recordings, and 4,725 DTDL duet test recordings, with a total of 5,550 test recordings. The identification performance was characterized by Acc. 1 and Acc. 2, as before. However, if a recording contains only single singer s_1 but the system hypothesizes two singers s_1 and s_4, then `#Correctly-identified Singer Pairs` = 0 and `#Correctly-identified Singers` = 1.

Table 5 shows the results obtained by the proposed two OSID systems. The singer-pair GMMs used in this experiment were generated using the PMC method. The number of Gaussian mixtures was set to 64 for both solo singer and singer pair GMMs. Compared to the results in Table 4, it is observed that while more uncertainties are added in the testing data, the resulting accuracies only decrease slightly. In addition, it is also found that the Two-Stage OSID system performed better than the Single-Stage OSID system. This indicates that although the Single-Stage OSID system takes advantage of the simplicity in design, it pays the loss of accuracy that can be achieved with the Two-Stage OSID system.

System	Acc. 1	Acc. 2
Two-Stage OSID System	76.2%	88.1%
Single-Stage OSID System	73.0%	87.9%

Table 5. Results of identifying both solo and duet recordings.

7. CONCLUSION

This paper examined the feasibility of overlapping singer identification and compared two approaches to the problem of detecting and identifying singers in duet recordings. The research presented here extends previous works on singer identification. The systems proposed up-to-date only focus on the identification of singers in solo recordings. In reality, many music recordings involve multiple singers such as duet love songs, gospel music, or folk and country music. Encouraging results arrived at this initial stage of investigation laid a good foundation for the future development of a robust automatic singer identification system. Regarding future work, a wider variety of music will be acquired to scale up and further test the system.

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CHARACTERISATION OF HARMONY WITH INDUCTIVE LOGIC PROGRAMMING

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ABSTRACT

We present an approach for the automatic characterisation of the harmony of song sets making use of relational induction of logical rules. We analyse manually annotated chord data available in RDF and interlinked with web identifiers for chords which themselves give access to the root, bass, component intervals of the chords. We pre-process these data to obtain high-level information such as chord category, degree and intervals between chords before passing them to an Inductive Logic Programming software which extracts the harmony rules underlying them. This framework is tested over the Beatles songs and the Real Book songs. It generates a total over several experiments of 12,450 harmony rules characterising and differentiating the Real Book (jazz) songs and the Beatles' (pop) music. Encouragingly, a preliminary analysis of the most common rules reveals a list of well-known pop and jazz patterns that could be completed by a more in depth analysis of the other rules.

1 INTRODUCTION

The explosion of the size of personal and commercial music collections has left both content providers and customers with a common difficulty: organising their huge musical libraries in such a way that each song can be easily retrieved, recommended and included in a playlist with similar songs. Most of the metadata provided are hand-annotated by experts such as in AllMusicGuide [1] or the result of a community effort such as in MusicBrainz [2]. Because classifying large amounts of data is expensive and/or time-consuming, people are gaining interest in the automatic characterisation of songs or groups of songs. Songs can be characterised by their rhythm, harmony, structure, instrumentation, etc. In this article we present the first step towards a framework able to automatically induce rules characterising songs by various musical phenomena. For this study we are interested in the automatic extraction of harmony patterns.

Many of the existing approaches for pattern extraction and recognition make use of statistical descriptions [14]. However some of the attempts to automatically build descriptors of music explore the idea of using logical rules [6]. Such rules could be expressed as follows:

$$(X_1 \wedge X_2 ... X_n) \Rightarrow \textit{Musical Phenomenon} \qquad (1)$$

where the X_i are structured descriptions of local musical content and *Musical Phenomenon* is the high level property we are interested in. The logical framework offers several advantages over the statistical framework, for example temporal relations between local musical events can easily be expressed. Moreover, logical rules are human-readable. Thus, automatically extracted musical patterns expressed in logical formulae can be transmitted as they are to musicologists who can in turn analyse them.

To induce such logical rules we can either use statistical methods such as the C4.5 algorithm, a statistical decision tree learning algorithm, or relational methods such as Inductive Logic Programming (ILP) [5]. We choose to focus on relational induction of harmony rules with ILP. To test our framework we study the chord sequences of the Beatles and Real Book songs starting from a symbolic representation of these songs. However, the primary focus of this paper is on methodology and knowledge representation, rather than on the presentation of new musical knowledge extracted by the system.

The paper is organised as follows: In Section 2, we review some existing studies using logical rules for MIR. In Section 3 we explain our methodology to automatically extract logical harmony rules from manually annotated chords. In Section 4 the details and results of our automatic analysis of the Beatles and Real Book songs with ILP are presented and discussed before concluding in Section 5.

2 RELATED WORK

We now review some previous work in which automatically induced logical rules are used for musical retrieval tasks.

In [6] logical rules for characterising melody in MIDI files are built with random forest classifiers, a statistical approach. A pattern-based first order inductive system called PAL set up by Morales [4] learns counterpoint rules. The

[1] http://www.allmusic.com/
[2] http://musicbrainz.org/

system looks for patterns in the notes (described by their tone, height and voice) using a background knowledge restricted to the classification of intervals between two notes into perfect or imperfect consonant and dissonant, valid and invalid intervals.

Also many research projects have focused on automatically building expressive performance rules. For instance Widmer [16] develops an inductive rule learning system identifying both relevant and unknown rules of expressive performance from MIDI recordings of Mozart's sonatas performed by different pianists. His rules are based on the tempo, dynamics and articulation properties of the notes. Dovey [1] analyses and extracts rules from piano performances of Rachmaninoff using ILP. Subsequently, Van Baelen and De Readt [13] extend this work by transforming the characterisation rules into generative rules for MIDI.

Finally several musical rule derivation studies were conducted by Ramirez et al. [8, 10, 9]. In [8] an ILP based application learns rules in popular music harmonisation. They are constructed at a bar level (and not at a note level) to capture chord patterns. The structure (i.e. musical phrases) of the songs given as examples is manually annotated, which provides the system with a rich background knowledge containing not only local but also global information. Subsequently, Ramirez et al. [10] study Jazz performance from audio data with an ILP system. The system induces rules related to the duration transformation, onset deviation and energy of a note and the alteration of the score melody by adding or deleting notes. A background knowledge composed of information about the neighboring notes and the Narmour group(s) to which each note belongs is provided to the system. Finally, Ramirez et al. [9] implements a framework which analyses classical violin performance by means of both an ILP technique (the relational decision tree learner called TILDE) and a numerical method. Another component then uses these results to synthesise expressive performance from unexpressive melody descriptions.

3 METHODOLOGY

In search of chord idioms, Mauch et al. [3] made an inventory of chord sequences present in the Real Book and in the Beatles' studio albums. Their approach is entirely statistical and resulted in an exhaustive list of chord sequences together with their relative frequencies. To compare the results of our relational methodology with their results obtained with a statistical method we examine RDF (Resource Description Framework) descriptions of the two manually annotated collections from [3]:

- Harte's transcriptions [2] of the 180 songs featured on the Beatles' studio albums,
- transcriptions of 244 Jazz standards from the Real Book [15] [3] .

These transcriptions constitute a compact symbolic representation of the songs: the chords are manually labelled in a jazz/pop/rock shorthand fashion and their start and end times are provided.

The steps to extract harmony rules from these songs transcriptions are summarised as follows: First the RDF representation of the harmonic events is pre-processed and transcribed into a logic programming format that can be understood by an Inductive Logic Programming system. This logic programming representation is passed to the ILP software Aleph [11] which induces the logical harmony rules underlying the harmonic events.

3.1 Harmonic content description

The RDF files describing the Beatles and Real songs we study contain a structured representation of the harmonic events based on the Music Ontology [7]. Each harmonic event (or chord) is associated with a start time, an end time and a web identifier we can crawl to get the RDF description of the chord based on the Chord Ontology [4] .

We implemented a RDF chord parser to transcribe RDF chord representation into Prolog files that can be directly given as input to Aleph. For each of these chords it extracts the root note, bass note, component intervals, start time and end time from the RDF description. It then computes the chord category and degree of a chord and the root interval and bass interval between two consecutive chords. Intervals are measured upwards.

For this study we limit the chord categories (or chord types) to 'Major', 'minor', 'augmented', 'diminished', 'suspended', 'Dominant', 'neutral' (when the 3rd is neither present nor suspended) and 'unknown' (for every chord that does not belong to the previous categories). For each chord, the intervals are analysed by the RDF chord parser which then assigns the chord to one of these categories. First it reduces the chord to a 7th chord and checks if this reduced chord is a dominant 7th, in which case the chord is labeled 'Dominant'. Otherwise the chord is reduced to a triad and the type of this triad is kept as the chord category.

The degrees are computed by our RDF chord parser using the current key. Key information was added by hand when available. We only had access to tonality information for the Beatles, so no degree details were added for the Real Book songs. For the Beatles we performed two studies: one without degree over the whole set of songs and one with degree in which only the songs where there was no key modulation were kept. We also filtered out the songs which were not tonal songs (i.e. songs that were not following major or minor scales). The remaining songs constitute 73.9% of the Beatles' songs.

[3] these RDF files are available on http://chordtranscriptions.net/

[4] http://purl.org/ontology/chord

Our sole interest is in sequences of chords between which there is a harmonic modification (i.e. at least the root, bass or chord category differs from one chord to the next one). Although harmonic rhythm is important we leave it for future work.

3.2 Rule induction with ILP

The inductive concept learning task of Inductive Logic Programming can be described as follows. We provide a set of examples E containing positive (set E^+) and negative (set E^-) examples of a concept C and a background knowledge B all expressed in logic programs. Given this we find a hypothesis H (also expressed in logic program form) such that every positive example $e \in E^+$ is covered by $H \wedge B$ (completeness) and no negative example $e \in E^-$ is covered by $H \wedge B$ (consistency).

We are interested in chord sequences of length 4 as in Mauch et al. [3]. Four chord sequences is a typical phrase length for the studied corpora. This choice is also the result of an empirical process: we also studied shorter sequences, but the results were less interesting, i.e. characteristic of the corpus and presenting well-known patterns. For longer sequences, the extracted patterns are less general, i.e. have a smaller coverage and thus less characteristic of the corpus. The concept we want to characterise is the harmony of a set of songs e.g. all the Beatles songs, all the Real Book songs. Therefore the positive examples given to the ILP system are all the chord sequences of length 4 (predicate `chord_prog_4/4`) found in such a set of songs. These chord sequences overlap. For instance from a chord sequence of length 8 we extract 5 overlapping chord sequences of length 4. The background knowledge is composed of the descriptions of all the chords previously derived by the RDF chord parser.

In the ILP system we use to induce harmony rules (Aleph [11]), we can either provide negative examples of a concept (in our case, chord progressions of length 4 from another set of songs) or force Aleph to explain the positive examples using a well-designed negative example (we will refer to this mode as the *one negative example mode*). In the latter case our negative example consists of the first chord sequence of our corpus in which we exchanged the position of the first and second chords. It is a valid negative example because in our background knowledge each chord in each song is uniquely identified by a Prolog atom and the position of each individual chord relative to the other chords is stored. We found out that by limiting the set of negative examples to this very simple one we obtained a more complete set of rules than when using the *positive examples only mode* of Aleph which randomly generates a limited number of negative examples.

To generate hypotheses Aleph uses inverse entailment. It consists of selecting an uncovered example, saturating it to obtain a bottom clause and searching the space of clauses that subsumes this bottom clause in a top-down manner starting from the shortest clauses. Saturating an example means looking for all the facts that are true about this example (using the example itself and the background knowledge). The bottom clause is the disjunction of all these facts. The clause that covers the maximum number of examples and the minimum number of negative examples (i.e. which maximises a score function based on the number of positive and negative examples covered by this clause) is kept as a hypothesis. The examples covered by the found hypothesis are removed and the next uncovered example is selected to be saturated, and so on until no uncovered example is left. Finally Aleph returns a set of hypotheses that covers all the positive examples. This process is not deterministic. The set of generated rules depends on the order in which the examples are selected by Aleph (which is the order in which the examples are given to Aleph). So the resulting set of rules is only one of the sets of rules that could be induced from the set of examples. However since Aleph looks for the most general rules at each step, the final set of rules is a sufficient description of the data (it explains all chord sequences) and is non-redundant (no subset of the rules explains all the chord sequences). This minimal sufficient description of a data set could be very useful for classification purposes since only a few characteristics needs to be computed to classify a new example. This is one of the advantages of our method against the purely statistical method employed in [3] which only computes the frequencies of each chord sequence and does not try to build a sufficient model of the corpora.

To obtain meaningful rules we also constrain Aleph to look for a hypothesis explaining the chord progressions in terms of root note progressions (`root_prog_4/8`), bass note progressions (`bassNote_prog_4/8`), chord category progressions (`category_prog_4/8`), root interval progressions (`rootInterval_prog_3/7`), bass interval progressions (`bassInterval_prog_3/7`) and degree progressions (`degree_prog_4/8`).

4 EXPERIMENTS AND RESULTS

4.1 Independent characterisation of the Beatles and Real Book chord sequences

We run two experiments. In the first experiment we want to characterise the chord sequences present in the Beatles' songs and compare them to the chord sequences present in the Real Book songs. Therefore we extract all the chord sequences of length 4 in the Beatles' tonal songs with no modulation (10,096 chord sequences), all the chord sequences of length 4 in all the Beatles' songs (13,593 chord sequences) and all the chord sequences of length 4 from the Real Book songs (23,677 chord sequences). Then for each of these sets

Rule	C_1	C_2
1. maj → maj → maj → maj	4752 (35%)	3951 (39%)
2. maj → maj → maj → min	632 (4.65%)	431 (4.27%)
3. min → maj → maj → maj	628 (4.62%)	448 (4.44%)
4. • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf5th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ •	586 (4.31%)	-
5. • $\overset{/\text{perfU}}{\longrightarrow}$ • $\overset{/\text{perfU}}{\longrightarrow}$ • $\overset{/\text{perfU}}{\longrightarrow}$ •	584 (4.30%)	-
6. maj → min → maj → maj	522 (3.84%)	384 (3.80%)
7. maj → maj → min → maj	494 (3.63%)	363 (3.60%)
8. • $\overset{/\text{perf5th}}{\longrightarrow}$ • $\overset{/\text{perf4th}}{\longrightarrow}$ • $\overset{/\text{perf5th}}{\longrightarrow}$ •	463 (3.41%)	346 (3.43%)
9. maj → maj → min → min	344 (2.53%)	217 (2.15%)
10. • $\overset{\text{perfU}}{\longrightarrow}$ • $\overset{\text{perfU}}{\longrightarrow}$ • $\overset{\text{perfU}}{\longrightarrow}$ •	336 (2.47%)	237 (2.38%)
11. min → min → maj → maj	331 (2.44%)	216 (2.14%)
12. maj → min → min → maj	308 (2.27%)	197 (1.95%)
13. • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{maj2nd}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ •	260 (1.91%)	209 (2.07%)
14. • $\overset{\text{maj2nd}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ •	251 (1.85%)	195 (1.93%)

Table 1. Beatles harmony rules whose coverage is larger than 1.75%. C_1 and C_2 represent the positive coverage over all the Beatles songs and over the Beatles tonal songs with no modulation respectively. "perfU" means perfect unison.

of chord sequences we induce rules characterising them using the *one negative example mode* in Aleph.

Our system induces a set of 250 rules for each of the Beatles collections (tonal songs, all songs) and a set of 596 rules for the Real Book. The positive coverage of a rule is the number of positive examples covered by this rule. We want to consider only the patterns occurring in multiple songs (i.e. the ones characteristic of the corpus). For that we leave out the rules with a too small coverage (smaller than 0.5%). Because of space limitation we also only show the top rules for each experiment but a complete list of rules is available upon request. The top rules for our first experiment are shown in Tables 1 and 2.

For readability purposes we only show a compact representation of the body of rules:

- degrees are represented with roman numerals,
- "/ " refers to the bass note as in jazz chord notation,
- the intervals between roots (written first) or bass notes of the chords (symbolised by "/") are put on top of the arrows,
- a bullet symbolises the absence of information about some characteristics of the chord.

In accordance with conclusions in [3], some patterns extracted in these experiments are very common pop and jazz harmonic patterns. For instance, the Beatles rule with the highest coverage (more than a third of the chord sequences) is maj → maj → maj → maj. The minor chord is the second

Rule	C
1. • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ •	1861 (7.86%)
2. min → dom → min → dom	969 (4.09%)
3. min → dom → maj → min	727 (3.07%)
4. dom → min → dom → min	726 (3.07%)
5. min → min → min → min	708 (2.99%)
6. dom → dom → dom → dom	674 (2.85%)
7. • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perfU}}{\longrightarrow}$ •	615 (2.60%)
8. • $\overset{\text{maj6th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ •	611 (2.58%)
9. • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf5th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ •	608 (2.57%)
10. dom → min → dom → maj	594 (2.51%)
11. dom → maj → min → dom	586 (2.47%)
12. • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perfU}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ •	579 (2.45%)
13. • $\overset{/\text{maj6th}}{\longrightarrow}$ • $\overset{/\text{perf4th}}{\longrightarrow}$ • $\overset{/\text{perf4th}}{\longrightarrow}$ •	547 (2.31%)
14. maj → min → dom → maj	478 (2.02%)
15. • $\overset{/\text{maj7th}}{\longrightarrow}$ • $\overset{/\text{perf4th}}{\longrightarrow}$ • $\overset{/\text{perf4th}}{\longrightarrow}$ •	477 (2.01%)
16. • $\overset{/\text{perf4th}}{\longrightarrow}$ • $\overset{/\text{maj6th}}{\longrightarrow}$ • $\overset{/\text{perf4th}}{\longrightarrow}$ •	440 (1.86%)
17. • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{perf4th}}{\longrightarrow}$ • $\overset{\text{maj6th}}{\longrightarrow}$ •	436 (1.84%)
18. min → dom → maj → dom	424 (1.79%)

Table 2. Real Book harmony rules whose coverage is larger than 1.75%. C is the positive coverage.

	Rule	C_1	C_2
1.	maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj	3.13%	3.79%
	I maj → IV maj → I maj → IV maj	-	2.47%
	V maj → I maj → V maj → I maj	-	1.00%
2.	maj $\overset{\text{perf5th}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj	2.94%	3.61%
	IV maj → I maj → IV maj → I maj	-	2.43%
	I maj → V maj → I maj → V maj	-	0.84%
3.	maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{maj2nd}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj	1.38%	1.75%
	I maj → IV maj → V maj → I maj	-	1.59%
4.	maj $\overset{\text{maj2nd}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj	1.21%	1.47%
	IV maj → V maj → I maj → IV maj	-	1.15%
5.	maj $\overset{\text{perf5th}}{\longrightarrow}$ maj $\overset{\text{min7th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj	1.04%	1.28%
	I maj → V maj → IV maj → I maj	-	0.69%
	IV maj → I maj → bVII maj → IV maj	-	0.52%
6.	maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{maj2nd}}{\longrightarrow}$ maj	0.93%	1.11%
	V maj → I maj → IV maj → V maj	-	1.03%
7.	maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj	0.91%	1.09%
	V maj → I maj → IV maj → I maj	-	0.83%

Table 3. Beatles root interval and chord category rules (whose coverage is larger than 1%) and the associated degree and chord category rules. C_1 and C_2 represent the positive coverage over all the Beatles songs and over the Beatles tonal songs with no modulation respectively.

Rule	C
1. maj $\overset{\text{maj6th}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ dom	190 (0.80%)
2. dom $\overset{\text{perf4th}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ dom $\overset{\text{perf4th}}{\longrightarrow}$ maj	176 (0.74%)
3. min $\overset{\text{perf4th}}{\longrightarrow}$ dom $\overset{\text{perf4th}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ dom	174 (0.73%)
4. min $\overset{\text{perf4th}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ dom $\overset{\text{perf4th}}{\longrightarrow}$ maj	171 (0.72%)
5. min $\overset{\text{perf4th}}{\longrightarrow}$ dom $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{maj6th}}{\longrightarrow}$ min	170 (0.72%)
6. dom $\overset{\text{perfU}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ dom $\overset{\text{perf4th}}{\longrightarrow}$ maj	133 (0.56%)
7. maj $\overset{\text{maj2nd}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ dom $\overset{\text{perf4th}}{\longrightarrow}$ maj	126 (0.53%)
8. min $\overset{\text{perf4th}}{\longrightarrow}$ dom $\overset{\text{perf5th}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ dom	124 (0.52%)
9. min $\overset{\text{perfU}}{\longrightarrow}$ min $\overset{\text{perfU}}{\longrightarrow}$ min $\overset{\text{perfU}}{\longrightarrow}$ min	124 (0.52%)
10. dom $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{maj6th}}{\longrightarrow}$ min $\overset{\text{perf4th}}{\longrightarrow}$ min	121 (0.51%)

Table 4. Top ten Real Book harmony rules when considering root interval progressions and chord category progressions. C is the positive coverage.

most frequent chord category in the Beatles and the dominant chord ranks quite low in the chord category rules (rule 25 not shown here). For the Real Book, the rule with the highest coverage is $\bullet \overset{\text{perf4th}}{\longrightarrow} \bullet \overset{\text{perf4th}}{\longrightarrow} \bullet \overset{\text{perf4th}}{\longrightarrow} \bullet$. An interpretation of this rule is that for most of the instances it certainly is ii-V-I-IV (a very common jazz chord progression). Another common jazz chord progression, I-VI-II-V (often used as "turnaround" in jazz), is captured by rule 8 in Table 2. Moreover the dominant chord is the most frequent chord category in the Real Book which clearly distinguishes the jazz standards of the Real Book from the pop songs of the Beatles.

Note that due to the fact that the chord sequences overlap and due to the cyclic nature of some of the pop and jazz songs, many rules are not independent. For instance rules 2, 3, 6 and 7 in Table 1 can represent the same chord sequence maj-maj-maj-min repeated several times.

Moreover Aleph can also derive rules that contain some degree information. For that we constrain Aleph to derive rules about the intervals between the chord roots which we believe capture degree patterns. The top root interval and category rules for each corpus are presented in Tables 3 and 4. Furthermore, since we have key information for some of the Beatles songs we can actually obtain degree rules for them and an analysis of the degree rules allows us to match each root interval rule (with no tonal centre information) with the degree rules which are covered by it. The result of this matching process between degree and root interval rules is presented in Table 3 (top rules only). So for instance in Table 3 the instances of the root interval rule 5:

$$\text{maj} \overset{\text{perf5th}}{\longrightarrow} \text{maj} \overset{\text{min7th}}{\longrightarrow} \text{maj} \overset{\text{perf5th}}{\longrightarrow} \text{maj}$$

Rule	C
1. maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{maj2nd}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj	188 (1.38%)
2. maj $\overset{\text{maj2nd}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj	165 (1.21%)
3. maj $\overset{\text{perf5th}}{\longrightarrow}$ maj $\overset{\text{min7th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj	141 (1.04%)
4. maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{maj2nd}}{\longrightarrow}$ maj	126 (0.93%)
5. A maj $\rightarrow$ D maj $\rightarrow$ A maj $\rightarrow$ D maj	114 (0.84%)
6. maj$/_A \rightarrow$ maj$/_A \rightarrow$ maj$/_A \rightarrow$ maj$/_A$	110 (0.81%)
7. maj $\overset{\text{min7th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj	108 (0.79%)
8. maj $\overset{\text{perf5th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj $\overset{\text{min7th}}{\longrightarrow}$ maj	102 (0.75%)
9. maj $\overset{\text{perfU}}{\longrightarrow}$ maj $\overset{\text{perf4th}}{\longrightarrow}$ maj $\overset{\text{perf5th}}{\longrightarrow}$ maj	99 (0.73%)
10. D maj $\rightarrow$ G maj $\rightarrow$ D maj $\rightarrow$ G maj	92 (0.68%)

Table 5. Top ten Beatles harmony rules when the Real Book is taken as negative example. C is the positive coverage.

are for 54% of them instances of the degree rule:

$$\text{I maj} \rightarrow \text{V maj} \rightarrow \text{IV maj} \rightarrow \text{I maj}$$

and for 41%, instances of the degree rule:

$$\text{IV maj} \rightarrow \text{I maj} \rightarrow \text{bVII maj} \rightarrow \text{IV maj}$$

4.2 Characterisation of the Beatles vs. Real Book songs

For the second experiment we want to know the Beatles chord sequences that are not present in the Real Book. Aleph is provided with all the Beatles chord sequences of length 4 as positive examples and all the Real Book chord sequences of length 4 as negative examples. It returns 1679 rules which characterise all the chord sequences that only appear in the Beatles songs. The top ten rules are shown in Table 5. Some of these rules are correlated. For instance the 3 chord cyclic pattern I-IV-V-I-IV-V-I..., very common in the early compositions of the Beatles (see for instance the song *Please Please Me* of the album *Please Please Me*), is covered by rules 1, 2 and 4. Similarly the cyclic pattern I-V-IV-I-V-IV-I... is covered by rules 3, 7 and 8. Note also that the "back and forth" pattern between the first and fourth degree mentioned in [3] and identified in rule 1 of Table 3 appears in rules 5 and 10 of Table 5.

As in the previous experiment we also try to characterise the chord sequences in terms of root intervals and chord categories and obtain a set of 1520 rules which are not presented here.

4.3 Considerations about the size of the corpora and the computation time

Such an ILP approach has never been applied on such a scale: we dealt with data sets a musicologist would typically

be interested in studying (unified corpora of songs commonly accepted as representative of a composer/band/genre).

Although ILP systems are known to usually be resource intensive, the computation time of the ILP system was not a limiting factor in this work. Aleph computed all the rules in less than a minute on a regular desktop computer. We see our framework as a useful tool for musicologists since manual harmonic annotation and analysis of the whole Beatles corpus takes several years of musicological work whereas the automatic extraction of the chord progression patterns using ILP takes only seconds, allowing the user to concentrate on the interpretation of the results.

5 CONCLUSION AND FUTURE WORK

In this paper we presented a framework to automatically extract harmony rules from manually labelled chords using Inductive Logic Programing. We also gave an overview of the most common harmony rules extracted on the Beatles and Real Book songs using this framework. This first analysis shows that very common jazz and pop patterns are present in these rules. We hope that an in-depth musicological analysis of the other rules will reveal other less common and more specific harmonic patterns. We identified patterns listed in [3] but our methodology does more than listing and counting chord sequences, it also builds a minimal rule set which describes a data set. We believe this technique can be used by musicologists to automatically characterise the harmony of large sets of songs in few seconds.

Furthermore, since our system builds a sufficient model of a data set in future work we intend to test whether such logical rules can efficiently be used for classification and clustering purposes. We also plan to use this technique to characterise other musical phenomena such as rhythm, melody, structure from symbolic data. In order to deal with more data and to avoid the time consuming task of manual annotation of collections we intend to use automatic symbolic analysis systems such as Melisma [12]. A further step will be to adapt our ILP framework to audio data.

6 ACKNOWLEDGMENTS

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TIMBRE AND RHYTHMIC TRAP-TANDEM FEATURES FOR MUSIC INFORMATION RETRIEVAL

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ABSTRACT

The enormous growth of digital music databases has led to a comparable growth in the need for methods that help users organize and access such information. One area in particular that has seen much recent research activity is the use of automated techniques to describe audio content and to allow for its identification, browsing and retrieval. Conventional approaches to music content description rely on features characterizing the shape of the signal spectrum in relatively short-term frames. In the context of Automatic Speech Recognition (ASR), Hermansky [7] described an interesting alternative to short-term spectrum features, the TRAP-TANDEM approach which uses long-term band-limited features trained in a supervised fashion. We adapt this idea to the specific case of music signals and propose a generic system for the description of temporal patterns. The same system with different settings is able to extract features describing either timbre or rhythmic content. The quality of the generated features is demonstrated in a set of music retrieval experiments and compared to other state-of-the-art models.

1 INTRODUCTION

As discussed in [2], most state-of-the-art algorithms dedicated to the high-level description of music signals rely on the same basic architecture with different algorithm variants and parameters, i.e. short-term audio features extraction followed by some supervised or unsupervised machine learning algorithm. The most successful approaches rely on features describing the shape of short-term spectral frames of audio signal. Spectral shape is indeed known to be correlated with the perceived timbre of sounds as confirmed by recent experiments in isolated instrument recognition (see notably [8]). As a matter of fact, short-term spectral envelopes have dominated similar research fields for years, like e.g. ASR. These short-term spectral envelopes are typically transformed in accordance with some constraining properties of human hearing such as the nonlinear (critical-band like) frequency resolution (Bark, Mel), the compressive nonlinearity between acoustic stimulus and its percept (logarithm) or the decreasing sensitivity of hearing at lower frequencies (equal-loudness curves). Moreover, these modified spectral frames are typically projected on spectral basis that decorrelate the feature space (cepstrum).

In both speech and music signals, the smallest unit of information, i.e. phonemes on the one hand and notes on the other hand (not only pitched notes but also hits of percussive instruments or whatever that produces sounds), spread on longer time intervals than the usual short-term audio frame. Indeed, typical ASR/MIR (Music Information Retrieval) systems considers slices of audio signals of length 20 to 50 ms (slightly longer when accurate pitch estimates are needed in the lower frequencies). On the contrary, phonemes were demonstrated to spread at least over the interval 200-300ms [26]. As a matter of fact, the minimum discriminable inter onset interval (IOI) is estimated to lie within the range 50-100ms (i.e. two sounds separated by less than the minimum IOI will be perceived as one) so that it is likely that at least 50-100ms of information is needed by a human listener to interpret the incoming sound. Studies on rhythm perception show that the rate of information in music signals is even less. Experiments [16] have indicated that pulse sensation cease to exist outside of the period range of 200-2000ms which is known as the *region of pulse sensation* while the most natural foot-tapping period is approximately 500-600ms. Given these observations, it is reasonable to think that it is probably more perceptually relevant to model audio signals with a longer context than the usual 20-50ms spectrum frames. To preserve information related to the short-term dynamics of sounds and to keep sufficient time-resolution when detecting e.g. musical note onsets, a trade-off consists in building a model of the sequence of short-term feature vectors over a longer time-scale. This process is sometimes referred to as temporal feature integration [14].

The simplest approach consists in computing simple statistics of feature vectors (means and variances) over *texture-windows*. This has been shown [25] to significantly improve music genres classification accuracy when using windows of approximately 1.0 second as opposed to the direct use of short-term frames. However, simple statistics discard dynamical changes of short-term features while the dynamics of sound and notably the attack time and the fluctuations

of the spectral envelope over time have proved to be of a great importance in the perception of individual instrument notes (see [11]). Meng [14] modeled the evolution of features over a texture window with an auto-regressive (AR) model and got improved genre classification results than with simple statistics. McKinney and Breebart [12] computed a periodogram for each short-term feature dimension over a frame corresponding to roughly 768ms. Each periodogram was then summarized by its power in 4 predefined frequency bands using a fixed filter bank. This approach was pursued by Arenas-Garcia et al. [1] who trained the filter-bank in a supervised fashion to optimally suit a particular music organization task. Rauber et al. [20] used critical band energies periodograms with a much longer context, i.e. 6 seconds. This longer context was considered to model rhythmic patterns and the range 0-10Hz was considered as higher values are beyond what humans can perceive as rhythm (see again the region of pulse sensation). These approaches to temporal feature integration model the dynamics of each feature independently. Though Meng [14] describes a general multivariate autoregressive model that does take into account correlations between feature trajectories, for the sake of simplicity he experiments in practice with a diagonal multivariate autoregressive model, i.e. an AR model of each feature dimension. Pohle et al. [19] use independent component analysis on short sequences of critical band energies and obtain time-frequency 2D filters that are reminiscent of cortical receptive fields [4]. Though this approach seems more appropriate to take into account correlations between feature trajectories, it is at best of similar quality as short-term features in genre classification experiments.

As a matter of fact, the use of long-term features has been investigated more in depth in the context ASR, notably by Hermansky [7]. These features are extracted in 2 steps. Firstly, rather long-term TempoRAL Patterns (TRAPs) of band-limited (1-3 Bark) spectral energies are considered. Though, a context of 200-300ms seems needed for ASR, an even longer time interval of 1 second is considered so that information about slowly varying noise can be removed from the data (i.e. mean/standard deviation normalization). Hermansky [7] argues that, consistently with color separation in vision, it seems likely that the frequency selectivity of the hearing system is used for separating the reliable (high SNR) part of the signals from the unreliable ones, so that it seems reasonable to use independent frequency localized processors. Consequently, each band-limited TRAP is processed individually by a specifically trained system to build as much knowledge as possible into the feature extraction module to minimize the complexity of the subsequent stochastically trained models. The second step, referred to as the TANDEM part, consists in training a system aiming at the combination of frequency-localized evidence into a set of robust features that can be used in conventional HMM-based ASR systems.

To our knowledge, there's been only one application of these TRAP-TANDEM features to music signals in the context of drum transcription [17]. In this paper, we further investigate some possible applications of the TRAP-TANDEM approach in the context of music information retrieval. More specifically, we describe in section 2 our own implementation of the TRAP-TANDEM feature extraction module, which slightly differs from the original method. As a matter of fact, we propose two different implementations to focus on two different aspects of music signals, namely timbre and rhythm. In section 3, we evaluate the validity of these features for music information retrieval in a set of music clustering experiments. Section 4 reaches conclusion.

2 MUSICAL TRAP-TANDEM FEATURES

The first step of the processing consists in converting the audio signal into some time-frequency representation. In practice, we use the typical short-term Fourier transform (STFT) with Hann windowing of the short-term audio frames. The resulting short-term spectra are projected onto the Mel scale to simulate human critical bands of hearing [13]. The perceptual relevance of this time-frequency representation is further improved by exploiting masking properties of the human ear (see [22]) and frequency response of the outer and middle ear (see [23]). The loudness in Sone of the spectra is finally evaluated according to [3].

These short-term spectra will be later used to build timbre related TRAPs. Rhythmic TRAPs are based on a slightly different representation. More specifically, each critical band of the time-frequency plane is smoothed with a kernel, which width is taken in the range of the minimum IOI so that rhythmically irrelevant details of the band envelopes will be smoothed while the peaks corresponding to two different note onsets will not be merged. The first order derivative of each smoothed critical band is then taken to emphasize sudden changes. Critical bands are finally combined as suggested by Scheirer [21] who demonstrated that the perceived rhythmic content of many types of musical excerpts was mainly contained in 4 to 6 larger critical bands.

To reduce further processing, we deploy a note onset detection algorithm and we will only compute one TRAP feature vector per onsets instead of using a constant and faster rate of feature extraction. By synchronizing the analysis on detected onsets, an important factor of variability of the data is strongly reduced, i.e. the data is made translation invariant, and consequently, we can expect that the task of learning relevant features will be simplified. The differentiated signals computed for rhythm description are used as a basis to detect note onsets. The signals are first half-wave rectified and peaks are detected by using an adaptive threshold to account for possible loudness changes over the course of the musical piece. Peaks are first combined over the different

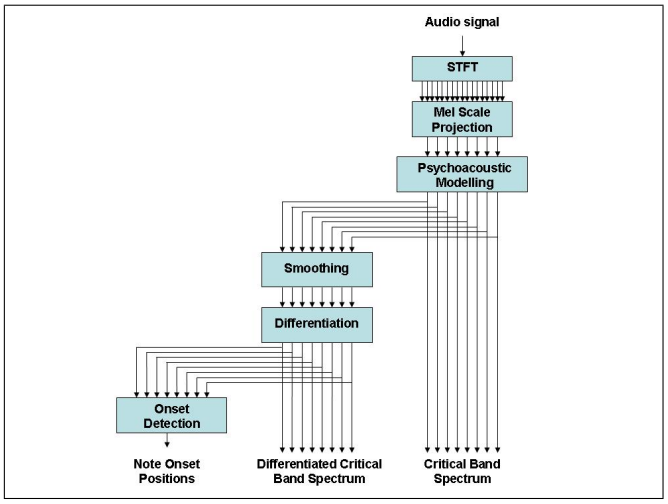

Figure 1. From audio signal to critical band spectra and onset positions.

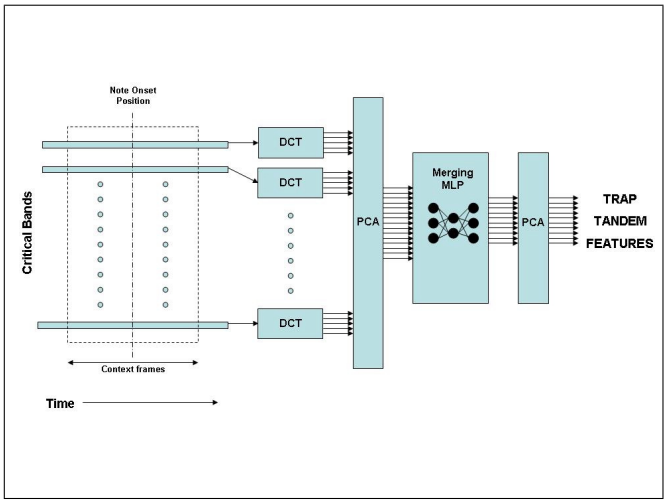

Figure 2. The TRAP-TANDEM feature extraction processing chain.

bands and those closer than the minimum IOI are merged together. Figure 1 illustrates the processing chain that goes from the audio signal to both critical band spectra and onset positions.

The data from each critical band and differentiated critical band surrounding each onset is then parameterized with the cosine transform, which has the good property of producing decorrelated signals. The cosine transform is simple to deploy since it does not need any training phase, plus it has the interesting property that it closely resembles the Principal Component Analysis (PCA) of such critical band signals [7]. The cosine transform also allows for a significant reduction of the dimensionality of the input data. The range of modulation frequencies of the cosines is carefully selected. For timbre description, modulations between 4 and up to 100 Hz can be considered. The lower limit of 4 Hz is set in accordance with the smallest perceivable sound unit discussed in section 1. The higher limit is in the range of modulations contributing to perceptual roughness, which is generally thought to be a primary component of musical dissonance [24]. As a matter of fact, the percept of roughness is considered to correlate with temporal envelope modulations in the range of about 20-150 Hz and maximal at 70 Hz. For rhythm description, lower frequencies are considered. Modulations between 1 and 4 Hz are interesting since they are of the order of typical beat rates, i.e. 60 to 240 Beats Per Minute (BPM).

In the original TRAP approach, for each critical band some algorithm, typically a non-linear feedforward multilayer perceptron (MLP), is trained to estimate posterior probabilities of the classes under consideration. In ASR, the targets are phonemes and there exist plenty of annotated datasets. Ideally, we would like to have instrument annotations to translate the TRAP idea to the case of music signals. Unfortunately, no such dataset exists for real-world polyphonic music and as a matter of fact, the annotation problem would become even more complex since we're considering mixture of instruments. For the time being, we left aside the use of critical band MLPs.

The TANDEM part of the system is in charge of combining evidence from the different frequency bands into a single estimate. A MLP is typically trained to combined these band limited features into a set of class posterior probabilities. Again, we lack appropriate annotated datasets. As an alternative, we use music genres annotations that are much cheaper to obtain, using e.g. some online music guide such as the AllMusic guide [1]. The merging MLP is fed with the concatenation of all band limited features, which are whitened with PCA to make them decorrelated. The MLP is trained to associate acoustic evidence with 50-dimensional binary vectors for which each dimension corresponds to a particular music genres. Notice that one song may be characterised by multiple genres. Once properly trained, the outputs of the MLP are decorrelated by PCA and can be used as a feature vector describing the different genres in which the acoustic data has been observed. Figure 2 illustrates the processing chain that goes from the critical band or differentiated critical band spectra to the final TRAP-TANDEM like features.

3 EVALUATIONS

The TRAP-TANDEM features describe the timbre and rhythmic context of a note onset. They may need to be aggregated into a song-level model in e.g. song retrieval applications. One application scenario consists in retrieving sets of songs

[1] http://www.allmusic.com

similar to some query song according to some similarity measure between song-level models. Though listening tests have proved to be a valid evaluation method of such music retrieval systems due to the consistency of judgements observed over different listeners [15], they are very demanding in terms of time and human resources. Previous works have shown that evaluations based on genre data correspond to evaluations based on similarity ratings gathered in listening tests [15]. Consequently, we will base our evaluation of the descriptive quality of our features on some genre annotated data. However, since we are more interested in music retrieval than automatic genre labelling, we will use measures of the ranking quality of the system rather than classification accuracy.

3.1 Timbre TRAP-TANDEM evaluation

To evaluate the timbre TRAP-TANDEM features, we have gathered a set of 210 songs annotated with genre and styles from the AllMusic guide. The quality of the labels was cross-checked by systematic listening tests. Each selected song is performed by a different artist to avoid the *album effect* [9] in the evaluation. Moreover, the songs selected were not previously used while training the TRAP-TANDEM feature extractor. Six main genre clusters of songs are considered and each genre cluster is composed of a set of smaller style clusters with 10 songs per style. The **Rock** cluster is composed of the *Grunge*, *British-Invasion*, *Punk-Blues*, *Glam-Rock*, *New-Wave*, *Folk-Rock* sub-clusters. The **Jazz** cluster is composed of the *Soul-Jazz*, *Swing*, *Hard-Bop*, *Free-Jazz* sub-clusters. The **Hip-Hop** cluster is composed of the *West-Coast*, *East-Coast*, *Turntablism*, *Old-School* sub-clusters. The **Electronica** cluster is composed of the *House*, *Trip-Hop*, *Drum'n'Bass* sub-clusters. The last two clusters, **Soul** and **Adult Contemporary**, are both divided into two sub-clusters according to the gender of the lead singer.

We will measure how well the timbre TRAP-TANDEM features are able to recover this organisation in terms of genre/style clusters. In practice, we summarize the distribution of TRAP-TANDEM feature vectors over each song by a simple average. Though more complex models, like e.g. HMM, could be deployed, our goal is to demonstrate the descriptive quality of the TRAP-TANDEM features so that we leave more complex song-level modelling strategies to future work. Two songs can then be compared by evaluating the cosine similarity of their average song-level TRAP-TANDEM vectors. The similarity between each pair of songs of the dataset is computed and the quality of the system is assessed by comparing the labels of a song and its nearest neighbours. More specifically, the precision at 1, 2 and 3 are computed for each query and averaged over the dataset. The precision at n is a quality measure commonly used to evaluate information retrieval systems. It accounts for the quality of ranking, i.e. it is high if the most relevant hits are in the top documents returned for a query. It is measured by computing the precision at different cut-off points (for example, if the top 10 documents are all relevant to the query and the next ten are all nonrelevant, we have 100% precision at a cut off of 10 documents but a 50% precision at a cut off of 20 documents).

Table 1. Timbre TRAP-TANDEM features.

	Prec. at 1	Prec. at 2	Prec. at 3
Genre	72.86	71.90	69.84
Style	50.00	43.10	39.37

Table 2. Full Gaussian of MFCCs.

	Prec. at 1	Prec. at 2	Prec. at 3
Genre	65.24	57.62	54.13
Style	35.71	30.00	25.87

To ease the comparison of our approach with the state-of-the-art, we have implemented one of the most popular timbre similarity measure based on spectral shape features. It simply consists of a Mel-Frequency Cepstrum Coefficients (MFCCs) frame-based parameterization of the audio signal (20 coefficients including the energy coefficient). These features are aggregated over the song as a Gaussian with full covariance matrix and compared using the symmetric version of the Kullback-Leibler (KL) divergence. This approach has been originally introduced by Mandel and Ellis [10] and was used in the winning algorithm of the 1st Annual Music Information Retrieval Evaluation exchange (MIREX 2005) [2] artist identification contest and ranked 3rd on 13 at the MIREX 2005 music genres classification contest while being almost 3 times faster than the first two winning algorithms. It can be considered as a simplified, yet competitive, implementation of Aucouturier's timbre model [2].

Tables 1 and 2 summarize the average precision at 1, 2 and 3 for both models. Results are given for both genres and styles targets, i.e. in the first case precision increases if the nearest neighbours are of the same genre, and in the second case precision increases if the nearest neighbours are of the same style.

It is clear from this experiment that the TRAP-TANDEM features are more reliable than the short-term spectral shape features in a music retrieval context. It is worth noticing that MFCCs with a simple average and cosine similarity leads to poorer results than MFCCs with mean, full covariance matrix and KL divergence, while for the TRAP-TANDEM features, the use of a KL-based distance function leads to slightly inferior results than the simpler cosine similarity.

[1] 1

[2] http://www.music-ir.org/mirex2005/index.php/Main_Page

Table 3. Rhythm TRAP-TANDEM features.

	Prec. at 1	Prec. at 2	Prec. at 3
Style	79.37	77.01	76.36

Table 4. 10-fold 1-NN classification accuracy.

	Accuracy
Rhythm TRAP-TANDEM features (without annotated tempo)	79.49
Gouyon & Dixon (without annotated tempo	67.60
Gouyon & Dixon (with annotated tempo)	82.10
Peeters (with annotated tempo)	90.40
Dixon et al. (with annotated tempo and semi-automated beat tracking)	96.00

This suggests that it is possible to have better retrieval results with a simpler—and especially faster—similarity function. Indeed, even if we're considering here 50-dimensional TRAP-TANDEM vectors against 20-dimensional MFCC vectors, the cosine similarity remains much faster than the symmetric KL of two full Gaussians since the later requires some matrix multiplications. This advantage becomes crucial when dealing with industrial databases with millions of songs.

3.2 Rhythm TRAP-TANDEM evaluation

We will evaluate the descriptive power of the rhythm TRAP-TANDEM features in a similar fashion. Rhythm TRAP-TANDEM features are averaged over each song and two songs are compared with the cosine similarity. We used here a well known dataset that contains 698 pieces of ballroom dance music divided into 8 sub-styles having different rhythmic characteristics, namely **Cha Cha Cha**, **Jive**, **Quickstep**, **Rumba**, **Samba**, **Tango**, **Viennese Waltz** and **Waltz**. It is interesting to notice that the TRAP-TANDEM system for rhythm was trained with the same 50 genres targets as the system for timbre, and that these genres are far from being as restricted as **Jive** or **Cha Cha Cha**. Table 3 summarizes the results obtained.

To ease the comparison with state-of-the-art algorithms, we also computed the classification accuracy on a 10-fold cross validation experiment with a 1-Nearest Neighbour classifier since various authors have reported results on this dataset using this particular evaluation procedure (see table 4). Gouyon and Dixon [6] reports up to 82.10% accuracy using a set of rhythmic features including the manually annotated tempo. The accuracy drops to 67.60% when using the tempo automatically extracted by their algorithm. Peeters [18] reaches up to 90.40% using *spectral rhythm patterns* normalised by the manually annotated tempo. While these two algorithms extract features from some periodicity representation of the audio signal, Dixon et al. [5] characterise the amplitude envelope of musical patterns, synchronised on musical bar positions and normalised to a reference tempo. They obtain an impressive 96.00% accuracy, but they also make use of the annotated tempo and a semi-automated beat tracking algorithm. On the contrary, our approach is fully automatic and reaches 79.49% classification accuracy. Moreover the results obtained by Dixon et al. with tempo normalised/bar synchronised temporal patterns suggest that the TRAP-TANDEM rhythmic features could become even more effective if synchronised on higher level musical events (musical bar positions instead of note onsets) and if made independent of the tempo. However, though on this particular dataset, a tempo normalisation may be needed since the clusters exhibit clearly defined rhythmic patterns with variable tempi, a tempo normalisation may not be so crucial for a general purpose music similarity engine since *slow/fast* songs should probably be similar to other *slow/fast* songs independently from the rhythmical pattern they're built on, i.e. the percept of speed would be more important than the perception of a particular rhythmical pattern.

4 CONCLUSION

We have presented a new set of features for music content description based on the work by Hermansky [7] in the context of ASR. The original approach has been adapted to the specific case of music signals and two different implementations based on the same architecture have been proposed to describe two apparently dissimilar dimensions of music, namely timbre and rhythmic patterns. Instead of using a relatively simple low-level characterization of the audio signal (like e.g. MFCCs), the TRAP-TANDEM approach is a complex feature extraction module that encodes temporal patterns and as much prior knowledge as possible. The distribution of TRAP-TANDEM features over a song can be described with simple models that can be used together with fast similarity measures. First experimental results confirm that the TRAP-TANDEM approach is competitive against state-of-the-art algorithms. Future work will focus on experimenting at a larger scale to confirm—or infirm—the descriptive quality of the TRAP-TANDEM features.

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USING THE SDIF SOUND DESCRIPTION INTERCHANGE FORMAT FOR AUDIO FEATURES

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ABSTRACT

We present a set of extensions to the Sound Description Interchange Format (SDIF) for the purpose of storage and/or transmission of general audio descriptors. The aim is to allow portability and interoperability between the feature extraction module of an audio information retrieval application and the remaining modules, such as training, classification or clustering. A set of techniques addressing the needs of short-time features and temporal modeling over longer windows are proposed, together with the mechanisms that allow further extensions or adaptations by the user. The paper is completed by an overview of the general aspects of SDIF and its practical use by means of a set of existing programming interfaces for, among others, C, C++ and Matlab.

1 INTRODUCTION

The *Sound Description Interchange Format* (SDIF) [1, 2] is an established standard for the precise, well-defined and extensible storage and interchange of a variety of sound descriptions including representations of the signal for analysis/synthesis like spectral, sinusoidal, time-domain, or higher-level models, audio descriptors like loudness or fundamental frequency, markers, labels, and statistical models. SDIF consists of a basic data format framework based on time-tagged frames containing matrices of binary numbers or text, and an extensible set of standard type declarations corresponding to different sound descriptions.

The SDIF standard was created in 1998 in collaboration between Ircam–Centre Pompidou in Paris, France, CNMAT at the University of Berkeley, USA, and the Music Technology Group (MTG) of the Universitat Pompeu Fabra, Barcelona, Spain. It arose out of the need to be able to store and exchange sound representation data between different analysis/synthesis programs, research teams, and institutions, and to enable anyone to interpret the information in the files correctly, needing as little external information as possible. With the previously widely used headerless ASCII formats, this goal was seriously compromised, as well as precision and space efficiency lost.

In the present contribution, we propose a set of extensions to the SDIF description types and conventions and best practices for the application of storing and/or transmitting a wide range of audio descriptors, e.g. as generated by the feature [1] extraction stage of an audio pattern recognition system. In the Audio or Music Information Retrieval fields, this is of interest to assure interoperability between feature extraction and further processing modules, such as feature selection, training, clustering or classification, possibly being developed independently from each other, or even by different institutions. Such a need for standardization can arise in the context of multi-partner evaluation, such as in the Music Information Retrieval Evaluation eXchange (MIREX) [3].

SDIF is an open format, unhampered by licensing or intellectual property restrictions, made by the research community for research and creation. There is a wide range of software available for its use: Libraries in C, C++, Java, Matlab, Perl, and Python, command-line and graphical tools, and plugins for audio-processing systems such as Max/MSP and PureData.

After an introduction to the basics of SDIF (section 2), and a general overview of common different types of audio descriptors (section 3), we will propose several format conventions for storing general descriptor information (section 4). The standard's type declarations are completely extensible and modifiable by the user (section 5), which makes the current proposal adaptable to special application needs. Finally, in section 7, we will present a selection of helpful tools distributed with the SDIF library, programming interfaces for reading and writing SDIF files, a number of applications and online resources.

2 OVERVIEW OF THE SDIF SPECIFICATION

SDIF is a binary meta-format based on streams, frames and matrices. The matrix content representation can be chosen to be text, integer, or floating point in different bit-widths. Several matrices are grouped into a frame, which has a precise 64 bit floating point time tag in seconds relative to the start of the file. Frames are always stored in increasing temporal order and are assigned to a stream. Streams separate different entities of the same type, such as different sound files, channels, or different analyses or descriptors, thus allowing to unite a database of several source sound files, plus various stages of analysis and descriptor data, in one single SDIF file. [2] For example, one could choose to store two channels of audio, their STFT representation, fundamental frequency and sinusoidal partial representation, markers, labels, and harmonicity audio descriptors in one file.

[1] We will use the terms *feature* and *descriptor* interchangeably.

[2] This was formerly achieved by separate files with different file extensions, grouped in the same directory, with the risk of loss of single files, and a certain implicitness of their relationships.

Frame and matrix types are identified by 4-byte ASCII *signatures*, inspired by the IFF format. Each frame starts with a header containing the frame signature, the number of matrices it contains, an integer identifying the stream it belongs to, and the time tag. Each matrix' header contains its signature, its content data type, and its number of rows and columns. Matrices are padded to 8 byte boundaries to achieve 64 bit word alignment for fast access using memory mapped files.

Special header-frames can store global textual meta-data in so-called Name–Value Tables (NVTs), together with relationships of different streams (see section 6). However, global information that is needed to interpret the data, for instance the sampling rate, is stored in information matrices, making SDIF files streamable.

There are a number of standard description types that specify the columns of matrices, and which matrices can occur in which frame type. Examples include `1FQ0` to store fundamental frequencies and `1TRC` to store partial tracks resulting from sinusoidal analysis. By convention, the signatures of standard frame and matrix signatures start with a version number, mostly `1`. It is always possible to declare new experimental description types, or to extend existing types by new matrices for a frame, or new columns for a matrix. New experimental types have signatures starting with `X`. SDIF is designed such that programs can always ignore additional data they don't know how to interpret.

3 OVERVIEW OF AUDIO DESCRIPTORS

Audio descriptors (also named audio features or audio attributes) are numerical values describing the contents of an audio signal according to various points of view, e.g. temporal, spectral or perceptual characteristics. They are used nowadays in many areas such as for the development of query-by-example (search-by-similarity), automatic indexing, segmentation or identification applications. These features are extracted from the audio signal using signal processing algorithms. Several taxonomies can be used to classify them, for example according to the type of content they can be applied to (single musical notes, percussion, sound effects, generic audio, etc.), the underlying signal model they rely on (source-filter, sinusoidal harmonic models, etc.), their abstractness, i.e. what the audio feature represents (spectral envelope, temporal envelope, etc.), their steadiness or variability, i.e., the fact that they represent a value extracted from the signal at a given time, or a parameter from a model of the signal behavior along time (mean, standard deviation, derivative or Markov model of a parameter), or the time extent of the description provided by them: some descriptors apply to only part of the object (e.g., description of the attack of the sound) whereas others apply to the whole signal (e.g., loudness). On the computer side, an audio feature is represented by a numerical value or a vector of numerical values when several audio features can be linked together (such as the coefficients of the MFCC, or various definitions of the spectral centroid).

Each of such numerical values describes a specific property of the signal around a specific time (the time of the segment corresponding to the analysis window). Most audio features are first computed on a small-time scale (often called short-term analysis) which for most of them correspond to the length of the analysis window used for FFT analysis (usually 40 ms to 80 ms). Other features are computed on a longer-time scale due to their semantics (the log-attack-time, the roughness or the fluctuation strength are associated to a whole musical note) or due to computational requirements (the estimation of a vibrato or tremolo at a frequency of 6 Hz necessitates a window larger than 80 ms). In this case the numerical value needs to be associated exactly to the segment used to compute it. Finally, it has become frequent to model the behavior of an audio feature over time. Using long-term analysis (often with window lengths of 500 ms or larger), it is common to compute the mean, variance or derivative values of the feature over the sliding window. This type of description is often named *temporal modeling*, and the corresponding large frames are sometimes called *texture windows* or *macroframes*. In this case, the segment over which the temporal model is computed does not need to have a specific semantic (e.g., "every 200 ms" has no meaning by itself), although it can have (for example for beat-synchronous analysis).

In the proposed SDIF extensions, we address these specific requirements concerning definition of different time horizons, multidimensionality and addition of semantic information. It is out of the scope of this article to present detailed mathematical definitions of the descriptors. A large set of them can be found in [4].

4 EXTENDED TYPES FOR GENERALIZED AUDIO DESCRIPTORS

A defining aspect of SDIF is its flexibility regarding data types. Using the data declaration syntax introduced later in section 5, the user is able to define or redefine the frame/matrix grouping rules and to add or modify type declarations at will. Throughout the paper, we will present different possibilities for different applications in the context of descriptor extraction. At the same time, however, we are proposing a ready-to-use set of type declarations designed according to what we consider appropriate to most audio content analysis and retrieval applications. The proposed type declarations, together with related documentation and several example SDIF files can be accessed online [3] .

In both short-time and temporal modeling cases, the values of a given descriptor at a given time are stored in a separate matrix. For example, spectral centroid values are stored in matrices of signature `1SCN` and for zero-crossing values the matrix signature is `1ZCR`. In analogy with the standard matrix types (e.g., with sinusoidal modeling), the number of rows of the matrix must correspond to the dimensionality of the descriptor, i.e., with the number of frequency bands (such as with spectral flatness) or with the number of coefficients (such as with MFCCs or autocorrelation features).

[3] http://sdif.sourceforge.net/descriptor-types

The columns are intended to store the values for different definitions or implementations of the same descriptor. For example, the first column of the spectral centroid could contain the values for a centroid defined on linear frequency and amplitude scales, and the second the values using logarithmic scalings. We will thus use *dimensions* to denote rows and *variations* to denote columns.

Both the number of rows and of columns are not fixed beforehand by the type declaration. In most cases, the number of rows will depend on a user-defined feature extraction parameter, such as number of frequency bands or cepstral coefficients. It is also most likely that the user will just store a single variation for each descriptor. In some training experiments, however, it can be interesting to store different variations at the same time and let an automatic feature selection algorithm select the most informative one.

Even if not strictly necessary for a clustering or statistical training algorithm, the feature extraction program should store additional information alongside the descriptor values, such as bandwidths, thresholds or labels describing the contents of the columns. This data will be included in information matrices, whose signatures will be equal to the descriptor matrix signatures, with the first character `1` replaced by `I`. Entries on information matrices are always stored column-wise. The next subsections will detail how to store such matrices.

To accommodate the different descriptor time-spans mentioned above (short-, mid-, long-term and temporal modeling) we propose two different schemes for SDIF storage. In fact, we simplify such categorization by only considering, on the one hand, the descriptors themselves (which are directly based on the signal) and, on the other hand, temporal modelings (which are based on descriptors), which will respectively be addressed in the next two subsections. We deliberately avoid a categorical distinction between the length of the time spans, since the only difference from the SDIF point of view is the definition of the time boundaries. Also, we will avoid the terminology "low-level" or "high-level" feature because of the lack of consensus about its meaning. By convention, an SDIF file containing only audio descriptors should end with the double extension `.descr.sdif`.

4.1 Format for Descriptors

We propose the following procedure to store descriptors that are directly based on the signal, independently of their time span. Each data matrix corresponding to a given analysis window of a given descriptor is contained within a frame of type `1DSC`, a *descriptor frame*. In order to be able to perform a one-to-one mapping between streams and descriptors, each descriptor frame must contain only a single descriptor data matrix. The time tag on each frame corresponds to the center of the analysis window. The optional information matrices must be stored at the beginning of the corresponding descriptor stream, and are valid until a new information matrix of the same type appears on the same stream. Figure 1 represents this approach schematically for the case of a file storing spectral centroid (`1SCN`) and perceptual tristimulus (`1PTR`) data. In this example, the spectral centroid is a unidimensional descriptor with a single variation, and two different variations of the three-dimensional tristimulus have been stored, each one defined with a different amplitude scaling.

The SDIF reading routines search first by frames, and then by matrices. Thus, if the performance in accessing individual descriptors is critical, an alternative approach would be to declare a pair of frame type and matrix type with the same signature for each descriptor. Its disadvantage however is that it requires to double the amount of type declarations.

4.2 Format for Temporal Modeling of Descriptors

There are three main reasons for considering temporal modeling of descriptors as a special case that requires dedicated storing conventions. First, they are *derived* features of typically the same dimensionality than their shorter-term counterparts they are based on, and thus they can reuse their matrix type declarations. This results in each temporal model being associated to a frame type, containing matrices of the types already declared for the descriptors. For example, a texture window of the weighted variance of the spectral centroid will correspond to a frame of type `1WVR` containing a matrix of type `1SCN`. An indicative list of types of temporal modeling is: mean, variance, standard deviation, weighted mean, weighted standard deviation, amplitude modulation.

Second, it is possible to assume that their time-span will be relatively long-term (in the range of seconds), and thus performance when accessing them will rarely be critical. If stored together with their short-term descriptors, they will predictably constitute a very small percentage of the file size. This allows more freedom in choosing the most appropriate way of frame and matrix grouping, depending on the application. This can be done either in the one-matrix-per-frame way, as before, or using *compound frames*, i.e. grouping several matrices corresponding to different descriptors under the same frame. In the first case, individual assignment to streams, and different window sizes between different descriptors will be possible. In the compound case, visual inspection of SDIF files (e.g., after ASCII conversion with the **sdiftotext** tool) will be clearer. To leave both possibilities open, the provided type declaration lists all descriptor matrices as optional members of each temporal modeling frame. If visual inspection is not important, we recommend using the one-matrix-per-frame alternative.

Finally, the third reason is that the segment the temporal modeling works on is not necessarily of constant size and overlap. For example, one can define a sequence of log-attack time features resulting from a note-wise segmentation of a melodic line. Thus, a mechanism must be provided to indicate the segment on which the descriptor evolution is modeled. This mechanism makes use of the standard frame type `1MRK` that represents a marker at the time of the frame, containing matrices `1BEG` and/or `1END` with a unique identifier of a segment. Now, the temporal modeling frame will also contain a matrix with the identifier of the segment whose descriptor evolution it models.

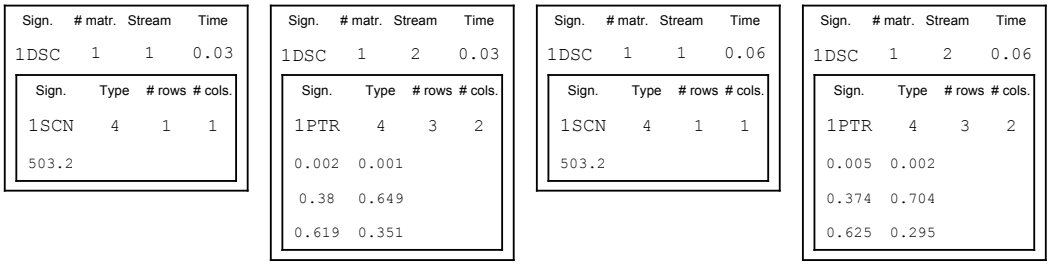

Figure 1. SDIF storage example for four consecutive frames of a short-time spectral centroid (`1SCN`) and a short-time perceptual tristimulus (`1PTR`).

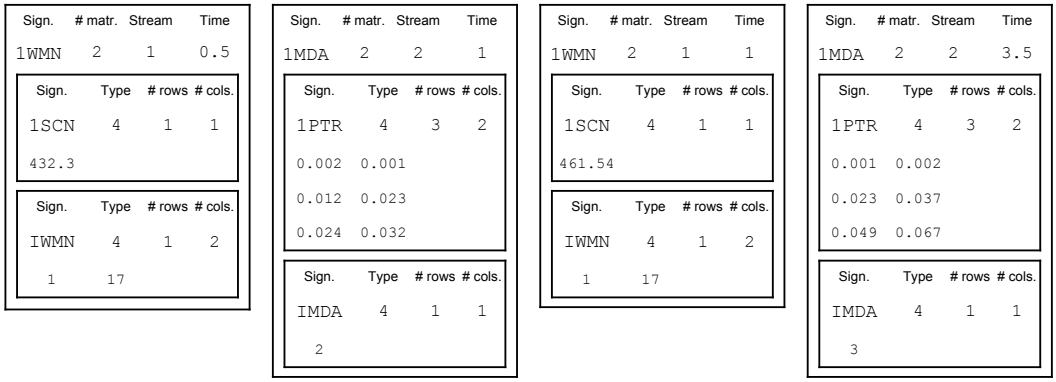

Figure 2. SDIF storage example for four consecutive temporal modeling segments: weighted mean (`1WMN`) of spectral centroid and modulation amplitude (`1MDA`) of perceptual tristimulus.

Each temporal modeling frame type has an associated information matrix type, e.g. `IWVR` for the weighted variance frame type `1WVR`. Then, each temporal modeling frame contains one information matrix indicating the duration, in seconds, of the window as its first element. For regularly spaced texture windows, the time tag of the frames corresponds to the middle of the temporal modeling segment. For irregular segments given by markers, the temporal modeling frame is written at the start time of the segment. Figure 2 shows an example of this approach, using single-matrix grouping. It shows the storage for two consecutive texture windows containing the weighted mean (`1WMN`) of the spectral centroid, and the amplitude modulation (`1MDA`) of the perceptual tristimulus. Like with any other matrix, the meaning of the columns is contained in the type declaration. As in the plain descriptors case, it is possible to store descriptor-related information matrices at the beginning of the corresponding streams. It should be noted that some temporal modelings, such as log-attack time or temporal increase, are almost always based on a very specific descriptor, in this case the energy envelope. It would be however possible to attach them to no matter which shorter-time feature (for instance, the temporal increase of the fundamental frequency could be an informative "sweep descriptor").

5 USER-DEFINED TYPES

SDIF allows to define new description types by declaring frame and matrix signatures, and matrix columns, or to extend existing types by new matrices, or new columns. The declaration of extended types is included in a special *type declaration* frame of signature `1TYP`. Each SDIF file must include a `1TYP` frame declaring the non-standard types, and is therefore self-contained. The standard types must not be declared in such a frame, since they are already recognized by the library. The following is an example of the type declaration for some existing types:

```
1TYP
{
  1MTD 1SCN    {SpectralCentroid}
  1MTD 1ZCR    {ZeroCrossingRate}
  1MTD IWMN    {WindowLength}
  1FTD 1WMN
       {
         1SCN  SpectralCentroidWeightedMean;
         1ZCR  SignalZeroCrossingRateWeightedMean;
         1FQ0  FundamentalFrequencyWeightedMean;
         IWMN  WeightedMeanInfo;
       }
}
```

Each line starting with `1MTD` is a matrix type declaration, containing the new matrix signature and, between curly

brackets, a list of the declared column names, separated by commas. The lines starting with `1FTD` are frame type declarations; after the frame signature (`1WMN` in the example), between the curly brackets, there is the list of the matrices that can appear inside that frame with their role, terminated by a semicolon. A matrix type must be declared before the frame type containing it. As a side note, we point out that even for standard types (like `1FQ0` in the example), the association with a non-standard frame *must* be created. All the descriptor types proposed here contain only one declared column. This does not hinder the user to store as many additional columns as needed.

6 DECLARATION OF STREAM RELATIONSHIPS

In the previous section we have seen that the proposed SDIF representation of audio descriptors will distribute the data over several streams of SDIF frames. For instance, we could have a stream of temporal modeling frames `1WMN`, modeling segments given by a stream of `1MRK` frames, based on various descriptors in different streams, that were extracted from the audio data in a stream of `1TDS` files.

While the relationships between these streams are clear in our context of use, we would like them to be expressed explicitly, in a machine-readable format, in order to be exploited by tools, visualisers, and other applications. The explicitness of the relationships and meanings of streams becomes even more important when several source sounds, channels, or analyses are included in the same SDIF file.

At the beginning of the SDIF standard, there has been an attempt [5] to formalise the relationships between streams in an XML based language, which has not been followed by concrete implementations. This is possibly due to the comparatively simple SDIF files that were used at that time, posing no real need for an explicit representation, and, what's more, to the problem of adding a dependency on an external library to parse XML to all SDIF tools.

We propose here a new, simple approach how to define relationships, meaning, and content of streams in a tabular text format, not unlike the existing Name–Value Tables. We will start with expressing only those relationships that are indeed needed for the applications described in this article, but we strive to keep the format open enough for other relationships to be added in the future.

The format associates one entity by a relationship to a list of entities. The possible entities are listed in table 1, the list of defined relationships in table 2. For stream IDs and frame and matrix signatures, the notation of entities follows the SDIF-selection syntax [2] to ease understanding and parsing, all other entities are labels, starting with a letter. We propose to store this table in a `2IDS` frame with text matrix.

Entity	Example	Description
# *stream id*	`#23`	Stream number 23
: *frame* / *matrix*	`:1WMN/1SCN`	stream contains weighted mean of spectral centroid
identifier	left-channel	Label

Table 1. List of entities in the proposed stream relationship declaration.

Relationship	Description
contains	left stream contains frame, matrix types given right
group	left label groups labels or streams to the right
segments	left stream contains segmentation information for right entity
derives	left entity is derived (by analysis or temporal modeling) from right entity

Table 2. List of relationships between two entities.

7 SDIF USAGE INFORMATION

7.1 SDIF Tools

In order to easily manipulate the produced SDIF files, the library provides some command-line tools together with the source code. Basically, there are tools to display, to extract and to convert the stored data; the most important are:

• **querysdif**: prints out a summary of the information stored in the file, e.g. the number and the types of stored frames and matrices,

• **sdifextract**: extracts selected data from a file; the output can be stored in many formats like SDIF itself, multi-bpf (text lines of frame time and all selected columns), etc.,

• **sdiftotext, tosdif**: convert data between SDIF and plain text, preserving frame-matrix relationships. These also exist as *droplets* for Macintosh, onto which one can drop a file, generating the complementary text or SDIF file.

7.2 Programming Interfaces

This section will describe the existing possibilities to manipulate SDIF data using the SDIF libraries that are publicly available on the SourceForge software repository (see location below). There exist a C-library (denoted just as the SDIF library) and a C++ library (called Easdif). Easdif incorporates the complete interface of the SDIF library. Additionally Easdif provides a high level programming interface to SDIF files, frames, and matrices including iterators over SDIF files. These iterators allow easy and efficient navigation within the SDIF file using an internal cache of the meta data (frame and matrix headers). The SDIF and Easdif libraries are designed to work on all major platforms (Linux, Mac OS X and Windows), and thanks to the use of the cmake [4] project generator they can be compiled with most common compilers (gcc, MSVC, Intel, . . .).

The Easdif library cannot only be used in C++ applications, it comes as well with a number of bindings for other applications and languages. A recent achievement is the ad-

[4] http://www.cmake.org

dition of sources for Matlab and Octave bindings (mex) that allow access to data stored in SDIF files for these two programing environments. The Matlab/Octave interface makes use of the frame and matrix header cache of Easdif to allow efficient random access to SDIF frames and matrices.

Additionally, the Easdif library includes support for the SWIG [5] wrapper generator and interface descriptions for bindings for python, perl and java. Starting from the existing interface descriptions the addition of other scripting languages that are supported by SWIG should be easy.

7.3 Applications using SDIF

Many applications developed nowadays integrate a general-purpose SDIF reader/writer. Here is a partial list:

• **AudioSculpt** [6] : tool to analyse and modify sounds through a visual approach,
• **ASAnnotation** [7] : sound analysis, visualisation and annotation, based on AudioSculpt,
• **CLAM** [8] : C++ library for music information retrieval,
• **Csound** [9] : language for sound manipulation and analysis,
• **Diphone Studio** [10] : tool for sonic morphing,
• **FTM** [11] for Max/MSP, jMax, PureData: realtime visual programming environment for sound analysis/synthesis,
• **Loris** [12] : sound modeling and processing package,
• **OpenMusic** [13] : visual programming language for computer-aided composition,
• **Open Sound World** [14] : programming environment to process sound in real-time,
• **SDIF-Edit** [15] : SDIF editor and visualisation tool,
• **SPEAR** [16] : sinudoidal analysis/resynthesis.

Moreover, in the developer tools included with the latest version of the Mac OS X operating system (version 10.5, *Leopard*), SDIF support has been included for the AudioUnit called *AdditiveSynth*.

7.4 Availability and Online Resources

SDIF is an open source project hosted on the SourceForge repository. The following resources are available:

• **SDIF Homepage.** (http://sdif.sourceforge.net)
Includes the format specification and main documents, and pointers to all other resources.
• **SDIF Download.** (http://sourceforge.net/projects/sdif)
Includes the sources and several binaries for the SDIF and Easdif libraries. For developers, CVS access is possible.
• **SDIF Wiki.** (http://sdif.wiki.sourceforge.net)
is intended for the documentation of application-specific usages, the proposal of extensions, and additional user-oriented documentation.
• **SDIF mailing lists.** There is one mailing list for users (http://listes.ircam.fr/wws/info/sdif) and one for developers (https://lists.sourceforge.net/lists/listinfo/sdif-devel).

[5] http://www.swig.org
[6] http://forumnet.ircam.fr/691.html
[7] http://www.ircam.fr/anasyn/ASAnnotation
[8] http://clam.iua.upf.edu
[9] http://www.csounds.com
[10] http://forumnet.ircam.fr/703.html
[11] http://ftm.ircam.fr
[12] http://www.cerlsoundgroup.org/Loris/
[13] http://recherche.ircam.fr/equipes/repmus/OpenMusic/
[14] http://osw.sourceforge.net
[15] http://recherche.ircam.fr/equipes/repmus/bresson/sdifedit/sdifedit.html
[16] http://www.klingbeil.com/spear/

8 CONCLUSION

In the context of the growing needs for standardization within the fields of audio-based content analysis and retrieval, we have proposed the use of the well-established and open SDIF format for storing, transmitting and sharing general audio features. We made use of SDIF's extension capabilities to declare a set of extended types addressing the needs of general short-term and temporal modeling descriptors. We are hoping to initiate a discussion among interested researchers, e.g. via the wiki and the mailing lists, to further assess the general needs, and eventually update the type proposals. Concerning future work, we are studying the use of SDIF for other audio-related pattern recognition data, such as feature transformation data and statistical models.

9 ACKNOWLEDGMENTS

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NON-NEGATIVE MATRIX DIVISION FOR THE AUTOMATIC TRANSCRIPTION OF POLYPHONIC MUSIC

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ABSTRACT

In this paper we present a new method in the style of non-negative matrix factorization for automatic transcription of polyphonic music played by a single instrument (e.g., a piano). We suggest using a fixed repository of base vectors corresponding to tone models of single pitches played on a certain instrument. This assumption turns the blind factorization into a kind of non-negative matrix division for which an algorithm is presented. The same algorithm can be applied for learning the model dictionary from sample tones as well. This method is biased towards the instrument used during the training phase. But this is admissible in applications like performance analysis of solo music. The proposed approach is tested on a Mozart sonata where a symbolic representation is available as well as the recording on a computer controlled grand piano.

1 INTRODUCTION

Transcription of polyphonic music is a difficult task even for humans after several years of musical training. In the computational field people have been working on the problem of extracting single note events from complex music recordings for more than three decades. In special cases like monophonic music some systems have proven to be successful [1]. But pieces where more than one note is present at a time are much more challenging. Consonant tones in Western music often have frequency relations close to simple integer ratios and therefore cause overlapping harmonics. So when considering a power spectrum the mapping of found energies to certain fundamental frequencies is usually ambiguous.

A review of transcription methods is given in [4] and [5], clustering them into three main approaches. The first systems were built on pure bottom-up principles without considering any higher level knowledge. Although these algorithms used to fit very specific cases only, recent works like [6] or [13] show that bottom-up methods have overcome those early restrictions. A second group of transcription methods, like used by [12], is based on blackboard systems. Here low-level information gathered by digital signal processing and frame-wise description of the auditory scene as well as high-level prior knowledge is used to support or discard hypotheses at multiple levels.

The third major approach to music transcription is made up of model based algorithms. Similar to blackboard systems they also include high-level information as well as low-level signal based features. The difference is that prior knowledge is fed into the system by introducing a model of the analyzed data. The signal is then processed in order to estimate the model's parameters. The results of these methods can only be as good as the assumed model fits the actual data. Works like [14] or [2] are examples of this class.

During the last years the methods of non-negative matrix factorization (NMF) [10], independent component analysis (ICA) [9] and sparse coding [13] became of increasing interest in audio analysis. The basic idea is the introduction of hidden, not directly observable atoms. The above cited methods decompose an input matrix into two factors where one is a dictionary describing the individual atoms and the other gives the activation of these components as a function of time. The non-negativity constraint is derived from the areas of application where single observations linearly add up to the whole. For instance in audio analysis it would not make sense to consider notes with negative loudness. Since the only prior knowledge is the maximum number of independent components these algorithms are part of the group of bottom-up methods as described above.

In this paper we propose a new method where the ideas of matrix factorization and sparse, independent components are adapted to follow a model based approach. Studies on the human approach to music transcription [5] have shown that trained musicians use lots of background information like the style of the piece or the instruments playing. They expect certain timbres and therefore would for example never search for distorted guitar tones in a classical piano piece. Applying this principle to the non-negative matrix factorization, prior knowledge about the dictionary is incorporated. The activation matrix will then remain the only

unknown component which needs an operation like a non-negative matrix division in order to be calculated.

Section 2 of this paper focuses on the NMF and its shortcomings in the context of transcription of solo music as a motivation for our method that is then explained in detail. Section 3 describes how the tone models representing a certain instrument can be extracted from sample recordings. The post processing step transforming the activation patterns yielded by the matrix division into discrete note events is described in section 4. Sections on the experimentation results and our conclusions complete the paper.

2 PITCH DECOMPOSITION

2.1 Non-negative matrix factorization

Non-negative matrix factorization as introduced in [9] decomposes a non-negative input V of size $m \times n$ into two non-negative output matrices W and H of size $m \times r$ and $r \times n$ respectively, such that

$$V \approx W \cdot H \tag{1}$$

where by convention W is regarded as a set of basis vectors and H as the aggregation of their activation patterns.

Since perfect factorization is not possible in almost all cases, a solution to equation (1) with minimal error of reconstruction is achieved by minimizing a cost function over the difference between V and $W \cdot H$. Common such functions are the Euclidean distance $E(V, WH)$ or the Kullback-Leibler divergence $D(V \parallel WH)$.

Applied to a power spectrum, as obtained by the short time Fourier transform, the basis components in W are weighted frequency groups that are found to sound together. Ideally they belong either to a single pitch played on a certain instrument or a group of pitches that are normally played together like the notes of a chord. If the number r of basis components is smaller than the number of different pitches played, some of the pitches have to be either omitted or grouped within one atom. In the reverse case where r is sufficiently large there can be atoms representing noise, or there is more than one atom per one single pitch. Here it is very likely that a component represents the sustained part of a note whereas another maps to the note onset with much richer harmonics. A detailed investigation on these effects can be found in [13].

The component activation data in H contains the strength of each atom at a certain time frame. Due to the non-negativity constraint the combination is additive only. This gives consideration to the fact that there is nothing like negative loudness of notes.

Effective algorithms for the calculation of the NNMF have been introduced in [10]. Multiplicative update rules are used in order to find local minima starting form randomly initialized matrices W and H. Using the Kullback-Leibler divergence $D(V \parallel WH)$ as cost function these update rules are

$$H_{a\mu} \leftarrow H_{a\mu} \frac{\sum_i W_{ia} V_{i\mu} / (WH)_{i\mu}}{\sum_k W_{ka}} \tag{2}$$

$$W_{ia} \leftarrow W_{ia} \frac{\sum_\mu H_{a\mu} V_{i\mu} / (WH)_{i\mu}}{\sum_\nu H_{a\nu}} \tag{3}$$

In [10] a proof for the convergence of this algorithm towards a local minimum is given. It is shown that the divergence is (i) non-increasing under above update rules and (ii) invariant if and only if W as well as H are at stationary points.

2.2 Drawbacks of NNMF

Several works like [13] or [15] have concentrated on applying matrix factorization using non-negativity and sparseness constraints to automatic music transcription. Although there are numerous advantages, an inherent problem of NMF based approaches is the determination of an appropriate number r of base components. This parameter has to be guessed since the number of different pitches present in a piece of music is not known in advance. An r that is too small cannot represent each pitch individually whereas too large values cause increased computational expenses as well as difficulties when mapping base vectors in W to transcribed pitches.

Another drawback is that there is no guarantee that each played pitch is represented at all. [13] reports that although using a more than sufficiently large number of base vectors in an NMF as well as in two sparse coding approaches a few notes are not represented. This is the case when chords or certain residual noise patterns become more significant than single tones that are played only very rarely.

Thirdly learning a dictionary of independent components while transcribing music played by a single instrument does not seem to be a natural way of approaching the problem. As shown in [5] humans start by detecting the genre and style of a piece, which allows them to limit the number of possible instruments and timbres to be expected. Learning the dictionary of independent components along with their activation is, as pointed out above, likely to model noise as well and therefore prone to overfitting. Restricting dictionary vectors to feasible values in advance is a reasonable means of preventing overfitting as well as unnecessary computational costs.

2.3 Non-negative matrix division

To overcome the above drawbacks we propose fixing the number r of independent components to the number of actual possibilities regarding the pitch range of the instrument in focus, and the single atoms of the dictionary W to stereotypical tone models of the corresponding pitches. One approach would be to use multiplicative updates on a random initialization of H applying the rule in (2) while omitting (3) or defining constraints on its computation like done in [16]. But exploiting the fact that W is fixed and therefore single vectors of V can be processed independently, equation (1) resolves to

$$v \approx \overline{W} \cdot h \tag{4}$$

where $\overline{W}$ is the fixed dictionary. v and h are column vectors representing one time frame of the spectrogram and the pitch activation respectively. In order to measure the quality of an approximation in (4) again a cost function is needed. A convenient measure is the mean square criterion where

$$f = \frac{1}{2} \parallel \overline{W}h - v \parallel^2 \tag{5}$$

has to be minimized while regarding the constraint of non-negativity. According to [8] this problem is solved by an iterative algorithm as follows.

1. Initialize all elements of h to zero and introduce two sets P and Z where P is empty and Z contains all indices within h.
2. Compute the gradient $\bigtriangledown_h f = \overline{W}^T \cdot (v - \overline{W}h)$ where f is the cost function as defined in (5).
3. If $Z = \{\}$ or $\forall i : i \in Z \Rightarrow (\bigtriangledown_h f)_i \leq 0$ then terminate.
4. Find the maximum element of $\bigtriangledown_h f$ and move its index from Z to P.
5. Solve the unconstrained linear least squares problem $\overline{W}_{sub} \cdot z = v$ where $\overline{W}_{sub}$ is a copy of $\overline{W}$ where all columns corresponding to indices in Z are set to zeros. Within the result z only those elements with indices contained in P are significant. The others are set to zero.
6. If $\forall i : i \in P \Rightarrow z_i \geq 0$ then z is a feasible solution to the subproblem, h is set to z and the main loop is continued at step 2.
7. If the above condition does not hold z can only contribute to the new temporary solution up to a certain amount. Therefore a factor α (a learning rate) is calculated as $\alpha = argmin_i(h_i/(h_i - z_i))$ where only the indices of negative elements in z are allowed as i.
8. Calculate the new temporary solution using α from the above step as $h = h + \alpha(z - h)$
9. Move all indices for which the corresponding element in h is zero from P to Z. Continue working on the subproblem at step 5.

Although the result of one frame is a useful hint for the computation of the next frame, single time frames can now be independently processed. This makes the method suitable for parallelization as well as online processing.

Reassembling the results of individual frames gives a complete activation matrix like H from equation (1) as an optimal non-negative quotient of an input power spectrogram V and a given tone model dictionary W. The method can therefore be seen as a non-negative matrix division in contrast to the uninformed matrix factorization.

3 TONE MODEL LEARNING

The method as pointed out so far requires a given dictionary of tone models that has to be learned in advance. In this work an approach is explained where the same algorithm as for the pitch decomposition is used. The necessary training data consists of recordings of single pitches played on the particular instrument. Starting from equation (1) again instead of fixing W to a given dictionary, H is chosen to have a number of components $r = 1$. This does justice to the facts that there shall be exactly one basis vector per midi pitch and in a single training instance there is only one tone present. The values of H are set to the corresponding values of the amplitude envelope expressing the current loudness of the sound. Then the tone model is calculated as described in section 2.3 but having a fixed H instead of a fixed W.

In cases where only recordings of some notes and not the whole pitch range are available interpolation is applied. Given a fundamental frequency f_0 for which the tone model is not known the starting point are the two nearest frequencies f_l and f_h with given energy distribution such that $f_l < f_0 < f_h$. The interpolation is then done in two steps. At first for each bin's center frequency f within the spectrum of f_0 the corresponding frequency within the known models is calculated. In the second step the energy of f' is estimated using parabolic interpolation. Finally the approximations using the lower and the upper frequency model are combined by taking the average.

4 FRAME-WISE CLASSIFICATION AND TRANSCRIPTION

The steps described in sections 2.3 and 3 lead to a piano-roll-like representation of the activation of individual pitches given by the matrix H. In order to close the gap between this representation and a symbolic transcription in midi-format a final step of post-processing is still needed since the activation data (i) contains low energy noise as well as higher level outliers and (ii) is not separated into discrete note events.

All the above calculations have been at the level of individual frames. Since musical notes are in most cases stable over a certain amount of time it makes sense to use the information given within a local neighborhood. We do so by applying a median filter in order to smooth the activation curves as well es to eliminate very short fragments. In our experiments window lengths of around 100 ms have been found to yield good results.

Since the smoothed activation data is very sparse, discrete note events can easily be extracted by identifying segments with above zero activation while filling out very small gaps (up to 30 ms) that might be there even after smoothing.

5 EXPERIMENTAL RESULTS

5.1 Tone model learning

As our prime test environment we have chosen solo piano music. In order to learn the dictionary for our experiments we took recordings of single tones played on a computer controlled Bösendorfer SE290 grand piano like they were also used by Goebl [3]. Recordings were available for every fourth midi pitch at different velocity levels. The power spectrum was computed using a short time Fourier transform with a window length of 4096 samples and a hop size of 441 frames or 10 ms when having input data sampled at a rate of 44.1 kHz. Additional zero padding by a factor of 2 was used in order to get narrower frequency bins. The window used was a Hamming window.

The resulting spectrum was then preprocessed in two steps. At first a silence detector sets all frames which have a total energy below a certain threshold to zeros. Then in order to further suppress noise as well as to remove the bias due to the tone model learned from one specific instrument, magnitude warping and spectral average subtraction as described in [7] is performed. The spectrum $X(k)$ of the signal is assumed to be a combination of $S(k)$ representing the sound that is originally excited, $H(k)$ being the frequency response of the environment like the instrument body and an additional noise component $N(k)$, giving the decomposition

$$X(k) = H(k)S(k) + N(k) \tag{6}$$

In order to equalize the factor $H(k)$ the power spectrum $X(k)$ is magnitude warped by applying

$$Y(k) = \ln\left(1 + \frac{1}{g}X(k)\right) \tag{7}$$

The purpose of the term g is to normalize the spectrum such that the level of the noise $N(k)$ is close to one whereas the spectral peaks are much larger. Assuming that the major part of $X(k)$ is just noise floor and the peaks of $H(k)S(k)$ are quite sparse any outlier resistible average measure can be taken in order to find a feasible g. In our tests just using the minimum of $X(k)$ gave satisfying results as well. Due to this warping the influence of $H(k)$ is reduced.

The additive noise is suppressed by subtraction of a moving average $\overline{N}(k)$ within the logarithmic scale. The size of the sliding window has a width of 100 Hz but increases at higher frequency bands such that it always covers at least a range of ± 4 semitones with regard to the currently processed coefficient k. The moving average $\overline{N}(k)$ representing the noise floor is then subtracted from $Y(k)$ leaving the preprocessed spectrum $Z(k)$ as

$$Z(k) = \max\left\{0, Y(k) - \overline{N}(k)\right\} \tag{8}$$

According to [7] using the logarithmic scale gives clearly better results than the linear scale.

The preprocessed recordings of the single tones are then passed to the model learner as described in section 3. Recordings were available for every fourth pitch using velocities of 30, 50, 70, 90 and 110. The influence of loudness of the training data is weakened by its explicit consideration during the learning algorithm. But since individual harmonics fade out unequally the initial loudness still influences the resulting model. We overcome this effect by taking into account the models learned from all different velocities and calculating a final one by taking the average spectral power at each frequency bin. To complete the dictionary containing all midi pitches from 21 to 108, which constitutes the whole pitch range of the piano, the missing tone models were interpolated.

5.2 Transcription

As test data we use the recording of Mozart's sonata KV279 played by a professional pianist on a computer monitored Bösendorfer SE290 grand piano, giving us a precise ground truth of played notes in a midi-like format. The test set of 10.000 time frames contains 1087 keystrokes and continuous pedal events that are known as well. Although this covers only less than the

first two minutes of the piece it is a respectable basis for cross validation experiments as will be described in this section.

The data was converted to be monaural and transformed into the frequency domain using the same parameters and preprocessing as used for the dictionary learning. The only difference was that the upper half of the frequency bands were dropped in order to reduce computational costs on further operations. Tests have shown that this reduction of data causes hardly any loss of quality. The power spectrum was then processed using the non-negative matrix division approach including smoothing of the result as well as the extraction of discrete note events. First tests have shown that the resulting set of identified notes includes 1077 out of the 1087 present notes, missing less than 1%. However the amount of spurious events being more than three times as high as the number of correct notes was unacceptable and made further post-processing necessary.

The data showed that the higher the pitches, the higher are the activation levels of occurring spurious events. A reason might be that due to the disregard of the higher half of the spectrum the higher notes within the dictionary are only represented by the fundamental frequency and the first one or two harmonics. Such a base vector is more likely to match a noise pattern than one containing the whole range of harmonics as is the case with models of lower pitches.

For this reason it is not adequate to use a single magnitude threshold on the activation data in order to distinguish correct note events from spurious ones. Instead we applied a very simple rule based classifier (RB) that has one magnitude threshold per frequency band as the only rules.

We compared this classifier to a second, instance based, one. We decided to use a nearest neighbor algorithm (IB) having a broader basis of decision-making. The features used were pitch, length, maximum energy and the sum of energy of a note. A 10-fold cross validation on our data set was done to test the ability of these two classifiers to separate correct notes from spurious ones. The results are listed in table 1 showing that the instance based classifier clearly outperforms the rule based one on all measures defined as

$$precision = \frac{TP}{TP + FP} \tag{9}$$

$$recall = \frac{TP}{TP + FN} \tag{10}$$

$$f = 2 \cdot \frac{precision \cdot recall}{precision + recall} \tag{11}$$

where TP are the correctly found notes, FN are the missed and FP the spurious notes. A note event is counted as correct if the transcribed and the real note do overlap. Cases where the sustain pedal is used are handled in the way that notes are allowed to last as long as the pedal is pressed but they do not need to since the actual length of the note cannot be told exactly.

	precision	recall	f
raw data	21.8%	99.1%	0.357
classifier (RB, notes)	85.9%	85.4%	0.856
classifier (IB, notes)	95.6%	88.1%	0.917
classifier (frames)	42.3%	68.8%	0.524

Table 1. Classification results on solo piano music

Determining note onsets and offsets by just considering if the activation is above zero is simple. However this usually leads to transcriptions into notes that are longer than the original ones. To overcome this drawback we applied another classifier to the raw activation data in order to decide whether a pitch is played or not for each frame individually. In our test set of 10.000 frames containing 88 pitches each, this leaves us with 880.000 instances with more than 20.000 instances belonging to an actually played note. The features given to the, again instance based, classifier were the ones used for note-wise processing with addition of the current activation of each instance.

The result is again shown in table 1. The recall means that almost 70% of the original sound is covered in the transcription. 42% of the play time within the result match with a real note whereas the remainder of 58% is made up by erroneous note elongations and spurious notes.

6 CONCLUSIONS

In this work we have proposed a modification to existing transcription approaches using non-negative matrix factorization. Instead of tackling the problem in an uninformed way the new method makes use of an a priori learned dictionary of base vectors or tone models. This transforms the problem from general factorization to a division problem. The methods for the calculation of activation magnitudes can also be applied to the initial model learning in order to yield an appropriate dictionary. An advantage over uninformed matrix factorization is that single time frames can be processed independently - a fact that can be utilized to reduce computational complexity.

Applied to solo piano music, the raw resulting activation patterns contained more than 99% of the original notes but more than three times as many spurious notes as well. A post processing step with quite simple classifiers achieved an overall f-value of about 90% for

note-wise detection of played notes. In comparison, the frame wise classification yields an f-value of about 0.5 which is significantly less accurate.

We believe that additional information like high-level musical knowledge could help to improve the final step of picking correct notes while neglecting spurious ones. Also the information from an additional onset detection could benefit the frame-wise detection accuracy. Steep slopes can be observed at the beginning of connected parts within the activation data. Yet their reliability for onset detection has not been investigated.

Another aspect that has not been considered yet is how to cope with situations where there is more than one instrument present, or pieces where the playing instrument is not known a priori. Although the preprocessing that was applied in the test environment does some spectral whitening and therefore reduces the influence of timbre we still expect the timbral correlation between the instrument used for the dictionary learning and the one that shall be transcribed to be essential.

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VOCAL SEGMENT CLASSIFICATION IN POPULAR MUSIC

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ABSTRACT

This paper explores the vocal and non-vocal music classification problem within popular songs. A newly built labeled database covering 147 popular songs is announced. It is designed for classifying signals from $1sec$ time windows. Features are selected for this particular task, in order to capture both the temporal correlations and the dependencies among the feature dimensions. We systematically study the performance of a set of classifiers, including linear regression, generalized linear model, Gaussian mixture model, reduced kernel orthonormalized partial least squares and K-means on cross-validated training and test setup. The database is divided in two different ways: with/without artist overlap between training and test sets, so as to study the so called 'artist effect'. The performance and results are analyzed in depth: from error rates to sample-to-sample error correlation. A voting scheme is proposed to enhance the performance under certain conditions.

1 INTRODUCTION

The wide availability of digital music has increased the interest in music information retrieval, and in particular in features of music and of music meta-data, that could be used for better indexing and search. High-level musical features aimed at better indexing comprise, e.g., music instrument detection and separation [13], automatic transcription of music [8], melody detection [2], musical genre classification [10], sound source separation [18], singer recognition [16], and vocal detection [4]. While the latter obviously is of interest for music indexing, it has shown to be a surprisingly hard problem. In this paper we will pursue two objectives in relation to vocal/non-vocal music classification. We will investigate a multi-classifier system, and we will publish a new labeled database that can hopefully stimulate further research in the area.

While almost all musical genres are represented in digital forms, naturally popular music is most widely distributed, and in this paper we focus solely on popular music. It is not clear that the classification problem can be generalized between genres, but this is a problem we will investigate in later work.

Singing voice segmentation research started less than a decade ago. Berenzweig and Ellis attempted to locate the vocal line from music using a multi-layer perceptron speech model, trained to discriminate 54 phone classes, as the first step for lyric recognition [4]. However, even though singing and speech share certain similarities, the singing process involves the rapid acoustic variation, which makes it statistically different from normal speech. Such differences may lie in the phonetic and timing modification to follow the tune of the background music, and the usage of words or phrases in lyrics and their sequences. Their work was inspired by [15] and [19], where the task was to distinguish speech and music signals within the "music-speech" corpus: 240 $15s$ extracts collected 'at random' from the radio. A set of features have been designed specifically for speech/music discrimination, and they are capable of measuring the conceptually distinct properties of both classes.

Lyrics recognition can be one of a variety of uses for vocal segmentation. By matching the word transcriptions, it is applicable to search for different versions of the same song. Moreover, accurate singing detection could be potential for online lyrics display by automatically aligning the singing pieces with the known lyrics available on the Internet. Singer recognition of music recordings has later received more attention, and has become one of the popular research topics within MIR. In early work of singer recognition, techniques were borrowed from speaker recognition. A Gaussian Mixture Model (GMM) was applied based on Mel-frequency Cepstral Coefficients (MFCC) to detect singer identity [20]. As briefly introduced, singing voices are different from the conventional speech in terms of time-frequency features; and vocal and non-vocal features have differences w.r.t. spectral distribution. Hence the performance of a singer recognition system has been investigated using the unsegmented music piece, the vocal segments, and the non-vocal ones in [5]. 15% improvement has been achieved by only using the vocal segments, compared to the baseline of the system trained on the unsegmented music signals; and the performance became 23% worse when only non-vocal segments were used. It demonstrated that the vocal segments are the primary source for recognizing singers. Later, work on automatic singer recognition took vocal segmentation as the first step to enhance

the system performance, e.g. [16].

Loosely speaking, vocal segmentation has two forms. One is to deal with a continuous music stream, and the locations of the singing voice have to be detected as well as classified, one example is [4]. The second one is to pre-segment the signals into windows, and the task is only to classify these segments into two classes. Our work follows the second line, in order to build models based on our in-house Pop music database. A detailed description of the database will be presented in section 4. The voice is only segmented in the time domain, instead of the frequency domain, meaning the resulting vocal segments will still be a mixture of singing voices and instrumental background. Here we will cast the vocal segments detection in its simplest form, i.e. as a binary classification problem: one class represents signals with singing voices (with or without background music); the other purely instrumental segments, which we call accompaniment.

In this paper we study this problem from a different angle. Several classifiers are invoked, and individual performance (errors and error rates) is inspected. To enhance performance, we study the possibility of sample-to-sample cross-classifier voting, where the outputs of several classifiers are merged to give a single prediction. The paper is organized as follows. Section 2 explains the selection of features. Classification frameworks are covered by section 3. With the purpose of announcing the Pop music database, we introduce the database design in section 4. In section 5, the experiments are described in depth, and the performance characteristics are presented. At last, section 6 concludes the current work.

2 ACOUSTIC FEATURES

2.1 Mel-Frequency Cepstral Coefficients

MFCCs are well-known in the speech and speaker recognition society. They are designed as perceptually weighted cepstral coefficients, since the mel-frequency warping emulates human sound perception. MFCCs share two aspects with the human auditory system: A logarithmic dependence on signal power and a simple bandwidth-to-center frequency scaling so that the frequency resolution is better at lower frequencies. MFCCs have recently shown their applicability in music signal processing realm, e.g. [1] for music genre classification, [16] and [5] for singer recognition, and [14] for vocal segmentation, and many more exist.

Features are extracted from short time scales, e.g. $20ms$, due to the stationarity of music signals. To process windows at longer time scales, temporal feature integration is needed. Features at different time scales may contain different information. A small frame size may result in a noisy estimation; and a long frame size may cover multiple sounds (phonemes) and fail to capture appropriate information.

2.2 Multivariate AR

During the course of searching for appropriate features, researchers have realized that system performance can be improved by combining short-time frame-level features into clip-level features. Feature integration is one of the methods to form a long-time feature, in order to capture the discriminative information and characterize how frame-level features change over longer time periods for a certain task. Often the mean and variance of several short-time features are extracted as the clip-level features [17], using multivariate Gaussian model or a mixture of them. However, both the mean-variance and mean-covariance model fail to capture the temporal correlations. A frequency band approach has been proposed in [9], and the energy of the features was summarized into 4 frequency bands. Even though this method can represent temporal development, it does not model the feature correlations.

The multivariate autoregressive model (MAR) was recently introduced to music genre classification [11], and a detailed comparison of different temporal feature integration methods was reported. MAR being able to capture both the temporal correlations and the dependencies among the feature dimensions, has shown its superiority for representing music. We adapt this model in the feature extraction phase on top of short-time MFCCs. Here, a brief description of MAR will be given, for detail, see [11].

Assume the short-time MFCC at time t is denoted as $\mathbf{x}_t$, which is extracted from a short period of stationary signals. The MAR can be stated as,

$$\mathbf{x}_t = \sum_{p=1}^{P} \mathbf{A}_p \mathbf{x}_{t-p} + \mathbf{u}_t, \tag{1}$$

where $\mathbf{u}_t$ is the Gaussian noise $\mathcal{N}(\mathbf{v}, \mathbf{\Sigma})$, assumed i.i.d. $\mathbf{A}_p$ is the coefficients matrix for order p; and if it is defined as a diagonal matrix, dependencies among dimensions will not be considered. P indicates the order of the multivariate auto regressive model, meaning that $\mathbf{x}_t$ is predicted from the previous P short-time features. It is worth to mention that the mean of MFFCs $\mathbf{m}$ is related to the mean of the noise $\mathbf{v}$ in the following way (note: $\mathbf{I}$ is an identity matrix),

$$\mathbf{m} = (\mathbf{I} - \sum_{p=1}^{P} \mathbf{A}_p)^{-1} \mathbf{v}. \tag{2}$$

3 CLASSIFICATION FRAMEWORKS

We have examined a number of classifiers: linear regression model (LR), generalized linear model (GLM), Gaussian mixture model (GMM), reduced kernel orthonormalized partial least squares (rKOPLS) and K-means.

As the problem is a binary task, only a single dimension is needed for linear regression, and the labels are coded as

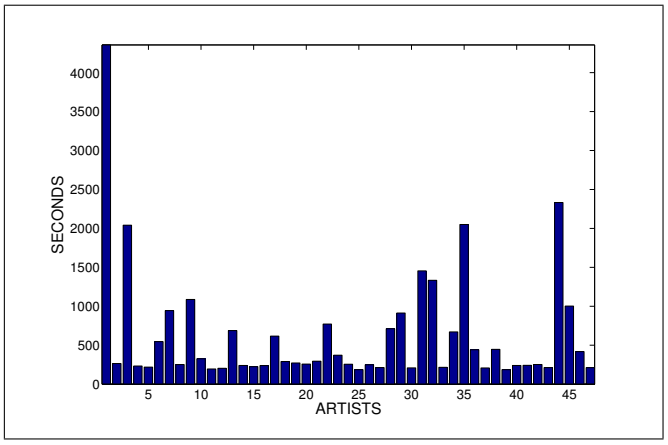

Figure 1. Distribution of Pop music among artists

± 1. The model is $l_n = \mathbf{w^T y}$. A 1 is added to the feature vector to model offset. Least squares is used as the cost function for training, and the minimum solution is the pseudo inverse. The prediction is made based on the *sign* of the output: we tag the sample as a vocal segment if the output is greater than zero; and as a non-vocal segment otherwise.

Generalized linear model relates a linear function of the inputs, through a link function to the mean of an exponential family function, $\mu = g(\mathbf{w}^T \mathbf{x}^n)$, where $\mathbf{w}$ is a weight vector of the model and $\mathbf{x}^n$ is the n'th feature vector. In our case we use the *softmax* link function, $\mu_i = \frac{e^{\mathbf{w}_i^T \mathbf{x}_i^n}}{\sum_j e^{\mathbf{w}_j^T \mathbf{x}_j^n}}$. $\mathbf{w}$ is found using iterative reweighted least squares [12].

GMM as one of the Bayesian classifiers, assumes a known probabilistic density distribution for each class. Hence we model data from each class as a group of Gaussian clusters. The parameters are estimated from training sets via the standard Expectation-Maximization (EM) algorithm. For simplicity, we assume the covariance matrices to be diagonal. Note that although features are independent within each mixture component due to the diagonal covariance matrix, the mixture model does not factorize over features. The diagonal covariance constraint posits the axes of the resulting Gaussian clusters parallel to the axes of the feature space. Observations are assigned to the class having the maximum *posterior* probability.

Any classification problem is solvable by a linear classifier if the data is projected into a high enough dimensional space (possibly infinite). To work in an infinite dimensional space is impossible, and kernel methods solve the problem by using inner products, which can be computed in the original space. Relevant features are found using orthonormalized partial least squares in kernel space. Then a linear classifier is trained and used for prediction. In the reduced form, rKOPLS [3] is able to handle large data sets, by only using a selection of the input samples to compute the relevant features, however all dimensions are used for the linear classifier, so this is not equal to a reduction of the training set.

K-means uses K clusters to model the distribution of each class. The optimization is done by assigning data points to the closest cluster centroid, and then updating the cluster centroid as the mean of the assigned data points. This is done iteratively, and minimizes the overall distances to cluster centroids. Optimization is very dependent on the initial centroids, and training should be repeated a number of times. Prediction is done by assigning a data point to the class of the closest cluster centroid.

4 DATABASE

The database used in the experiments is our recently built in-house database for vocal and non-vocal segments classification purpose. Due to the complexity of music signals and the dramatic variations of music, in the preliminary stage of the research, we focus only on one music genre: the popular music. Even within one music genre, Berenzweig et al. have pointed out the 'Album Effect'. That is songs from one album tend to have similarities w.r.t. audio production techniques, stylistic themes and instrumentation, etc. [5].

This database contains 147 Pop mp3s: with 141 singing songs and 6 pure accompaniment songs. The 6 accompaniment songs are not the accompaniment of any of the other singing songs. The music in total lasts $8h\ 40min\ 2sec$. All songs are sampled at 44.1 kHz. Two channels are averaged, and segmentation is based on the mean. Songs are manually segmented into $1sec$ segments without overlap, and are annotated second-by-second. The labeling is based on the following strategy: if the major part of this $1sec$ music piece is singing voice, it is tagged as vocal segment; otherwise non-vocal segment. We believe that the long-term acoustic features are more capable of differentiating singing voice, and $1sec$ seems to be a reasonable choice based on [14]. Furthermore labeling signals at this time scale is not only more accurate, but also less expensive.

Usually the average partition of vocal/non-vocal in Pop music is about 70%/30%. Around 28% of the 141 singing songs is non-vocal music in the collection of this database. Forty-seven artists/groups are covered. By artists in Pop music we mean the performers (singers) or bands instead of composers. The distribution of songs among artists is not even, and Figure 1 gives the total number of windows (seconds) each artist contributes.

5 EXPERIMENTS AND RESULTS

We have used a set of features extracted from the music database. First, we extracted the first 6 original MFCCs over a $20ms$ frame hopped every $10ms$. The 0^{th} MFCC representing the log-energy was computed as well. The means were calculated on signals covering $1sec$ in time. MAR were afterwards computed on top of the first 6 MFCCs with $P = 3$, and we ended up with a 6-by-18 $\mathbf{A}_p$ matrix, a 1-by-6

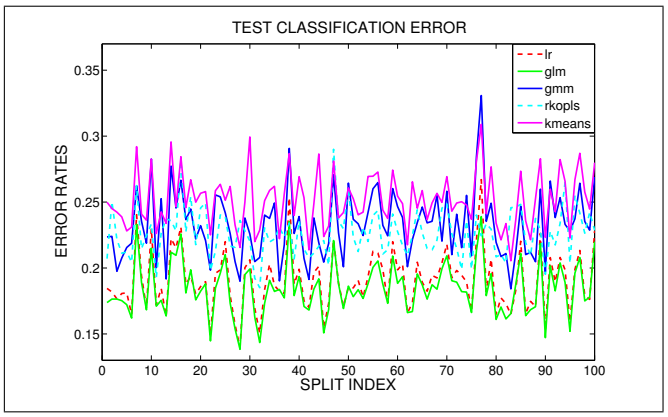

Figure 2. Classification error rates as a function of splits of five classifiers on test sets.

vector $\mathbf{v}$ and a 6-by-6 covariance matrix Σ. Since Σ is symmetric, repetitions were discarded. $\mathbf{A}_p$, $\mathbf{v}$ and Σ all together form a 135-dimensional feature set. The choice for 6 MFCC is on one hand empirical, and on the other hand to reduce the computational complexity. All in all, for $1sec$ music signal we concatenated 135-d MAR, the means of both 0^{th} and 6 original MFCCs to form a 142-d feature vector.

5.1 Data Dependency and Song Variation

We used one type of cross-validation, namely holdout validation, to evaluate the performance of the classification frameworks. To represent the breadth of available signals in the database, we kept 117 songs with the 6 accompaniment songs to train the models, and the remaining 30 to test. We randomly split the database 100 times and evaluated each classifier based on the aggregate average. In this way we eliminated the data set dependencies, due to the possible similarities between certain songs. The random splitting regarded a song as one unit, therefore there was no overlap song-wise in the training and test set. On the other hand artist overlap did exist. The models were trained and test set errors were calculated for each split. The GLM model from the Netlab toolbox was used with *softmax* activation function on outputs, and the model was trained using iterative reweighted least squares. As to GMM, we used the generalizable gaussian mixture model introduced in [7], where the mean and variance of GMM are updated with separate subsets of data. Music components have earlier been considered as 'noise' and modeled by a simpler model [16], thus we employed a more flexible model for the vocal than non-vocal parts: 8 mixtures for the vocal model, and 4 for the non-vocal model. For rKOPLS, we randomly chose 1000 windows from the training set to calculate the feature projections. The average error rates of the five classification algorithms are summarized in the left column of Table 1.

A bit surprisingly the performance is significantly better for the linear models. We show the performance of the chosen classifiers as a function of splits in Figure 2. Each curve represents one classifier, and the trial-by-trial difference is quite striking. It proved our assumption that the classification performance depends heavily on the data sets, and the misclassification varies between 13.8% and 23.9% for the best model (GLM). We envision that there is significant variation in the data set, and the characteristics of some songs may be distinguishing to the others. To test the hypothesis, we studied the performance on individual songs. Figure 3 presents the average classification errors of each song predicted by the best model: GLM, and the inter-song variation is obviously revealed: for some songs it is easy to distinguish the voice and music segments; and some songs are hard to classify.

	Error Rates	
Artists	overlap	no overlap
LR	19.03±2.25%	20.52±3.5%
GLM	18.46±2.02%	19.82±2.81%
GMM	23.27±2.54%	24.50±2.99%
rKOPLS	22.62±1.85%	24.60±3.14%
K-means	25.13±2.11%	NA

Table 1. The average error rates (mean ± standard deviation) of 5 classifiers on test sets.

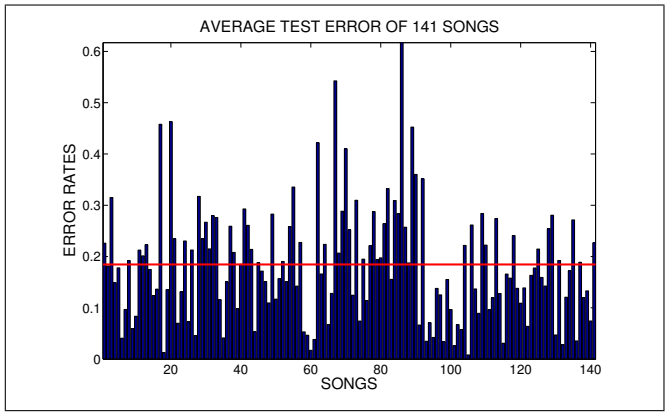

Figure 3. Test classification error rates for individual songs by GLM model. The dash line gives the average error rates of the 100-split cross-validation.

5.2 Correlation Between Classifiers and Voting

While observing the classification variation among data splits in Figure 2, we also noticed that even though classification performance is different from classifier to classifier, the tendency of these five curves does share some similarity. Here we first carefully studied the pair-to-pair performance correlation between the classification algorithms. In Table 2 the degree of matching is reported: 1 refers to perfect match; 0 to no match. It seems that the two linear classifiers have a very high degree of matching, which means that little will be gained by combining these two.

The simplest way of combining classification results is

	LR	GLM	GMM	rKOPLS	K-means
LR	1.0000	0.9603	0.8203	0.8040	0.8110
GLM	0.9603	1.0000	0.8141	0.8266	0.8091
GMM	0.8203	0.8141	1.0000	0.7309	0.7745
rKOPLS	0.8040	0.8266	0.7309	1.0000	0.7568
K-means	0.8110	0.8091	0.7745	0.7568	1.0000

Table 2. A matrix of the degree of matching.

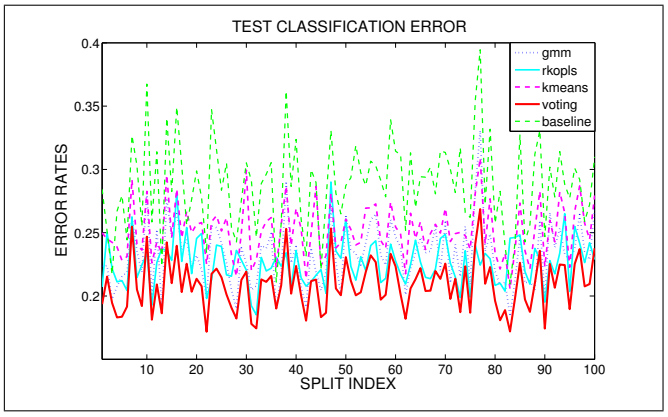

Figure 4. Voting results. It gives the voting performance among GMM, rKOPLS and K-means. The light dash line shows the baseline of random guessing for each data split.

by majority voting, meaning that the class with the most votes is chosen as the output. The voting has been done crossing all five classifiers, unfortunately the average voting results (error rates) on the test sets was 18.62%, which is slightly worse than the best individual classifier. The reason seems to be that even though the other classifiers are not so correlated with the linear ones, the miss classification rate is too high to improve performance.

However voting does help enhance the performance, if it performs among not so correlated classification results. Figure 4 demonstrates the sample-to-sample majority voting among three classifiers: GMM, rKOPLS and K-means. The similar tendency was preserved in the voting results, and there were only 10 splits out of 100, where the voting results were worse than the best ones among these three. The average performance of voting on test sets was $20.90 \pm 2.02\%$.

Here we will elaborate on the performance on individual songs, by looking at the predicted labels from each classifier and voting predictions. Figure 5 demonstrates how voting works, and how the prediction results correlate. Two songs: 'Do You Know What You Want' by M2M, and 'A Thousand Times' by Sophie Zelmani, have been chosen to illustrate the 'good' and 'bad' cases, i.e. when voting helps and fails. Vocal segments are tagged with '1', and '0' for non-vocal ones. The ground truth is given as a reference. The voting was carried out among GMM, rKOPLS and K-means, and their predictions are shown. If the classifiers make mistakes in a similar pattern, the voting cannot recover the wrong predictions, e.g. area B. If the predictions are not correlated to a high degree voting helps, e.g. area A.

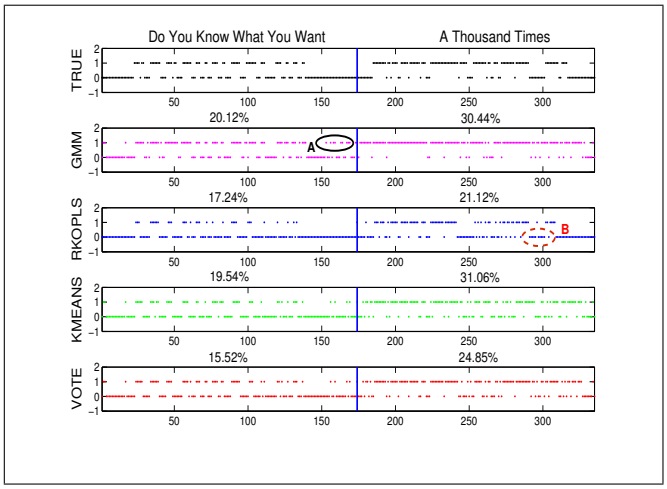

Figure 5. Sample-to-sample errors and voting results. Two songs represent the 'good' and 'bad' voting cases. Individual error rates for each classifier and voting results are given. Two areas marked A & B indicate the scenarios when voting helps and fails.

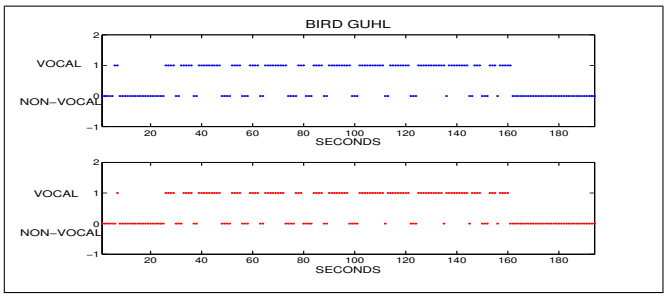

Figure 6. Two manual label results of the same song: 'Bird Guhl'. It is obvious that the disagreement only appears in the transition parts.

Moreover, we noticed that it is very likely for classifiers to make wrong predictions in the transition sections, meaning the changing from vocal to non-vocal parts, and vice versa. We found this is reasonable comparing with manual labels by different persons, shown in Figure 6. The song was labeled carefully by both people, the absence of mind or guessing should not be a concern. The mismatch indicates the perception or judging difference, and it only happens in the transition parts. The total mismatch is about 3% for this particular song: 'Bird Guhl' by Antony and the Johnsons.

5.3 'Artist Effect'

In previous experiments, we randomly selected songs to form training and test sets, hence the same artist may appear in both sets. Taking the previous results as a baseline, we studied the 'artist effect' in this classification problem. We tried to keep the size of test sets the same as before, and carefully selected around 30 songs in order to avoid artist overlap for each split, and formed 100 splits. The second column of Table 1 summarizes the average error rates for 4 classi-

fiers. The average results are a little worse than the previous ones, and they also have bigger variance along the splits. Therefore we speculate that artists do have some influence in vocal/non-vocal music classification, and the influence may be caused by different styles, and models trained on particular styles are hard to be generalized to other styles.

6 CONCLUSION AND DISCUSSION

We have investigated the vocal/non-vocal popular music classification. Experiments were carried out on our database, containing 147 popular songs. To be in line with the label set, the classifiers were trained based on features at $1sec$ time scale. We have employed 142-d acoustic features, consisting MFCCs and MAR, to measure the distinct properties of vocal and non-vocal music. Five classifiers have been invoked: LR, GLM, GMM, rKOPLS and K-means.

We cross-validated the entire database, and measured the aggregate average to eliminate the data set dependency. GLM outperformed all the others, and provided us with 18.46% error rate on the baseline of 28%. The performance has great variation among data splits and songs, indicating the variability of popular songs. The correlations among classifiers have been investigated, and the proposed voting scheme did help among less correlated classifiers. Finally we looked into the 'artist effect', and it did degrade the classification accuracy a bit by separating artists in training and test sets. All in all vocal/non-vocal music classification was found to be a difficult problem, and it depends heavily on the music itself. Maybe classification within similar song styles can improve the performance.

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PERCEPTUALLY-BASED EVALUATION OF THE ERRORS USUALLY MADE WHEN AUTOMATICALLY TRANSCRIBING MUSIC

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ABSTRACT

This paper investigates the perceptual importance of typical errors occurring when transcribing polyphonic music excerpts into a symbolic form. The case of the automatic transcription of piano music is taken as the target application and two subjective tests are designed. The main test aims at understanding how human subjects rank typical transcription errors such as note insertion, deletion or replacement, note doubling, incorrect note onset or duration, and so forth. The Bradley-Terry-Luce (BTL) analysis framework is used and the results show that pitch errors are more clearly perceived than incorrect loudness estimations or temporal deviations from the original recording. A second test presents a first attempt to include this information in more perceptually motivated measures for evaluating transcription systems.

1 INTRODUCTION

In the benchmarking of Information Retrieval systems, performance is often evaluated by counting and classifying errors. Classically the ratio of relevant items that are returned out of the full set of original ones, referred to as *recall*, measures the completeness of the system performance whereas the proportion of relevant items that are retrieved, or *precision*, indicates the correctness of the answer. The F-measure, combining precision and recall, offers a single score to assess the performance. When music processing systems are involved, the question arises as to how to complement such a quantitative assessment by incorporating a certain amount of perceptually motivated criteria or weights.

This paper investigates the perceptual importance of typical errors occurring when transcribing polyphonic music excerpts into a symbolic form, *e.g.* converting a piece recorded in a PCM (.wav) format into a MIDI file. This particular Music Information Retrieval (MIR) task and its related subtasks (onset detection, multipitch estimation and tracking) have received a lot of attention [9] from the MIR community since the early works of Moorer [14] in the mid 70s. The approaches used to accomplish the goal are very diverse [4, 5, 14, 15, 16] and the evaluation of the performance for such systems is almost as varied. Some papers [4, 14] focus on a couple of sound examples, to probe typical errors such as octave errors, or deviations from ground truth such as duration differences, and so forth. However, the most widely used criteria for assessing automatic transcription are quantitative, even if the evaluation framework is not always similar (frame-based [15], note-based [16] or both [1]).

In the practical context of piano music for instance, the evaluation task is often handled by generating the PCM format piece from an original MIDI file which makes it possible to compare the input (ground truth) and output MIDI files. For that particular case, in this study, a perception test has been designed for subjectively rating a list of typical transcription errors (note insertions, deletions, incorrect onsets or duration...). The test is based on pairwise comparisons of sounds holding such targeted errors. The results are then analyzed by means of the Bradley-Terry-Luce (BTL) method [3].

In a second step, the question emerged of finding a way to take into account the perceptual ranking of the discomfort levels we obtained. Another test was designed to subjectively compare transcriptions resulting from different systems. It aimed at deriving more perceptually relevant metrics from the preceding BTL results by synthetically combining their main findings, and at checking their compliance with the test results. We worked in two directions: perceptually weighting typical errors, countable by comparing the input and output MIDI files, and adaptating similarity metrics [17].

2 THE EVALUATION MEASURES

The commonly-used F-measure is defined by:

$$f \triangleq 2\frac{rp}{r+p} = \frac{\#\text{TP}}{\#\text{TP} + \frac{1}{2}\#\text{FN} + \frac{1}{2}\#\text{FP}} \qquad (1)$$

The authors thank all the subjects involved in the perceptive test for their participation. They also thank M. Castellengo, A. de Cheveigné, D. Pressnitzer and J. Benenson for their useful remarks, and M. Marolt, N. Bertin and P. Leveau for sharing their programs. The research leading to this paper was supported by Institut TELECOM under the *Automatic Transcription of Music: Advanced Processing and Implementation - TAMTAM* project and by the French GIP ANR under contract ANR-06-JCJC-0027-01, *Décomposition en Éléments Sonores et Applications Musicales - DESAM*.

where r denotes the recall, p the precision, #TP the number of true positives (TP), #FN the number of false negatives (FN) and #FP the number of false positives (FP). f is equivalent to the quantity a, that is referred to as either accuracy or score [5], since $f = \frac{2}{\frac{1}{a}+1}$. The F-measure is useful to obtain the error rate for individually counted errors, but does not consider aspects like sequentiality, chords, harmonic or tonal relationships, etc.

Another evaluation approach comes from the problem of finding the similarity between two (musical) sequences. At the moment, these methods are commonly used to search for similar melodies in large databases, rather than in the field of the evaluation of transcriptions.

Let us assume that one must compare two sequences of symbols, A and B. The Levenshtein's distance, or edit distance [11], is a metric that counts the minimal number of operations necessary to transform A to B. The possible operations on symbols are: deletion from A, insertion into B, or replacement of a symbol in A by another one in B.

Mongeau and Sankoff [13] proposed adapting this distance to the case of monophonic musical sequences, in order to define a similarity metric between two melodies. The two sequences of notes are ordered according to the onset of each note. Each note is characterized by its pitch and duration, which are used to compute the cost of the following possible operations: insertion, deletion, replacement, with costs depending on tonal criteria, fragmentation and consolidation of several notes with the same pitch. These operations reflect typical mistakes in transcriptions. The minimum distance between the sets of notes is then estimated using the edit distance framework.

This melodic edit distance being applicable only to monophonic sequences, an extension to the polyphonic case has been recently proposed [8]. In order to represent the polyphonic nature of musical pieces, quotiented sequences are used. So far, this representation has only been applied to chord sequences, which constitute a restricted class of musical pieces: the notes within a chord must have the same onset and duration.

Another way to compute the similarity between two musical sequences [17] consists in considering each set of notes as points in a multidimensional space, *e.g.* the pitch/time domain. The algorithm is based on two choices. First, each point must be assigned a weight, *e.g.* the note duration. Second, a distance between a point in the first set and a point in the second one is defined, *e.g.* the euclidian distance in the time/pitch space. Then, the overall distance can be computed with the *Earth Movers Distance* (EMD) or the *Proportional Transportation Distance* (PTD). It is related to the minimum amount of work necessary to transform one set of weighted points to the other using the previously-defined distance, making it possible to transfer the weight of a source note towards several targets.

In all of these methods, the setting of the parameters is a crucial point. Indeed, the weighting between the time and the pitch dimensions, for instance, depends on music perception. The tests presented in this paper aim at assessing the trends of the perceptive impact of typical errors and the distribution of their related weights.

3 EXPERIMENTAL SETUP

3.1 Overview

The perception test consists of two tasks, which are detailed below. It was available on the Internet in the spring of 2007 for two weeks and was announced by e-mail. Before accessing the tasks, the subject is given instructions and information on the recommended audio device (high-quality headphones or loudspeakers, and a quiet environment) and on the estimated duration of the test. He or she is then invited to complete the tasks. Both of them consist in hearing a musical excerpt and several transcriptions of it, and in focusing on the discomfort caused by the transcriptions, with respect to the original. Task 1 uses artificial transcriptions, *i.e.* some copies of the original piece into which errors were inserted whereas task 2 uses transcriptions obtained by automatic transcription systems. In both cases, the transcriptions are resynthesized in the same recording conditions as the original piece in order to be heard and compared by the subject. At the end, the subject was asked to describe the criteria he used to compare files and to add any comments. Due to the total duration of the test a subject can possibly endure (about 40' here), we limited the scope of the study to pieces of classical piano music, from different periods, with different tempi and harmonic/melodic content.

3.2 Test 1: Subjective Evaluation of Typical Transcription Errors

3.2.1 Principle

Test 1 aims at obtaining a specific score for typical transcription errors. In order to achieve this, the transcriptions to be evaluated are made by inserting one and only one kind of error into an original excerpt. The error is chosen among the following list of typical errors: note deletion, random-pitched note insertion (1 to 11 half-tones), random-pitched note replacement (1 to 11 half-tones), octave insertion, octave replacement, fifth insertion, fifth replacement, note doubling, onset displacement, duration change (offset modification) and loudness modification (MIDI velocity).

These errors are inserted into three excerpts from *Studies, op 10 / Study 1 in C Major* by Chopin (8 seconds), *Suite Bergamasque / III. Clair de Lune* by C. Debussy (20 seconds), and *Sonata in D Major KV 311 / I. Allegro con Spirito* by W.A. Mozart (13 seconds).

Ideally, we would like to obtain a ranking of the typical errors. Due to the large number of files, asking the subjects

Figure 1. Test 1: for each pair of audio files, the subject selects the one causing more discomfort.

to give a score to each of them is not feasible. We preferred to set up a pairwise comparison task, as shown in Figure 1 and derived the full scale as described in the next section.

3.2.2 Protocol and Settings

For each kind of error, several test files are created with various error rates. The number of modified notes is parametered by the Modified Note Rate (MNR), which is set to either 10%, or 33%. For some kinds of error, the error intensity (EI) is also parametrized. This is quantified as a ratio of the note duration for duration changes and onset changes, and as a ratio of the MIDI velocity for loudness modifications. The EI is set to either 25%, or 75%. Modified notes are randomly chosen using the MNR. Intensity changes are made randomly, uniformly in the range centered on the true value and with the EI as radius.

To derive a ranking scale from pairwise comparisons, we choose the BTL method which uses hidden, "true" values associated to the transcriptions, along a given dimension (here, the discomfort). For a given pair of transcriptions, the subject's answer is a comparison of a noisy version of the two true values, the noise modeling the subjectivity and the variable skill of subjects. Thanks to this statistical framework, the full subjective scale is then obtained by processing all the pairwise comparisons. For this test, 20 pairs out of 812 are randomly chosen and presented to each subject for each musical excerpt. This number has been chosen in order to adjust the test duration and is not critical for the results, as long as the number of subjects is high enough.

3.3 Test 2: Subjective Evaluation of Transcriptions of Musical Pieces

Test 2 aims at obtaining a perceptive score for a series of transcriptions from several pieces of music. Three original excerpts from *Prelude in C minor BWV 847* by J.S. Bach (13 seconds), *Suite Bergamasque / III. Clair de Lune* by C. Debussy (20 seconds), and *Sonata in D Major KV 311 / I. Allegro con Spirito* by W.A. Mozart (13 seconds) were chosen. For each excerpt, five transcriptions are presented, as shown in Figure 2. The subject has to assign a non-negative value to each transcription. These values express the discomfort caused by transcription errors in comparison with its reference. The subject can listen as many times as needed to each transcription and reference.

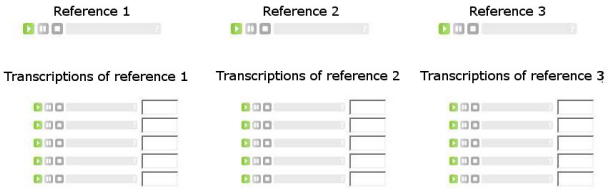

Figure 2. Test 2: the subject scores transcriptions with non-negative values.

In this test, all subjects are presented exactly the same audio files, in random order for each subject. One of the five transcriptions is the original piece in order to check whether the answers are consistent. The other four were obtained by automatic transcription systems, namely SONIC [12], available on the author's website, Bertin's system [2], a homemade system by P. Leveau based on [10] and an early version of [7]. The error rates and kinds of error thus depend on the specific behaviors of the transcription systems.

4 RESULTS

Thirty-seven subjects (24 musicians and 13 non-musicians) took part in this test. The results of Tests 1 and 2 are detailed here. The subjects' comments show that the instructions were understood correctly. They pointed out tone errors as a major cause of discomfort, while they seldom mentioned loudness and duration errors in an explicit way.

4.1 Test 1

Results of Test 1 are given in Figure 3. The BTL method makes it possible to obtain, from the pairwise comparisons of all the subjects, a subjective scale of discomfort for typical errors. A BTL perception value is thus assigned to each modification, which can be ordered according to this scale.

Different forms of evidence show the consistency of the obtained scale. First, increasing scores are obtained with increasing error rates, either MNR or EI, and decreasing harmonicity (octave, fifth, random pitches). Second, a minimum discomfort is obtained for the reference (taking into account its confidence interval). Third, as described in [6], the above 90% confidence intervals are related to a 5% risk. Thus, they are narrow enough to distinguish error types and to assert that the answers make sense, although adjacent error types should be considered perceptually equivalent.

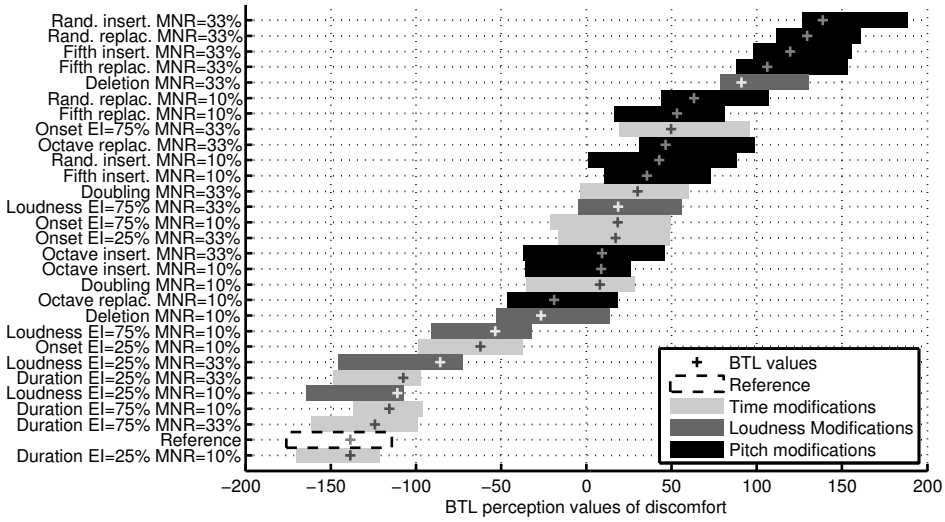

Figure 3. Test 1 : perceptive scale for typical errors. Crosses account for the related BTL value. Horizontal bars depict the 90% confidence intervals, obtained by a bootstrap method [6] using 100 resamplings of the data (because the data is not gaussian, confidence intervals may not be centered on BTL values).

Globally, as expected from the subjects' comments, the highest discomfort values are obtained with pitch modifications; loudness and time modifications cause low to medium discomfort. Regarding pitch changes, octave errors are judged much less serious than fifth changes, which cause a slightly lower discomfort than random changes. In each case, replacements and insertions are judged as equivalent, which would indicate that the discomfort is more induced by the added note than by the deleted note of a replacement. The lower values obtained for deletions confirm this hypothesis, which is commonly observed when working on transcription systems: a false negative usually sounds better than a false positive.

Results with time errors show that the modified onsets cause much more discomfort than duration changes. While one can expect that moving the beginning of an event causes a significant subjective change, it seems that subjects just did not perceive most of the modifications of duration. This can be explained by a specific feature of the piano: the ends of sounds generated from its freely-vibrating strings are less perceptible than for a musical instrument with forced vibrations. Thus current results for perception of note duration cannot be generalized to all instruments.

Finally, additional analysis of the results should be reported, which are not represented in Figure 3. First, similar results were obtained from subjects that were musicians and from non-musicians. Second, the three scales obtained for the three excerpts are also similar, with a little difference for the excerpt by Debussy in which deletions have a lower score and duration changes cause higher discomfort, probably because of the long durations in this slow piece of music.

4.2 Test 2

For each subject, scores of Test 2 are normalized by the maximum score he/she gave. Six subjects were removed since they scored a discomfort greater than 20% for the reference. Average scores and variances were then computed, with respect to the results from all the remaining subjects.

Results are represented in Figure 4. As the test is not a contest between existing algorithms, the systems were made anonymous, numbered from 1 to 4. The confidence in the results is assessed thanks to a 3 (composers) $\times$ 5 (algorithms) factorial ANOVA test, passed for each dimension and for interactions using a $p = 0.01$ test level. Thereby, the scores and the ranking of the algorithms are very dependent on the piece of music. This confirms that the performance of a transcription system is related to the musical content of pieces and thus depends on the test database. Large standard deviations indicate that the evaluation of a musical transcription depends greatly on proper subjective criteria. An important overlap between the answers makes it impossible to obtain a definitive ranking among the different algorithms even if for each excerpt, systems 2 and 3 are judged as the worst and the best one respectively.

5 EVALUATING THE TRANSCRIPTIONS

When comparing the results given by one of the objective evaluation methods proposed in Section 2 to the perceptive results of Test 1, several aspects are differentiated in the former case while they are not in the latter case, and vice versa. For instance, octave, fifth and random pitch changes have

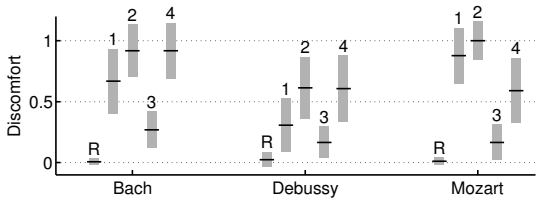

Figure 4. Results of Test 2: perceptive evaluation of the reference (R) and of transcriptions from four systems (1-4), with standard deviations (gray bars).

similar F-measures but cause an increasing discomfort. On the contrary, perceptive results are equivalent for replacements and insertions, whereas the F-measure is higher for insertions. Besides, perceptive results provide a balance between time and pitch influences, which is not taken into account in objective evaluation methods.

In this part, we estimate weighting coefficients of typical errors from Test 1 results, and we then apply them to adapt two existing metrics: the F-measure and the PTD. These modified methods are validated by applying them to the excerpts used in Test 2 and by comparing the results with the discomfort expressed by the subjects.

5.1 Extraction of the Weighting Coefficients

To extract the weighting coefficients, the results of Test 1 are normalized between 0 and 1. We only used results with MNR 33%, and the results were averaged for pitch modifications, insertions and replacements. Six criteria [1] are obtained, to be integrated into metrics. Their related weighting coefficients are given in the following table:

Octave	Fifth	Other intervals	Deletion	Duration	Onset
$\alpha_1 =$	$\alpha_2 =$	$\alpha_3 =$	$\alpha_4 =$	$\alpha_5 =$	$\alpha_6 =$
0.1794	0.2712	0.2941	0.2475	0.0355	0.4687

The coefficients are normalized so that $\frac{1}{3}\sum_{i=1}^{3}\alpha_i + \sum_{i=4}^{6}\alpha_i = 1$, since octave, fifth and random pitch represents alternative false positive errors.

5.2 Perceptive F-measure

In eq.(1), errors are the number of FP and FN, with an equal weight ($\frac{1}{2}$). We thus define the perceptive F-measure by:

$$f_{\text{percept}} \triangleq \frac{\#\text{TP}}{\#\text{TP} + \sum_{i=1}^{6}\alpha_i w_i \#E_i} \qquad (2)$$

[1] Loudness was not considered since the results were not satisfying, probably due to the difficulty of having a trustworthy loudness scale. Doubled notes were not used either because they could not be integrated into metrics.

where $\#E_i$ is the number of errors of type i (see below), $w_1 = w_2 = w_3 = w_4 = 1$, w_5 is the average duration error in the transcription, and w_6 is the average onset error. (The average errors are computed as the square root of the mean square error.) Note that a similar perceptive accuracy could be defined by using the equivalence mentioned in Section 2 and that the expression (2) equals the F-measure in the case $\alpha_1 = \alpha_2 = \alpha_3 = \alpha_4 = \frac{1}{2}$ and $\alpha_5 = \alpha_6 = 0$.

Errors from MIDI files are extracted as follows:

1. TP are estimated as notes with correct pitch (rounded to the nearest semitone) and onset deviation lower than 150 ms. For each TP, the relative onset deviation and the relative duration deviation (both with respect to the original note parameters) are extracted. Then, let $\#E_5 = \#E_6 = \#\text{TP}$.

2. FP are transcribed notes which are not TP. The set of FP is split as follows: for each FP, (a) if there is an original note at the octave or sub-octave, at the same time (*i.e.* with any overlap of both time supports), the FP is added to the set E_1 of octave FP; (b) otherwise, if there is an original note at the upper or lower fifth at the same time, the FP is added to the set E_2 of fifth FP; (c) otherwise, the FP is added to the set E_3 of other pitch FP.

3. FN are the set E_4 of original notes that are not associated with one TP.

5.3 Perceptive PTD

The PTD is originally used to evaluate melodic similarities (see Section 2). In this context, note duration as weights to transfer and the euclidian distance in the time/pitch space seem to be appropriate choices. Nevertheless, when comparing generic musical pieces, both of these choices should be changed. PTD weights should be defined in a musical sense but this is beyond the scope of the current work and we thus chose to assign an equal and unitary PTD weight to each note. Using the perceptual coefficients introduced in Section 5.1, the distance between two notes is then defined in the multidimensionnal space of criteria composed of pitch (octave, fifth or others), duration and onset modifications.

5.4 Results

Figure 5 shows the results of the application of the original two objective measures and of their perceptive versions to the musical excerpts from Test 2. In order to compare them to the discomfort results, F-measures were changed by applying the function $x \mapsto 1 - x$. In order to best fit the discomfort scale, all results were scaled by a multiplicative coefficient obtained by minimizing the mean square error.

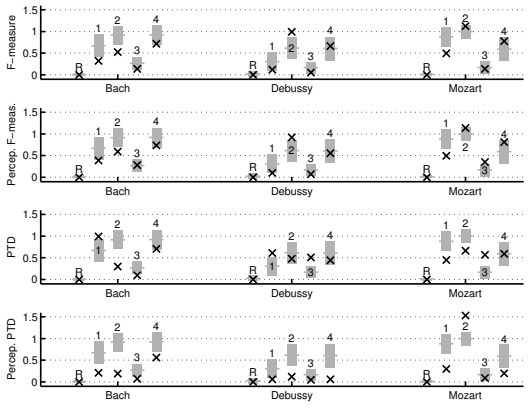

Figure 5. Transcription evaluation results with several objective and perceptive measures: in each case, crosses show the normalized error related to a measure, and the gray bars indicate the discomfort obtained in Test 2.

Results with the perceptive F-measure are slightly closer to the discomfort values than the original F-measure. Moreover, the ranking of the 15 excerpts is also closer to the discomfort-based ranking. Results of the perceptive PTD do not look better than the original, due to a high isolated value for the excerpt with highest discomfort (Mozart, System 2), that makes it difficult to scale the results adequately. However, the achieved ranking is dramatically better than the ranking by the original PTD, and also slightly better than the ranking by the perceptive F-measure. Thus, even if the relation between the discomfort and the perceptive PTD may be non-linear, the latter is appropriate in a ranking task.

6 CONCLUSIONS

The main idea of these tests was to get a ranking of the typical automatic transcription errors, to extract perception weights, and to integrate them into several musical sequence distance metrics. These primary results are consistent and the proposed perceptive metrics give satisfying results.

However further investigations should focus on a number of aspects, such as non-linear relations between specific error rates and discomfort, musical-based typical errors (taking into account tonality, melody, chords, etc.), and more specific algorithms to identify them.

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MULTIPLE-INSTANCE LEARNING FOR MUSIC INFORMATION RETRIEVAL

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ABSTRACT

Multiple-instance learning algorithms train classifiers from lightly supervised data, i.e. labeled collections of items, rather than labeled items. We compare the multiple-instance learners mi-SVM and MILES on the task of classifying 10-second song clips. These classifiers are trained on tags at the track, album, and artist levels, or granularities, that have been derived from tags at the clip granularity, allowing us to test the effectiveness of the learners at recovering the clip labeling in the training set and predicting the clip labeling for a held-out test set. We find that mi-SVM is better than a control at the recovery task on training clips, with an average classification accuracy as high as 87% over 43 tags; on test clips, it is comparable to the control with an average classification accuracy of up to 68%. MILES performed adequately on the recovery task, but poorly on the test clips.

1 INTRODUCTION

There are many high quality sources of metadata about musical material such as Last.fm, the All Music Guide, Pandora.com, etc. Typically, however, each source provides metadata only at certain granularities, i.e. describes the music only at certain scales. For example, the All Music Guide provides metadata about many artists and albums, but few tracks. Similarly, Last.fm users have described a large proportion of artists, a smaller proportion of albums, and an even smaller proportion of tracks. Furthermore, there are no publicly accessible, large-scale sources of metadata describing parts of tracks known as *clips*, here taken to be 10-second excerpts. This paper describes the use of clip-level classifiers to refine descriptions from one granularity to finer granularities, e.g. using audio classifiers trained on descriptions of artists to infer descriptions of albums, tracks, or clips.

Many descriptions of music apply at multiple granularities, like *rap*, or *saxophone*, although certain descriptions are valid only at specific granularities like *seen live* or *albums I own*. Descriptions valid at one granularity, however, might only apply to certain elements at a finer granularity. For example, at the artist level, the Beatles could very reasonably be tagged *psychedelic*. This tag would certainly apply to an album like *Sgt. Pepper's Lonely Hearts Club Band* but would not apply to one like *Meet the Beatles*. Similarly, the John Coltrane track "Giant Steps" could very reasonably be tagged *saxophone*. While valid for most clips in the track, it most notably is not valid during the piano solo. This paper describes systems capable of deriving, from feature similarity and a list of *psychedelic* artists or *saxophone* tracks, the clips for which these tags are most appropriate.

By considering tags one at a time, as either being present or absent from a clip, we pose the automatic tagging (autotagging) problem as clip classification. In this framework, the task of metadata refinement is known as *multiple instance learning* (MIL) [6]. In MIL, classifiers are trained on labels that are only applied to collections of *instances*, known as *bags*. Positive bags contain one or more positive instances, while negative bags contain no positive instances. Labeled bags provide less information to the learner than labeled instances, but are still effective at training classifiers. For the purposes of this paper, clips are the instances to be classified, and artists, albums, and tracks, in turn, are the bags.

There are two problems addressed in the multiple-instances learning (MIL) literature, the classification of bags and the classification of instances. As we are interested in refining musical metadata from bags to the instances within them, we only concern ourselves with multiple-instance learners that are capable of classifying instances. A related problem that we also examine is the training of a general instance-level classifier from bag-level labels. This task is slightly different in that the instances to be labeled are not in the training bags, and are unseen at training time.

To evaluate the applicability of MIL to music, we use the data from our MajorMiner game [10]. The game has collected approximately 12,000 clip-level descriptions of approximately 2,200 clips from many different tracks, albums, and artists. The most popular descriptions have been applied to hundreds of clips, and there are 43 tags that have been applied to at least 35 clips. Previous authors have generally used datasets that were labeled at the bag level, making it difficult to evaluate instance-level classification. Sometimes a subset of the data was laboriously annotated to allow the evaluation of instance-level classification. In the MajorMiner dataset, however, tags are applied directly to clips, making it

possible to test instance-level classification in both the training set and a separate test set. By deriving bag tags from clip tags in the training set, we can directly test the ability of multiple instance learners to recover metadata at the instance level from the bag level. This derivation adheres to the MIL formulation, labeling a given bag positive as a positive example of a tag if any of its clips have been labeled with that tag. In addition, a held-out test set allows us to evaluate the generalization of these classifiers to instances outside the training set.

1.1 Previous work

A number of authors have explored the link between music and text. Whitman and Ellis [14] trained a system for associating music with noun phrases and adjectives using a collection of reviews from the All Music Guide and Pitchfork Media. This work was based on the earlier work described in [15]. More recently, [12] used a naive Bayes classifier to both annotate and retrieve music based on an association between the music and text. Eck et al. [7] used boosted classifiers to identify the top k tags describing a particular track, training the classifiers on tags that the users of Last.fm had entered for the track's artist.

The MIL problem was first formulated for the task of digit recognition [8], in which a neural network was trained with the information of whether a given digit was present, but not where it was present. In this case, the bags were regions in which the digit was known to be present and the instances were shifted and windowed sub-regions. Another early application of MIL was to the problem of drug discovery [6], in which the bags were molecules and the instances were conformations of those molecules.

MIL has also been applied to object detection in images, in which the bags were images and the instances were either automatically segmented image regions [5] or automatically identified interest points [3]. It has been applied to video classification to match names and faces [16], in which the instances were (name, face) pairs, the bags were scenes, and the task was to determine whether a face had any names associated with it or not. And it has been applied to text classification [1], in which the bags were documents and the instances were sentences or paragraphs.

Many learning frameworks have been applied to MIL, including boosting [13], Markov chain Monte Carlo [3], and both 1-norm [4] and 2-norm support vector machines [1]. In this work, we compare the performance of two SVM-based MIL methods, mi-SVM [1] and MILES [4], on the task of autotagging 10-second song clips. These two algorithms are explained in greater detail in the next section.

2 MULTIPLE-INSTANCE LEARNING

The following notation is common to both mi-SVM and MILES. We denote the ith bag as B_i, of size ℓ_i, and the jth instance in that bag as x_{ij} where $j \in 1 \dots \ell_i$. The label for bag i is $Y_i \in \{-1, 1\}$ and the label for instance x_{ij} is y_{ij}. The set of positive bag indices is defined as $I^+ \equiv \{i : Y_i = 1\}$ and similarly the set of negative bag indices $I^- \equiv \{i : Y_i = -1\}$. All instances in a negative bag are negative and a single positive instance in a bag forces the bag to be positive, meaning that

$$Y_i = \max_j y_{ij}. \tag{1}$$

2.1 The mi-SVM algorithm

The mi-SVM algorithm is the instance-level MIL support vector machine classifier presented in [1]. Support vector machines generally maximize the margin around a hyperplane separating positive from negative examples. In the MIL setting, however, the optimal labeling of the points in positive bags must be computed as well, creating the following mixed integer quadratic program (QP)

$$\begin{aligned} \min_{\{y_{ij}\},\mathbf{w},b,\xi} \quad & \tfrac{1}{2}\|\mathbf{w}\|_2^2 + C\sum_{ij}\xi_{ij} \\ \text{subject to} \quad & \forall i : y_{ij}(\langle \mathbf{w}, \mathbf{x}_{ij}\rangle + b) \geq 1 - \xi_{ij} \\ & y_{ij} \in \{-1, 1\} \\ & \xi_{ij} \geq 0 \\ & \forall i \in I^+ : \sum_{j=1}^{\ell_i} \tfrac{1}{2}(y_{ij} + 1) \geq 1 \\ & \forall i \in I^- : y_{ij} = -1. \end{aligned} \tag{2}$$

Unfortunately, it is difficult to solve this integer program directly and [1] presents a heuristic means of approximating it that converges to a local optimum. This heuristic solves the standard SVM quadratic program with the labels fixed, and then uses the solution of the QP to impute the missing labels for instances in positive bags. This alternation continues until the labels no longer change. The instance labels are initialized from the bag labels, so that all instances in positive bags are initially labeled positive and all instances in negative bags are initially (and subsequently) labeled negative.

The number of iterations required for convergence depended on the number of instances in each bag, but was generally on the order of 10-20. We use the dual domain formulation of the standard SVM QP so that a nonlinear kernel could be used to define the similarity between instances; in particular, we used the radial basis function (RBF) kernel. In this dual domain, the classifier is a linear combination of a hopefully sparse subset of the training instances, the support vectors.

2.2 The MILES algorithm

Another SVM-based multiple-instance learner is Multiple-Instance Learning via Embedded Instance Selection (MILES)

[4]. While MILES is mainly a bag classifier, it is also able to derive classifications for instances. Classification with MILES proceeds in three steps. In the first step, bags are projected into a large feature space by computing the similarity between each bag and all of the training instances. The similarity between a bag and an instance is defined as the maximum similarity between any of the instances in that bag and the instance in question,

$$K(B_i, \mathbf{x}) \equiv \max_{k \in 1 \ldots \ell_i} K(\mathbf{x}_{ik}, \mathbf{x}). \quad (3)$$

The feature vector for one bag is then

$$\mathbf{m}_i \equiv \left[K(B_i, \mathbf{x}_{11}) \ldots K(B_i, \mathbf{x}_{N\ell_N})\right]^T. \quad (4)$$

In the second step, a 1-norm support vector machine [2] simultaneously learns a classifier and selects discriminative "features," i.e. instances. The 1-norm SVM solves the following linear program

$$\begin{aligned} \min_{\mathbf{w},b,\xi} \quad & \|\mathbf{w}\|_1 + C\mu \sum_{i \in I^+} \xi_i + C(1-\mu) \sum_{i \in I^-} \xi_i \quad (5) \\ \text{subject to} \quad & \forall i : Y_i(\langle \mathbf{w}, \mathbf{m}_i \rangle + b) \geq 1 - \xi_i \\ & \xi_i \geq 0. \end{aligned}$$

This optimization can be solved as written in the primal domain by representing $\mathbf{w}$ as the difference of two non-negative vectors. The ℓ_1-norm in the objective function encourages sparsity in the elements of $\mathbf{w}$ forcing many of them to 0. Since $\mathbf{w}$ multiplies instance-similarities, only the instances corresponding to nonzero elements of $\mathbf{w}$ affect the classification of new bags. These instances can be considered the support vectors in this case. Thus, the ℓ_1 SVM is also sparse in its selection of instances.

In the third step, classifications of some of the instances are derived from the bag classifications. There is one instance in every bag selected by (3) as the closest to each support vector. A bag's classification is not affected by instances that are not the closest to any support vector, and those instances can be ignored. Any remaining instance that contributes more than a threshold amount to the classification of the bag is considered to be that class itself. We treat unclassified instances as negatively classified even though [4] allows instances to remain unclassified.

While MILES has a number of parameters that need to be tuned, we found its performance quite consistent across a wide range of parameter settings. The μ parameter controls the penalty for misclassifying an instance from a positive bag versus misclassifying an instance from a negative bag. We found it to have little effect in our experiments except when within 10^{-4} of 1, so we kept it at 0.5. The C parameter controls the trade-off between increasing the margin and violating class constraints. We found that it could affect performance when it was small and determined that a value of 10 worked well. The σ parameters is the standard deviation of the RBF kernel. With feature vectors of unit norm, as described in the next section, $\sigma = 1$ worked well. Similarly, mi-SVM has C and σ parameters for which $C = 1$ and $\sigma = 1$ worked well. All of these parameters were tuned on a subset of the tags using different train/test breakdowns of the artists from our main experiments.

3 MUSICAL FEATURES

We use two types of features to describe musical audio. The spectral features come from our earlier work [9] and capture timbral aspects of the music related to instrumentation and production quality. The temporal features are novel, and summarize the beat, tempo, and rhythmic complexity of the music in four different frequency bands. All of these features are calculated on 10-second long clips of songs.

The temporal features are similar to those described in [11]. They are calculated on the magnitude of the Mel spectrogram, including frequencies from 50 Hz to 10,000 Hz, using a window size of 25 ms and a hop size of 10 ms. The mel bands are combined into four large bands at low, low-mid, high-mid, and high frequencies giving the total magnitude in each band over time. The bands are windowed and their Fourier transforms are taken, from which the magnitude of the 0-10 Hz modulation frequencies are kept. The DCT of these magnitudes is then taken and the bottom 50 coefficients of this *envelope cepstrum* are kept for each band. The four bands' vectors are then stacked to form the final, 200-dimensional feature vector.

The spectral features consist of the mean and unwrapped covariance of a clip's mel-frequency cepstral coefficients (MFCCs). The MFCCs are calculated from the mel spectrogram used in the temporal features above. Since the on-diagonal variance terms are strictly positive, their log is taken to make their distribution more Gaussian. We use 18-dimensional MFCCs, and only keep the unique elements of the covariance matrix, for a total of 189 dimensions.

The 0-th cepstral coefficient, the DC modulation frequency, and the on-diagonal variance terms tend to be much larger than other features. We therefore scale each feature to be zero-mean and unit-variance across the set of all instances to make the features more comparable to each other. This scaling improves classification because those dimensions that are biggest are generally not the most discriminative. After the feature dimensions are scaled, the temporal features are multiplied by $\frac{1}{\sqrt{2}}$, a value that was determined empirically to balance the spectral and temporal features well. Finally, the feature vector for each clip is normalized to have unit length to avoid problems with degenerate clips.

4 EXPERIMENTS

The data used in our experiments come from our MajorMiner game [10]. In this game, players label 10-second clips with

arbitrary textual descriptions called *tags*, scoring points when others describe the same clips with the same tags. The rules of the game encourage players to use original, yet relevant tags. In our experiments, we only include tags that have been verified by at least two different players on at least 35 clips, ensuring that the concept is relevant overall and to the individual clips. There are 43 such tags and they have been verified approximately 9000 times in total on approximately 2200 clips selected at random from 3900 tracks.

Note that these data do not include strict negative labels. While many clips are tagged *rock*, none are tagged *not rock*. Frequently, however, a clip will be tagged many times without being tagged *rock*. We take this as an indication that *rock* does not apply to that clip. More specifically, a negative example of a particular tag is a clip on which another tag has been verified, but the tag in question has not.

Certain data-handling issues required extra consideration. Before using the tags in our experiments, we performed a number of normalization steps on them, eliminating variations in punctuation, spelling, and suffixing. Around 45 silent clips were culled from the dataset and the last clip in each track was also discarded, as the irregular length of these clips can cause problems and they are generally silent. Finally, a number of tracks came from albums where the artist was indicated as "Various Artists" and "Original Soundtrack." We excluded these clips from any bags used in our experiments, but allowed them in the instance-level test set, where bags were ignored anyway.

4.1 Procedure

The experiment was five repetitions of a two-fold cross-validation procedure, where artists were assigned to one fold or the other to make sure tags were learned independent of artists. Each tag was evaluated independently as a binary classification task. To create the training set for a particular tag the positive bags with the most instances of that tag were selected from the training fold until either they were all selected or 400 instances had been selected. The negative bags with the most instances were then selected until there were at most as many negative instances as positive instances. On average, the selected track, album, and artist bags contained 2.47, 4.44, and 8.17 instances, respectively. In the training set, bag labels Y_i were calculated from instance labels y_{ij} according to (1).

As an illustration, to create the training and testing datasets for the tag *saxophone* at the artist granularity, all of the artists with any clips tagged *saxophone* are considered positive bags. So Sonny Rollins would be one positive bag, John Coltrane another, etc. All artists without any clips tagged *saxophone* are considered negative bags. To create the training set, the positive artists with the most labeled clips are selected, followed by the negative artists with the most labeled clips.

In addition to mi-SVM and MILES, two control algorithms were included in the evaluation. The first, referred to as the naïve approach, assumes that bag labels apply to all of the instances in a bag and trains an instance-labeling SVM on those labels directly. The second control, referred to as the cheating approach, trains on instance labels directly and serves as a ceiling on performance for instance classification in the training set. Since there are many negative instances in positive bags and this algorithm needs to train on an equal number of positive and negative examples, it only selects a subset of the negative instances to train on. This selection causes its slightly imperfect performance.

The maximum number of positive instances, 400, was chosen to give a reasonable running time for all algorithms. With 400 labeled instances in positive bags, the largest data sets would have 800 examples. Training an individual 1-norm or 2-norm SVM on this many examples takes only a few seconds, but there were about 5000 SVMs to be trained over 10 repetitions, on 43 tags, with 3 different bag sizes for each of 4 algorithms. The slowest algorithm was mi-SVM, which took 6.7 hours to run on one Intel Xeon 1.89 MHz CPU. Second was the naïve approach, which took 67 minutes, followed by MILES at 42 minutes and the cheating approach at 24 minutes.

We evaluate the classifiers on two different tasks. The first, is the recovery of instance labels in the training set. Since the classifier is trained only on bag labels and not directly on instance labels, this task is not trivial. To fix a performance baseline of 0.5, accuracy is evaluated on an equal number of positive and negative instances from the training bags. To increase the precision of the measurement, the maximum number of examples are selected while still maintaining balance, although this causes the variance of the estimates to differ between classes.

While recovering instance labels is one useful application of MIL, it does not allow the classifiers' performance on novel instances to be measured accurately, because of the partial supervision. We instead measure this using instances from the held-out test set, ignoring any notion of bags. To facilitate comparison between the accuracy of classifying each tag, the same number of test examples are selected for each tag, namely 11 positive and 11 negative (setting the baseline accuracy again to 0.5). When combined across two cross validation folds and five repetitions of the experiment, each class is tested on 220 examples. Since MILES classifies bags before classifying instances in those bags, each test point is presented to MILES in its own bag.

5 RESULTS

The overall classification accuracy of the various algorithms can be seen in Table 1. On the training instances, the cheating algorithm was the most accurate, followed by mi-SVM, the naïve algorithm, and MILES. Since the cheating algorithm is an upper bound on training set accuracy, mi-SVM is the most accurate realistic classifier, if only by a small margin.

	Training set			Test set		
	Trk	Alb	Art	Trk	Alb	Art
Cheat	85.9	83.9	82.6	64.7	65.7	66.2
mi-SVM	87.0	80.9	78.0	66.8	67.8	65.4
Naïve	85.0	79.5	77.3	67.4	67.7	66.0
MILES	81.2	73.2	70.0	51.7	50.9	50.6

Table 1. Overall classification accuracy percentages on labeled points in the train and test sets at track, album, and artist granularities.

The mi-SVM algorithm outperformed the naïve algorithm because the naïve algorithm generally returned the bag labels for the instances, while mi-SVM was able to identify some of the negative instances in positive bags.

On the test data, mi-SVM and the naïve algorithm performed equally well, followed by the cheating algorithm, and MILES. The inversion of the cheating and naïve algorithms on the test set was unexpected. It could possibly be caused by the naïve algorithm's ability to use more of the training examples presented to it, or it could indicate that the instance labels do contain some amount of noise that the bag labels smooth over. Such noise could be due to the lack of unambiguous negative labels in the MajorMiner data.

MILES was not very accurate on the test set. It was reasonably accurate on tags with many examples and small training bags, but otherwise it classified all test instances identically, resulting in an accuracy of 0.5. This might be due to a mismatch between the size of the bags in the training and test sets, since the test "bags" were all single instances. In experiments where MILES was tested on bags of a similar size to those it was trained on, its test accuracy more closely followed its training accuracy. In these experiments, MILES seemed to be able to rank test bags well, even when they differed in size from the training set, possibly indicating that the bias, b, was learned incorrectly. In the training set, its bag classifications were almost always correct, while its instance classifications suffered slightly from over-sparsity. In particular, as bag size increased, a smaller proportion of instances in the bag contributed to its classification, leaving all of the others classified as negative by default.

As expected, classification was more accurate on instances in training sets than in test sets. This indicates that even though training instances might not be labeled individually, bag labels still convey useful information. Also as expected, accuracy generally decreases for coarser-granularity bags, more so for training instances than for testing instances. This indicates that larger bags, while still allowing the training of classifiers, constrain the labels of training instances less.

In Figure 1, results on the test set are broken down by tag for each algorithm and bag size. This figure reveals the differences between the tags. Some tags perform better using track bags, while others perform better using artist or album bags. These differences could indicate the granularities at which each tag is appropriate. In particular, one can compare the performance of the naïve algorithm or mi-SVM trained on different bag granularities to determine if one granularity is significantly different from another.

The naïve algorithm trained on artist bags make the assumption that a tag that is appropriate for some of an artist's clips is also appropriate for the rest of them. The accuracy of the resulting classifier, relative to other granularities, should indicate how well this assumption holds. Since mi-SVM is initialized with the same labels as the naïve algorithm, a similar property might hold. We have found some evidence of this phenomenon in our results, but not enough to conclusively prove its existence. For example, the tags *saxophone*, *synth*, *piano*, *soft*, and *vocal* should be most relevant at the track level, and indeed, they train mi-SVM classifiers that are much better for track bags than for artist bags. A counterexample is provided by *trumpet*, however, which is better classified by mi-SVM trained on artist bags.

6 CONCLUSION

We have formulated a number of music information related multiple-instance learning tasks and evaluated the mi-SVM and MILES algorithms on them. By using clip-level tags to derive tags at the track, album, and artist granularities, we have created three different ground truth datasets. These datasets are suitable for testing the learners' ability both to recover the original tags for clips in the training set and to tag clips in a held-out test set. We found that mi-SVM was the most accurate in recovering tags in the training set, followed by the naïve approach of assuming bag tags applied to all instances, and then MILES. In predicting tags for test clips, we found that mi-SVM and the naïve approach were quite comparable, and both were much more accurate than MILES. While these results are promising, many multiple-instance learners have been formulated and it is possible that another one is more appropriate to the task of predicting tags.

6.1 Future work

The most straightforward extension of this work is to larger datasets with more tags and more labeled examples. It should be possible to evaluate the refinement of tags from the artist and album level to the track level, so data from sources like Last.fm, the All Music Guide, and Pandora could be used in the evaluation of MIL. It would also be interesting to qualitatively examine the results of refining labels from these data sources down to the clip level. Many other tasks in music information retrieval could also benefit from the decreased cost of collecting training data within the MIL framework, including polyphonic transcription, singing voice detection, structure finding, and instrument identification.

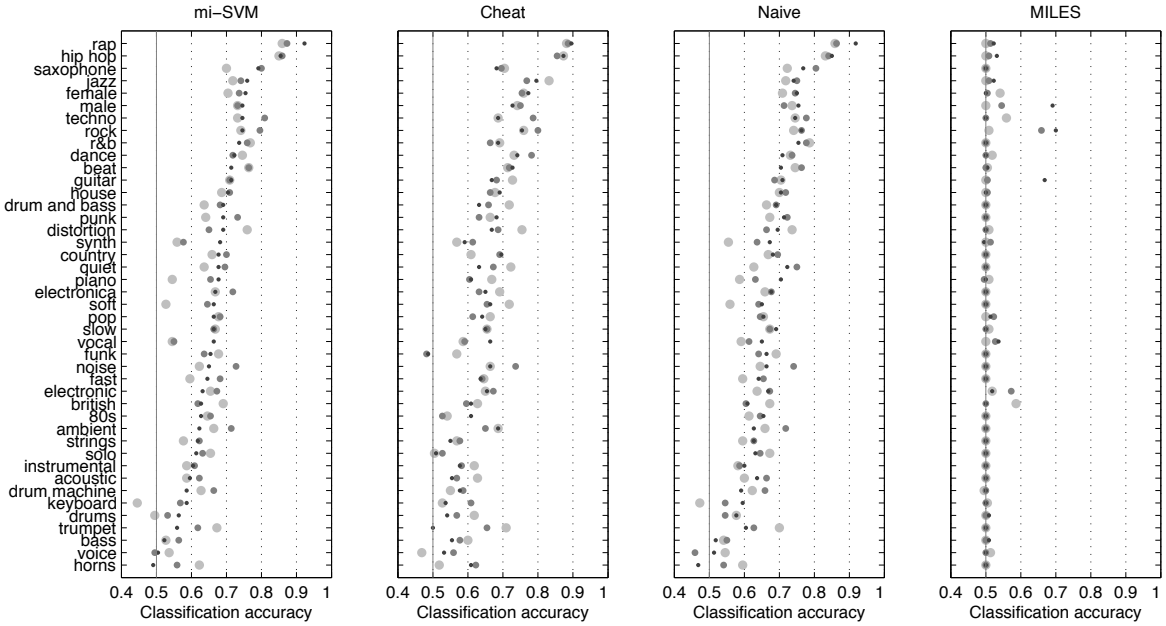

Figure 1. Classification accuracy on the test set, broken down by tag. Each pane is an algorithm, and each style of dot is a different bag granularity. Dots get larger and lighter for coarser granularities: track, album, artist. The tags are ordered by the accuracy of mi-SVM with track bags. For every dot, N=220, meaning that a difference of around 0.06 is statistically significant.

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OH OH OH WHOAH! TOWARDS AUTOMATIC TOPIC DETECTION IN SONG LYRICS

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ABSTRACT

We present an algorithm that allows for indexing music by topic. The application scenario is an information retrieval system into which any song with known lyrics can be inserted and indexed so as to make a music collection browseable by topic. We use text mining techniques for creating a vector space model of our lyrics collection and non-negative matrix factorization (NMF) to identify topic clusters which are then labeled manually. We include a discussion of the decisions regarding the parametrization of the applied methods. The suitability of our approach is assessed by measuring the agreement of test subjects who provide the labels for the topic clusters.

1 INTRODUCTION

The past few years have seen a tremendous amount of scientific work in the field of music information retrieval. Much, if not most of this work was concentrated on the acoustic properties of music, while the semantic content conveyed by the words of the songs was mostly ignored. There has been some work on song lyrics, but the field has not gained momentum comparable to work on acoustic features of music. In much of the older work lyrics were used to side or contrast with acoustic-based techniques in genre classification of songs [8] or artists [7]. In newer work, lyrics have become sources for metadata generation [9] and, probably inspired by the evolution of Web 2.0, lyrics were found useful as a basis for keyword generation for songs, a technique that may ultimately lead to automatic tagging [12].

In our view, any music browsing or recommendation system is incomplete if it does not incorporate the dimension of the songs' semantic content. It is therefore our goal to create elements of such a system based on the analysis of song lyrics. In the work at hand, we present the building blocks of a system that allows for searching a collection of lyrics by selecting from a set of topics. We describe the formal details of the procedure which we propose for making a collection of songs browseable by topic. Centrally to our algorithm, we apply NMF for clustering the lyrics documents as proposed in [13]. The resulting clusters are labeled manually so that they can be used as the dimensions of a topic space in which the documents are arranged.

The remainder of this paper is organized as follows: Section 2 describes the dataset we work with, in Section 3 we explain the details of our algorithm. Section 4 gives an account of the quality assessment we performed.

2 THE DATA

The music archive we work with is a subset of the collection marketed through Verisign Austria's [1] content download platform. This subset comprises approximately 60.000 of the most popular audio tracks [2] by some 6.000 artists. The tracks are affiliated with one or more of 31 genres. The most important genres are *"Pop"* (10817 songs), *"Alternative"* (9975 songs), *"Hip-Hop"* (4849), *"Rock"* (4632), *"Country"* (3936) and *"Dance/Electronic"* (3454).

The lyrics to these songs were extracted from the internet using the method presented in [5], which worked for roughly two thirds of the songs. The reason for this rather low rate of success is that the method we employed is not suitable for some classes of tracks. Such classes comprise of course the purely instrumental songs, but also works that bear a composite *artist* attribute, e.g., *"Mint Condition/feat. Charlie Wilson of the Gap Band"* or older and only locally known music. After removal of duplicates, which are also present in our dataset, we get a corpus of 33863 lyrics.

The lyrics are in 15 different languages, though the vast majority is in English. We found about 300 Spanish songs, roughly 30 in Italian, French, Gaelic and Latin; all other languages are even less frequent.

3 ALGORITHM

The method we propose requires the lyrics collection to be transformed into a vector space model in which each document is represented as a vector defining its affiliation to a set of topics. This transformation encompasses the following steps:

[1] `http://www.verisign.at/`

[2] i.e., the most frequently accessed tracks within a certain time span

1. *Preprocessing*
   Reading text from text files and creating a term-document matrix (TDM).

2. *Term Selection*
   Shrinking the TDM by dropping terms and documents.

3. *Term Weighting*
   Changing the values for the terms in the TDM.

4. *Clustering*
   Clustering the TDM using NMF.

5. *Labeling*
   Manually assigning labels to the topic clusters.

6. *Calculating Topic Affiliation*
   Automatically assigning topics to documents.

After this preparation the collection can be searched for topics using a query vector that defines the weight for each topic.

In the remainder of this section, we explain this algorithm and its parameters and name techniques for optimizing the results.

3.1 Preprocessing

To begin with, text files are read from the file system, the text is tokenized and a TDM is created without the application of stemming. For this step we use the R package *tm* [10, 3].

A closer look into our corpus reveals a correlation between the length of the text and its quality: the shorter it is, the more unlikely it is to contain the complete lyrics, but rather only a small part of it or, in some cases, only snippets of html code or text on a web page. A viable solution for improving the quality of the corpus is therefore to filter short lyrics. However, some songs actually do have very short lyrics (see Example 3.1); moreover, this is more common in some genres than in others, so filtering introduces a genre bias into the corpus. We decided to accept this trade-off on the one hand because our research is not focused on genres and on the other hand because lyrics that short do not convey much semantic content to analyze, at least not in textual form. [3] Our corpus was created using a lower threshold of 200 characters for the length of lyrics. The number of songs in the collection was thus reduced by approximately 3% to 32831.

[3] Note that brevity of lyrics may, however, be a valuable piece of information in a general music recommendation scenario.

Oh oh oh whoah
Oh oh oh whoah
Oh oh oh whoah

"Switch 625" by
Def Leppard

That's right
Have more rhythm
Woooo!
More rhythm

"Cherry Twist" by
The Chrystal Method

Example 3.1: Two songs with very short lyrics.

Our lyrics harvesting system assigns a confidence value to each extracted text (see [4]), which reflects the strength of agreement found in the different versions of the text. We use this value to identify lyrics to be deleted due to insufficient agreement and hence doubtful quality. The confidence value is a real value $\in [0, 1]$. Informal analysis showed that this value indicates poor quality quite realiably below 0.7. For lyrics with higher confidence value, a difference in this value did not seem correlated with a corresponding difference in the observed quality. These findings led us to set the lower threshold to 0.7, which further reduced the number of songs by 1.5% to 32323.

3.2 Term Selection

Reducing the number of terms is a powerful measure to influence the outcome of the subsequent clustering stage. On the one hand, this decreases the size of the TDM, making any subsequent computation faster and thus allowing more thorough experimentation. On the other hand, experiments showed that term selection can have tremendous influence on the kind of clusters that are produced.

Two approaches are used to cut down on the number of terms. First, stopword lists for a number of languages help to remove the most frequent words that are not considered to be helpful for topic clustering. We used stopwords [4] for English, Spanish, French and German and a custom stopword list for lyrics [5]. Second, terms and documents are deleted if they do not meet conditions defined by upper and lower thresholds for the document frequency [6] of a term (f_{max}, f_{min}) and by a minimal term count for documents (t_{min}).

Terms with particularly high or low document frequency are natural candidates for removal from the TDM. Terms occurring in only one document can hardly be useful for clustering, as they only add dimensions to the document representation in which documents are equally dissimilar from one another. We therefore chose to eliminate all terms with a document frequency of 1, which in our case reduced the memory consumed by the TDM by about 3%.

[4] These lists come with the R package *tm* and are part of the snowball stemmer (`http://snowball.tartarus.org/`)

[5] This list contains mainly exclamations like *"uuh"* and non-lyrics terms such as *"songlyrics"* or *"inurl"*

[6] i.e., the number of documents a term occurs in.

Terms with a very high document frequency account for dimensions of the document representation in which a great amount of documents are similar. With respect to clustering, these dimensions may be regarded as noise that makes the real clusters more difficult to discern. This rationale suggests that the clustering result improve when frequent terms are removed. Moreover, removing the frequent terms greatly reduces the space consumed by the TDM: in our case, deleting terms with a document frequency > 500 reduced the memory consumption by 42%. However, the disadvantage of this strategy may be that clusters mainly defined by frequent terms are lost.

Early findings suggested that the use of an upper limit for the document frequency makes the resulting clusters more diverse, but more systematic experimentation convinced us that it was better to use $f_{max} = \infty$.

3.3 Term Weighting

It has been proposed to use TF×IDF [11] as the weighting scheme prior to NMF clustering [13]. In text retrieval applications, the purpose of the weighting function is to amplify the weights of the terms that are most typical for a document and to lower the weights of the other terms. This is because the main use case of the method is *retrieval*, i.e., finding documents that match a user-defined query. In our algorithm, however, the use case to optimize for is not retrieval but *clustering*. During our experiments, we learned that this requires different properties of the weighting function.

We investigated the usefulness of three different weighing schemes in our context: term frequency weighting (i.e., no change to the TDM at all), TF×IDF and binary (i.e., replacing all nonzero entries in the TDM by 1). We found that binary weighting yielded the best results both in terms of individual cluster quality and evenness of the document-to-cluster distribution. Clustering using term frequency weighting produces term clusters in which the frequent terms are clearly overrepresented. Similarly, when applying TF×IDF weighting before clustering, the terms with low document frequency are too important in the resulting term clusters.

3.4 Clustering

We decided to use NMF [6, 13] for automatic topic detection as it is a clustering technique that results in additive representation of items [7] [6], a property that distinguishes it from most other clustering techniques. We also evaluated latent semantic analysis (LSA) [2], but found that it is not a suitable techniques for large sparse matrices due to its space complexity. Using NMF, the TDM is approximated by the matrix product of two matrices of appropriate dimensionality. NMF is parametrized most prominently by the number of clusters that it shall produce, k. More formally, let T be the TDM, W and H the factor matrices, and

$$T_{documents \times terms} = W_{documents \times k} H_{k \times terms} \quad (1)$$

The parameter k is the *inner dimension* of the factorization, i.e. the dimension that both factor matrices share. The approximation of the TDM by the NMF is depicted in Figure 1. For our purposes, the more important matrix of the two factor matrices is H, which contains the weight of each term in each cluster.

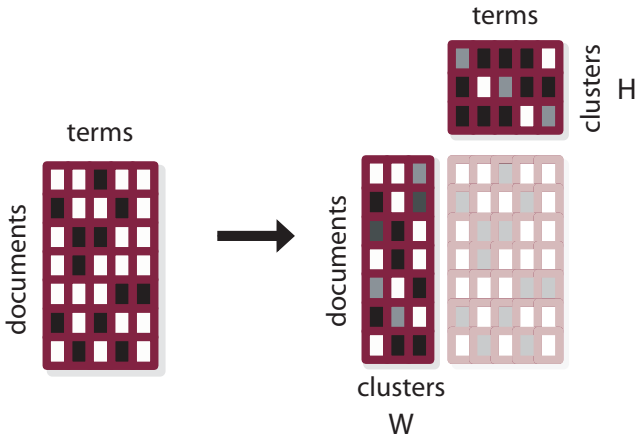

Figure 1. Non-negative factorization of the TDM. The TDM is approximated by the product WH.

A low value for k causes NMF to produce clusters that are actually mixtures of multiple topics, which may be related hierarchically, but this is not neccessarily the case. For instance, one cluster may describe the topic *"love"*, and on a closer look, the sub-topics *"loss"*, *"happy"* and *"family"* are recognizable, while another cluster could at first glance contain only the *"gangsta"* topic but at the same time be the strongest cluster for all spanish songs. In the first case, the clustering result is acceptable - none of the songs that fall into the cluster would really be misclassified; the only valid criticism is the lack of exactness. In the latter case, a portion of the lyrics that fall into that cluster are clearly misclassified because the cluster combines multiple different "real" topics.

When using high values for k, NMF tends to produce more specific clusters, most of which are quite interesting and useful. One of the drawbacks of this setting, however, is a tendency of NMF to find the same topics multiple times. Another noticeable side effect of high k values is that the important terms simply tend to be strong co-occurrents of the first term in the cluster, so the cluster represents not really a topic but rather contains the standard result of co-occurrence analysis.

[7] e.g., song X is represented as 10% topic A, 30% topic B and 60% topic C

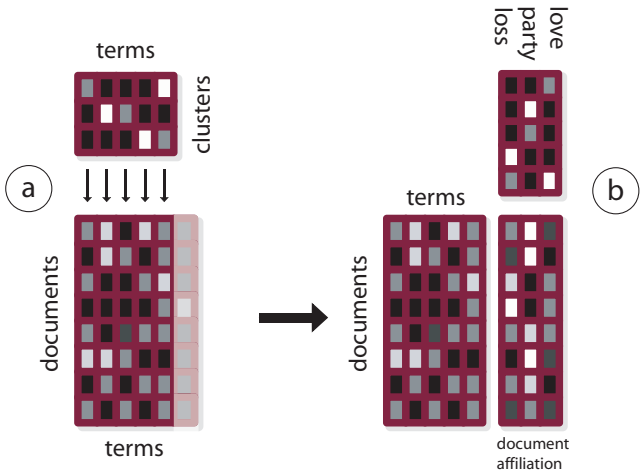

Figure 2. Computation of the documents' affiliation strength to the clusters. The manually assigned cluster labels in this example are 'love', 'party' and 'loss'.

After carefully weighing the benefits and drawbacks of both approaches, we decided to use a high value for the number of clusters in NMF. We conducted experiments for optimizing f_{max} and k, and as a result, we chose to set k to 60 (and f_{max} to ∞, as stated above).

3.5 Labeling

In order to make the clustering useful to an end user, the clusters need some kind of identifiers associated with them that hint at the nature of each cluster's content. Labeling a cluster was done manually by reading its most important terms and assigning one or more words that best summarised those terms. The exact procedure followed in labeling the clusters, and the results are described in Section 4.

3.6 Calculating Topic Affiliation

The degree of membership of a document in each cluster is computed using the orginal TDM weighted with term frequencies. This step is shown in Figure 2. First, only the columns (i.e. terms) that were used in the NMF are selected (ⓐ). The resulting matrix is multiplied by the transposed factor matrix H from the NMF that contains the term weights per cluster (ⓑ). Hence, for each cluster and each document, each term belonging to the document is multiplied by the weight of the term in the cluster and the sum over these products is regarded as the weight of the cluster for the document. After the calculation of this document affiliation matrix, its rows are normalized to a length of 1 in the euclidean norm.

3.7 Query Formulation and Document Retrieval

Retrieval of documents requires defining a query specifiying which clusters the documents should belong to. Resulting documents are ranked with respect to that query. A query vector is defined, assigning a weight to each cluster. The cosine similarity values between the rows of the document affiliation matrix and the query vector define the rank of all documents in the query result. The result is sorted in descending order; the first results are the most relevant ones.

4 ASSESSMENT OF LABELING

4.1 Setup

Stages 1-4 of the described algorithm were applied to our lyrics database with the settings $k = 60$ and f_{max} =∞, as explained above. The resulting 60 clusters were used as basis for a two-phased experiment that was inspired by the delphi method [1] which is a common method for obtaining opinion consensus from a group of experts.

In the first phase, test subjects were shown the most important terms [8] of each cluster and were asked to provide tags that summarise the terms.

In the second phase, the same word lists were shown to the same test subjects, but this time the task was to choose the best tags from those collected during the first phase, but not more than two. These tests were carried out with 6 subjects, all male between 20 and 32 years of age with strong background in computer science and little in music.

4.2 Evaluation Measure

The strength of agreement among test subjects cannot be measured after the first phase because they are completely free in their production of tags, so it is very unlikely that identical tags be used. In phase 2, when the subjects have to choose from the tags produced during phase 1, this is possible because everyone has to perform the very same task. For estimating the significance of the labeling outcome, we compute the probability of the actual result being attained by completely random behaviour on behalf of the subjects. The rationale is similar to that of methods for assessing intercoder agreement: The lower this probability is, the more evidence there is that the result is due to intelligible features of the data.

During phase 2, there was a given number of tags (m) associated with a given cluster [9]. If a person chose a tag at first position, we assigned a grade of 1 to that tag. If the person chose it for the second position, the grade was

[8] Those terms with a weight stronger than mean + std.dev. of all term weights in the cluster, sorted by weight. Not more than 30 terms were shown.

[9] There were at least 2 tags for each cluster, at most 10; mean tag count per cluster was 6.25.

2, all other tags were assigned grade 3. Thus, In the whole session, a $(n \times m)$ grading matrix was created containing the grades for all m tags created by all n test subjects.

The behaviour of the subjects was modeled as follows: For a given cluster, i.e., for a given m, a subject could choose one of the tags as the best tag with a probability of $P(first) = p_1$ and chose none with probability $P(none) = 1 - p_1$. If a tag was chosen best, each tag was equally probable to be chosen with a probability of $1/m$. Then, the person could pick another tag as second best with probability $P(second|first) = p_2$ and no second best tag with probability $P(nosecond|first) = 1 - p_2$. If a tag was chosen as second best, again, all tags were equally probable for this choice with probability $1/(m-1)$.

The model parameters p_1 and p_2 are computed based on the behaviour of the test subjects. p_1 is defined as the percentage of cases in which at least one tag was chosen, p_2 as the percentage of the former cases in which also a second tag was picked.

Consequently, given m, p_1 and p_2, the probability p(g) for a tag to get grade $g \in \{1, 2, 3\}$ is

$$p(1) = \frac{p_1 p_2}{m} + \frac{p_1(1-p_2)}{m} \quad (2)$$

$$p(2) = \frac{p_1 p_2}{m} \quad (3)$$

$$p(3) = \frac{p_1 p_2 (m-2)}{m} + \frac{p_1(1-p_2)(m-1)}{m} + 1 - p_1 \quad (4)$$

As the result of phase 2, we get a grading matrix for each cluster. The strength of agreement is assessed by computing how probable the column means of such a matrix are a priori. As each value in each column takes the values 1,2 or 3 with the probabilities explained above, the probability for a column to contain g_1 times the grade 1, g_2 times 2 and g_3 times 3 is

$$\frac{n!}{g_1! g_2! g_3!} p(1)^{g_1} p(2)^{g_2} p(3)^{g_3} \quad (5)$$

The likelihood of reaching a given column mean (i.e., average grade for a tag) is the sum of the probabilities of all grade combinations that result in the same or a better mean.

4.3 Discussion

The result of the assesment procedure is a number of tags for each cluster. The tags are associated with a grade and a value which indicates the likelihood for the grade to result from chance agreement among subjects. Table 1 shows the best-graded tag of each cluster, provided that the said likelihood is at most 10%. Due to this selection criterion, only 41 out of 60 tags are shown. As stated in Section 3.4, the same topic are found multiple times; this is reflected here in tags occurring more than once (e.g., *"dream"*, *"emotion"* or *"gangsta"*).

appearance	boys_and_girls	boys_and_girls	broken_hearted
clubbing	conflict	crime	dance
dance	depression	dream	dream
emotion	emotion	family	feelings
future	gangsta	gangsta	gangsta
gangsta	going_out	gospel	hard_times
hiphop	home	leave	listen
loneliness	loss	love	love
love	music	music	nature
party	sorrow	talk	weather
world			

Table 1. Winning tags at a significance level of 10%

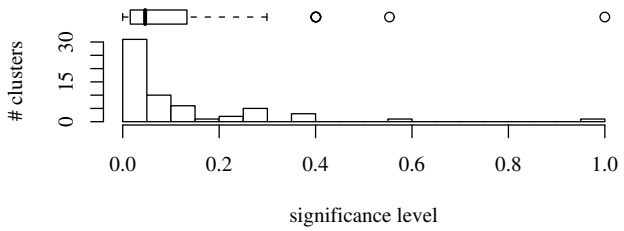

Figure 3. Number of clusters per significance level of the winning tag

Figure 3 shows the levels of significance for the most popular tag of each cluster in 5% steps. In 31 out of 60 clusters, the best tag was agreed on with a less than 5% likelihood of chance agreement. For another 10 clusters, the significance level was between 5 and 10%.

These results suggest that a reasonable portion of the clusters describes discernable topics and that they are reliably tagged.

5 CONCLUSION

The work at hand explains the structure and parametrization of an algorithm for the application of NMF to song lyrics. The focus is on showing the distinct stages of the algorithm and the considerations concerning the choice of parameter values for each stage. The most interesting choices for parameter values, in our view, are a) the high value for k (60 may still not be high enough) and b) the use of binary weighting prior to NMF clustering.

We also present an assessment of the clustering outcome indicating that most of the topic clusters resulting from our algorithm are useful for indexing our music collection. The procedure used for assessment is at the same time an integral part of the algorithm, the labeling stage, which has the convenient property that a statistically interpretable confidence value is calculated for each cluster so that it can be rejected or accepted for use in the subsequent stages.

6 FUTURE WORK

We plan to integrate the proposed method into an exising MIR system [10]. Only then the effects of certain parameter settings can be evaluated effectively. We also plan on evaluating our system in a user study with more test subjects than in the work at hand so as to produce statistically significant results. In addition to that, the question arises how similar clusters can be combined or hierarchically arranged for display to end users. Another interesting route of inquiry is hierarchical or iterative NMF clustering with the goal of finding less frequent topics. Taking the whole approach one step further, using NMF may turn out to be only one approach among many for creating weighted word lists (i.e., clusters) for indexing song collections. We consider the use of term co-occurrences most promising for the semi-manual definition of such lists for topics that are known to be contained in the archive but are not found by clustering. Ultimately, this task could be done by end users, allowing them to define and fine-tune topic classifiers themselves.

7 ACKNOWLEDGEMENTS

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[10] `http://musicexplorer.researchstudio.at/`

A DISCRETE MIXTURE MODEL FOR CHORD LABELLING

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ABSTRACT

Chord labels for recorded audio are in high demand both as an end product used by musicologists and hobby musicians and as an input feature for music similarity applications. Many past algorithms for chord labelling are based on chromagrams, but distribution of energy in chroma frames is not well understood. Furthermore, non-chord notes complicate chord estimation. We present a new approach which uses as a basis a relatively simple chroma model to represent short-time sonorities derived from melody range and bass range chromagrams. A chord is then modelled as a mixture of these sonorities, or subchords. We prove the practicability of the model by implementing a hidden Markov model (HMM) for chord labelling, in which we use the discrete subchord features as observations. We model gamma-distributed chord durations by duplicate states in the HMM, a technique that had not been applied to chord labelling. We test the algorithm by five-fold cross-validation on a set of 175 hand-labelled songs performed by the Beatles. Accuracy figures compare very well with other state of the art approaches. We include accuracy specified by chord type as well as a measure of temporal coherence.

1 INTRODUCTION

While many of the musics of the world have developed complex melodic and rhythmic structures, Western music is the one that is most strongly based on harmony [3]. A large part of harmony can be expressed as chords. Chords can be theoretically defined as sets of simultaneously sounding notes, but in practice, including all sounded pitch classes would lead to inappropriate chord labelling, so non-chord notes are largely excluded from chord analysis. However, the question which of the notes are non-chord notes and which actually constitute a new harmony is a perceptual one, and answers can vary considerably between listeners. This has also been an issue for automatic chord analysers from symbolic data [16]. Flourishing chord exchange websites [1] prove the sustained interest in chord labels of existing music. However, good labels are very hard to find, arguably due to the tediousness of the hand-labelling process as well as the lack of expertise of many enthusiastic authors of transcriptions. While classical performances are generally based on a score or tight harmonic instructions which result in perceived chords, in Jazz and popular music chords are often used as a kind of recipe, which is then realised by musicians as actually played notes, sometimes rather freely and including a lot of non-chord notes. Our aim is to translate performed pop music audio back to the chord recipe it supposedly has been generated from (*lead sheet*), thereby imitating human perception of chords. A rich and reliable automatic extraction could serve as a basis for accurate human transcriptions from audio. It could further inform other music information retrieval applications, e.g. music similarity. The most successful past efforts at chord labelling have been based on an audio feature called the *chromagram*. A chroma frame, also called pitch class profile (PCP), is a 12-dimensional real vector in which each element represents the energy of one pitch class present in a short segment (frame) of an audio recording. The matrix of the chroma frame columns is hence called chromagram. In 1999, Fujishima [5] introduced the chroma feature to music computing. While being a relatively good representation of some of the harmonic content, it tends to be rather prone to noise inflicted by transients as well as passing/changing notes. Different models have been proposed to improve estimation, e.g. by tuning [6] and smoothing using hidden Markov models [2, 11]. All the algorithms mentioned use only a very limited chord vocabulary, consisting of no more than four chord types, in particular excluding silence (*no chord*) and dominant 7th chords. Also, we are not aware of any attemps to address chord fragmentation issues.

We present a novel approach to chord modelling that addresses some of the weaknesses of previous chord recognition algorithms. Inspired by word models in speech processing we present a chord mixture model that allows a chord to be composed of many different sonorities over time. We also take account of the particular importance of the bass note by calculating a separate bass chromagram and integrating it into the model. Chord fragmentation is reduced using a duration distribution model that better fits the actual chord duration distribution. These characteristics approximate theoretic descriptions of chord progressions better than

[1] e.g. http://www.chordie.com/

previous approaches have.

The rest of this paper is organised as follows. Section 2 explains the acoustical model we are using. Section 3 describes the chord and chord transition models that constitute the hierarchical hidden Markov model. Section 4 describes how training and testing procedures are implemented. The result section 5 reports accuracy figures. Additionally, we introduce a new scoring method. In section 6 we discuss problems and possible future developments.

2 ACOUSTIC MODEL

2.1 Melody and Bass Range Chromagrams

We use mono audio tracks at a sample rate of 44.1 kHz and downsample them to 11025 kHz after low-pass filtering. We calculate the short-time discrete Fourier transform for windows of 8192 samples ($\approx 0.74s$) multiplied by a Hamming window. The hop-size is 1024 samples ($\approx 0.09s$), which corresponds to an overlap of $7/8$ of a frame window. In order to map the Fourier transform at frame t to the log-frequency (pitch) domain magnitudes $Q_k(t)$ we use the constant Q transform code written by Harte and Sandler [6]. Constant Q bin frequencies are spaced $33\frac{1}{3}$ cents (a third of a semitone) apart, ranging from 110 Hz to 1760 Hz (four octaves), i.e. the k^{th} element of the constant Q transform Q_k corresponds to the frequency

$$2^{\frac{k-1}{36}} \cdot 110\,\text{Hz}, \tag{1}$$

where $k \in 1, \ldots, (4 \cdot 36)$. In much the same way as Peeters [15], we smooth the constant Q transform by a median filter in the time direction (5 frames, $\approx 0.5s$), which has the effect of attenuating transients and drum noise.

For every frame t we wrap the constant Q magnitudes $Q(t)$ to a chroma vector $\mathbf{y}^*(t)$ of 36 bins by simply summing over bins that are an octave apart,

$$y^*_j(t) = \sum_{i=1}^{4} |Q_{36\cdot(i-1)+j}(t)|, \quad j = 1, \ldots, 36. \tag{2}$$

Similar to Peeters [15], we use only the strongest of the three possible sets of 12 semitone bins, e.g. $(1, 4, 7, \ldots, 34)$, thus "tuning" the chromagram and normalise the chroma vector to sum to 1,

$$y_k(t) = y^*_{3k+\nu}(t) \Bigg/ \sum_{i=1}^{12} y^*_{3i+\nu}(t)\ , \tag{3}$$

where $\nu \in \{0, 1, 2\}$ indicates the subset chosen to maximise $\sum_t \sum_k y^*_{3k+\nu}(t)$. A similar procedure leads to the calculation of the bass range chromagrams. The frequency range is 55 Hz to 220 Hz. The number of constant Q bins per semitone is 1, not 3. We linearly attenuate the bins at the frequency range borders, mainly to prevent a note just above the bass frequency range from leaking into the bass range.

2.2 Data

Harte has provided chord transcriptions for 180 Beatles recordings [7], the entirety of the group's 12 studio albums. Some of the songs have ambiguous tuning and/or do not pertain to Western harmonic rules. We omit 5 of these songs [2]. In a classification step similar to the one described by Mauch et al. [13] we map all chords to the classes *major*, *minor*, *dominant*, *diminished*, *suspended*, and *no chord* (which account for more than 94% of the frames) as well as *other* for transcriptions that do not match any of the classes. We classify as *dominant* the so-called dominant seventh chords and others that feature a minor seventh. We exclude the chords in the *other* class from all further calculations. Hence, the set of chords has $n = 12 \times 6$ elements.

2.3 Subchord Model

We want to model the sonorities a chord is made up of mentioned in Section 1 and call them *subchords*. Given the data we have, it is convenient to take as set of subchords just the set of chords introduced in the previous paragraph, denoting them S_i, $i = 1, ..., n$. In this way, we have a heuristic that allows us to estimate chroma profiles for every subchord [3]; in fact, for every such subchord S_i we use the ground truth labels G_t to obtain all positive examples

$$\mathbf{Y}_i = \{\mathbf{y}_t : G_t = S_i\}$$

and calculate the maximum likelihood parameter estimates $\hat{\theta}_i$ of a Gaussian mixture with three mixture components by maximising the likelihood

$$\prod_{\mathbf{y} \in \mathbf{Y_i}} L(\theta_i|\mathbf{y}).$$

Parameters are estimated using a MATLAB implementation of the EM algorithm by Wong and Bouman [4] with the default initialisation method. From the estimates $\hat{\theta}_i$, we obtain a simple subchord score function

$$p(S_i|\mathbf{y}) = \frac{L(\hat{\theta}_i, \mathbf{y})}{\sum_j L(\hat{\theta}_j, \mathbf{y})} \tag{4}$$

and hence a subchord classification function

$$s(\mathbf{y}) := \underset{S_i}{\operatorname{argmax}}\, p(S_i|\mathbf{y}) \in \{S_1, ..., S_n\}. \tag{5}$$

These will be used in the model with no bass information.

[2] *Revolution 9* (collage), *Love You Too* (Sitar-based), *Wild Honey Pie* (tuning issues), *Lovely Rita* (tuning issues), *Within You Without You* (Sitar-based)

[3] We only fit one Gaussian mixture for each chord type (i.e. *major*, *minor*, *diminished*, *dominant*, *suspended*, and *no chord*) by rotating all the relevant chromagrams, see [15]).

[4] http://web.ics.purdue.edu/~wong17/gaussmix/gaussmix.html

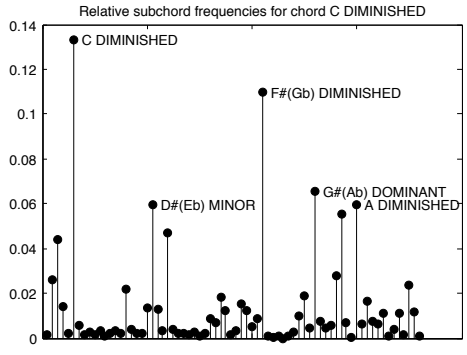

Figure 1. Example of subchord feature relative frequencies $b_{S|C}$ for the *diminished* chord. The five most frequent features are labelled. The subchord corresponding to C *diminished* most likely to be the best-fitting feature is indeed C *diminished*.

2.4 Subchord Model including Bass

In order to model the bass from the bass range chromagrams, every subchord S_i has a set $B_i \subset \{1, \ldots, 12\}$ of valid pitch classes coinciding with chord notes. The score for the bass range chromagram of subchord S_i at a bass chroma frame $\mathbf{y}^{\mathrm{b}}$ is the maximum value

$$p^{\mathrm{b}}(S_i|\mathbf{y}^{\mathrm{b}}) = \frac{\max_{j \in B_i} \left\{ y_j^{\mathrm{b}} \right\}}{\sum_k \max_{j \in B_k} \left\{ y_j^{\mathrm{b}} \right\}} \in [0, 1], \qquad (6)$$

the bass chromagram assumes in any of the pitch classes in B_i, b stands for *bass range*.

In order to obtain a model using both melody range and bass range information the two scores are combined to a single score

$$p(S_i|\mathbf{y}, \mathbf{y}^{\mathrm{b}}) = p^{\mathrm{b}}(S_i|\mathbf{y}^{\mathrm{b}}) \cdot p(S_i|\mathbf{y}). \qquad (7)$$

Analogous to Equation 5 we obtain a second subchord classification function

$$s(\mathbf{y}, \mathbf{y}^{\mathrm{b}}) := \operatorname*{argmax}_{S_i} p(S_i|\mathbf{y}, \mathbf{y}^{\mathrm{b}}) \in \{S_1, ..., S_n\}. \qquad (8)$$

2.5 Discrete HMM Observations

We discretise the chroma data $\mathbf{y}$ (and $\mathbf{y}^{\mathrm{b}}$) by assigning to each frame with chroma $\mathbf{y}$ the relative subchord, i.e. $s(\mathbf{y}, \mathbf{y}^{\mathrm{b}})$, or $s(\mathbf{y})$ depending on whether we want to consider the bass chromagrams or not. That means that in the HMM, the only information about a frame $\mathbf{y}$ we keep is which subchord fits best.

3 LANGUAGE MODEL

In analogy to speech processing the high-level processing in our model is called language modelling, although the language model we are employing is a hidden Markov model (HMM, see, e.g. [9]). Its structure can be described in terms of a chord model and a chord transition model.

3.1 Chord Model

The chord model represents one single chord over time. As we have argued above, a chord can generate a wealth of very different subchords. The HMM takes the categorical data $s(\mathbf{y}) \in \{S_1, \ldots, S_n\}$ as observations, which are estimations of the subchords. From these, we estimate the *chords*. The chords $C_1, \ldots, C_n$ take the same category names (C *major*, C# *major*,...) as subchords, but describe the perceptual concept rather than the sonority [5] . Given a chord C_i the off-line estimation of its emission probabilities consists of estimating the conditional probabilities

$$P(C_i|S_j),\ i, j \in 1, \ldots, n \qquad (9)$$

of the chord C_i conditional on the subchord being S_j. The maximum likelihood estimator is simply the relative conditional frequency

$$b_{i|k} = \frac{|\{t : s(\mathbf{y}_t) = S_i \wedge C_k = G_t\}|}{|\{t : C_k = G_t\}|}, \qquad (10)$$

where G_t is the ground truth label at t. These estimates are the (discrete) emission distribution in the hidden Markov model. A typical distribution can be seen in Figure 1, where C_k is a C *diminished* chord.

In hidden Markov models, state durations follow an exponential distribution, which has the undesirable property of assigning the majority of probability mass to short durations as is shown in Figure 2. The true distribution of chord durations is very different (solid steps), with no probability assigned to very short durations, and a lot between one and three seconds. To circumvent that problem we apply a variant of the technique used by Abdallah et al. [1] and model one chord by a left-to-right model of three hidden states with identical emission probabilities $b_{i|k}$. The chord duration distribution is thus a sum of three exponential random variables with parameter λ, i.e. it is gamma-distributed with shape parameter $k = 3$ and scale parameter λ. Hence, we can use the maximum likelihood estimator of the scale parameter λ of the gamma distribution with fixed k:

$$\hat{\lambda} = \frac{1}{k} \bar{d}_N, \qquad (11)$$

where $\bar{d}_N$ is the sample mean duration of chords. The obvious differences in fit between exponential and gamma modelling are shown in Figure 2. Self-transitions of the states in the left-to-right model within one chord will be assigned probabilities $1 - 1/\lambda$ (see also Figure 3).

[5] In fact, the subchords could well be other features, which arguably would have made the explanation a little less confusing.

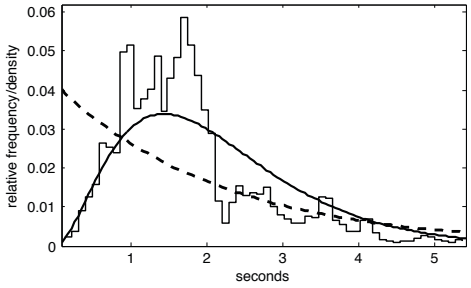

Figure 2. Chord duration histogram (solid steps) and fitted gamma density (solid curve) with parameters $\hat{\gamma}$ and $k = 3$ used in our model. Exponential density is dashed.

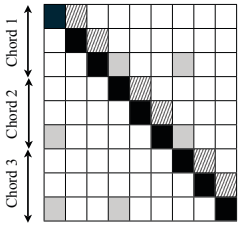

Figure 3. Non-ergodic transition matrix of a hypothetical model with only three chords. White areas correspond to zero probability. Self-transitions have probability $1 - 1/\hat{\lambda}$ (black), inner transitions in the chord model have probability $1/\hat{\lambda}$ (hatched), and chord transitions (grey) have probabilities estimated from symbolic data.

3.2 Chord Transition Model

We use a model that in linguistics is often referred to as a bigram model [9]. For our case we consider transition probabilities

$$P(C_{k_2}|C_{k_1}) \tag{12}$$

employing the estimates $\{a'_{k_1 k_2}\}$ derived from symbolic data smoothed by

$$a_{k_1 k_2} = a'_{k_1 k_2} + \max_{k_1, k_2}\{a'_{k_1 k_2}\}. \tag{13}$$

increasing probability mass for rarely seen chord progressions. The chord transition probabilities are symbolised by the grey fields in Figure 3. Similar smoothing techniques are often used in speech recognition in order not to underrepresent word bigrams that appear very rarely (or not at all) in the training data [12].

The initial state distribution of the hidden Markov model is set to uniform on the starting states of the chord, whereas we assign zero to the rest of the states.

4 IMPLEMENTATION

4.1 Model Training

We extract melody range and bass range chromagrams for all the songs in the Beatles collection as described in Section2.1. The four models that we test are as follows:

no bass, no duplicate states	no bass, duplicate states
bass, no duplicate states	bass, duplicate states

We divide the 175 hand-annotated songs into five sets, each spanning the whole 12 albums. For each of the four models we performe a five-fold cross-validation procedure by using one set in turn as a test set while the remaining four are used to train subchord, chord and chord transition models as described in sections 2.3 and 3.1.

4.2 Inference

For a given song from the respective test set, subchord features for all frames are calculated, thus obtaining a feature sequence $s(\mathbf{y}_t)$, $t \in T_{\text{song}}$, and the resulting emission probability matrix is

$$B_k(y_t) = b_{s(y_t)|k}, \tag{14}$$

where $b_{s(y_t)|k} = b_{i|k}$ with $i : S_i = s(y_t)$. In order to reduce the chord vocabulary for this particular song we perform a simple local chord search: B is convolved with a 30 frame long Gaussian window, and only those chords that assume the maximum in the convolution at least once are used. This procedure reduces the number of chords dramatically, from 72 to usually around 20, resulting in a significant performance increase. We use Kevin Murphy's implementation [6] of the Viterbi algorithm to decode the HMM by finding the most likely complete chord sequence for the whole song.

5 RESULTS

We calculate the accuracy for the set of chord classes. As we have six chord classes (or types), rather than two [11] or three [10] we decided to additionally provide results in which *major*, *dominant*, and *suspended* chords are merged. The calculation of accuracy is done by dividing summed duration of correctly annotated frames by the overall duration of the song collection. Similarly, in the case of one particular chord type (or song), this has been done by dividing the summed duration of correctly annotated frames of that chord type (or song) by the duration of all frames pertaining to that chord type (or song).

[6] http://www.cs.ubc.ca/~murphyk/Software/HMM/hmm.html

5.1 Song-Specific Accuracy

It is obvious that any chord extraction algorithm will not work equally well on all kinds of songs. Table 1 shows overall accuracy figures in both the merged and full evaluation mode for all four models. The models including bass information perform slightly better, though not significantly, with a mean chord recognition rate (averaged over songs) of 66.84% / 51.6% in the case of merged / full evaluation modes. The use of duplicate states has very little effect on the accuracy performance.

		without bass		with bass	
		std.	dupl.	std.	dupl.
merged	mean	64.74	64.96	66.46	**66.84**
	std. deviation	11.76	13.21	11.59	13.00
	max	86.35	89.15	86.99	88.81
full	mean	49.87	49.37	**51.60**	51.17
	std. deviation	13.70	14.85	13.93	15.65
	max	79.55	82.19	78.82	81.93

Table 1. Accuracy with respect to songs. Full and merged refer to the evaluation procedures explained in Section 5. The labels "without bass" and "with bass" denote if information from the bass chromagrams has been used or not, whereas "dupl." denotes the model in which the duplicated states have been used (see Section 3).

5.2 Total and Chord-specific Accuracy

Our top performance results (50.9 % for full evaluation mode, 65.9 % for merged evaluation mode) lie between the top scoring results of Lee and Slaney [11] (74 %) and Burgoyne et al. [4] (49 %). This is encouraging as we model more chord classes than Lee and Slaney [11], which decreases accuracy for either of the classes, and their figures refer to only the first two Beatles albums, which feature mainly *major* chords. Unfortunately, we cannot compare results on individual chords. We believe that such an overview is essential because some of the chord types appear so rarely that disregarding them will increase total accuracy, but delivers a less satisfying model from a human user perspective.

5.3 Fragmentation

For a human user of an automatic transcription not only the frame-wise overall correctness of the chord labels will be of importance, but also—among others properties—the level of fragmentation, which would ideally be similar to the one in the ground truth. As a measure for fragmentation we used the relative number of chord labels in the full evaluation mode. One can see in Table 3, the gamma duration modelling has been very successful in drastically reducing the fragmentation of the automatic chord transcription. This sheds a new light on the results as presented in Tables 1 and 2: the new duration modelling retains the level of accuracy but reduces fragmentation.

		without bass		with bass	
		std.	dupl.	std.	dupl.
merged	**total**	63.85	64.04	65.59	**65.91**
	major (merged)	70.31	72.04	72.58	**74.43**
	minor	48.57	43.93	**50.27**	45.63
	diminished	**14.63**	13.22	11.51	10.35
	no chord	**34.58**	27.42	25.59	19.48
full	**total**	49.17	48.64	**50.90**	50.37
	major	52.16	52.92	54.56	**55.45**
	minor	48.57	43.93	**50.27**	45.63
	dominant	44.88	46.42	**46.51**	46.42
	diminished	**14.63**	13.22	11.51	10.35
	suspended	**16.61**	11.04	13.22	9.04
	no chord	**34.58**	27.42	25.59	19.48

Table 2. Accuracy: Overall relative duration of correctly recognised chords, see also Table 1.

	without bass		with bass	
	std.	dupl.	std.	dupl.
fragmentation ratio	1.72	**1.12**	1.68	1.13

Table 3. Fragmentation.

6 DISCUSSION

6.1 Different Subchord Features

In the model presented in this paper, the subchord features coincide with the chords and emission distributions are discrete. This is not generally necessary, and one could well imagine trying out different sets of features, be they based on chromagrams or not. Advances in multi-pitch estima-

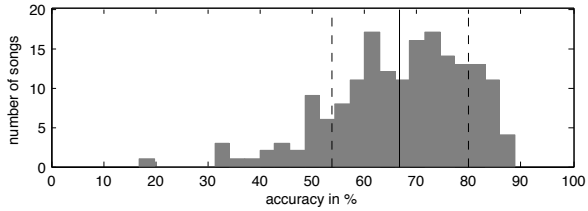

Figure 4. Histogram of recognition accuracy by song in the model using both gamma duration modelling and bass information, merged *major*, *minor*, and *suspended* chords, with mean and standard deviation markers.

tion [7] may make it feasible to use features more closely related to the notes played.

6.2 Hierarchical Levels and Training

While our duration modelling is a very simple form of hierarchical modelling, additional approaches are conceivable. Modelling song sections is promising because they could capture repetition, which is arguably the most characteristic parameter in music [8, p. 229]. Another option is key models, and a combination of the algorithms proposed by Noland and Sandler [14] and Lee and Slaney [11] is likely to improve recognition and enable key changes as part of the model. Such higher level models are needed to make on-line training of transition probabilities sensible as otherwise frequent transitions will be over-emphasised.

7 CONCLUSIONS

We have devised a new way of modelling chords, based on the frequency of *subchords*, chord-like sonorities that characterise a chord by their frequency of occurrence. A hidden Markov model based on this chord model has been implemented to label chords from audio with 6 chord classes (resulting in an overall vocabulary of 6×12 chords), while previous approaches never used more than four. The algorithm has shown competitive performance in five-fold cross-validation on 175 Beatles songs, the largest labelled data set available. In addition to the chord model we used a bass model, and more sophisticated state duration modelling. The use of the latter results in a reduction of the fragmentation in the automatic trancription while maintaining the level of accuracy. We believe that the novelties presented in this paper will be of use for future chord labelling algorithms, yet improvement in feature and model design provide plenty of room for improvement.

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[7] e.g. http://www.celemony.com/cms/index.php?id=dna

SPEEDING MELODY SEARCH WITH VANTAGE POINT TREES

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ABSTRACT

Melodic search engines let people find music in online collections by specifying the desired melody. Comparing the query melody to every item in a large database is prohibitively slow. If melodies can be placed in a metric space, search can be sped by comparing the query to a limited number of vantage melodies, rather than the entire database. We describe a simple melody metric that is customizable using a small number of example queries. This metric allows use of a generalized vantage point tree to organize the database. We show on a standard melodic database that the general vantage tree approach achieves superior search results for query-by-humming compared to an existing vantage point tree method. We then show this method can be used as a preprocessor to speed search for non-metric melodic comparison.

1. INTRODUCTION

Music is a popular category of online multimedia. Example collections include the millions of recordings in iTunes, emusic, and amazon.com. New melodic search engines let one search a music collection by specifying the desired melody. Typically, the query is entered by singing, playing a virtual piano keyboard or entering a symbolic representation, such as notes on a music staff. For an overview of recent approaches to melodic search, see [1, 2] or the results of the MIREX competitions [3].

Currently deployed systems compare the query melody to each melodic search key in the database. While this works for small collections, direct comparison to every database element becomes prohibitively slow as the size of the collection increases. Placing database melodies in a metric space lets one leverage a large body of research on efficiently finding objects [4].

Given a metric for melodies, we can pre-compute the distance of each melody in the database to a small set of melodies taken from the database, called vantage points. By comparing the query's distance to the vantage points with those of the database melodies, many melodies can be removed from consideration without need for direct comparison to the query, speeding search.

Recently, several authors [5-7] have organized melodic databases with vantage points and a metric. Typke et al [6] made an excellent first step, encoding melodies as a piecewise constant functions on the pitches of the chromatic scale and then applying a variant of Earth Mover Distance (EMD) [8]. EMD forms a pseudometric (non-identical elements may have a distance of 0). Their work did not, however, use a true metric, nor did it focus on learning to optimize the metric from real query data. They also did not explore the potential benefits of organizing vantage points in a structure.

Parker et al. [5] approximated an edit-distance-based metric for melodies based on a data set. They used a recursively organized structure, called a *vantage point tree* [4, 9], to prune the search space. Their system showed good results, but they did not learn a true metric, potentially compromising effectiveness. As with Typke et al., their pitch quanta are the notes on the chromatic scale. Such quantization can introduce error when used with pitch-tracker output from audio recordings (like sung queries) [10]. Also, they only tested a single tree structure, leaving open the possibility that other tree structures could improve performance.

This paper describes an un-quantized melodic encoding (Section 2) and a simple metric for melodic comparison (Section 3). We use this metric to organize a melodic database with a more general vantage point tree architecture than previous work has used (Section 4). The simplicity of the metric lets us tune its parameters (Section 6) from a small set of sung queries (Section 7). We explore the space of vantage point trees to find architectures that greatly improve search speed while sacrificing little in terms of search quality (Section 8). We then show a vantage point tree can be an effective preprocessor to speed search for a non-metric melodic comparison approach (Sections 5 and 9).

2. MELODIC ENCODING

In a typical melodic search system, the query is provided to the computer quantized to some musical alphabet (e.g. the note staff entry at http://musipedia.org) or as a sung example (http://midomi.com). In the case of a sung example, the melody is typically transcribed into a time-frequency representation where the fundamental frequency and amplitude of the audio is estimated at short fixed intervals (on the order of 10 milliseconds).

Our system accepts both music-staff entry and sung queries, so we use a representation scheme that is useful for both these cases. We encode all queries and all melodies in the database as sequences (strings) of note intervals. Each *note interval* is represented by a pair of values: the pitch interval (PI) between adjacent notes (measured in units of musical half-steps) and the log of the ratio between the length of a note and the length of the following note (LIR) [11] (where note lengths are inter-onset-intervals). We use note intervals because they are transposition invariant (melodies that differ only in key appear the same) and tempo invariant (melodies that differ only in tempo appear the same).

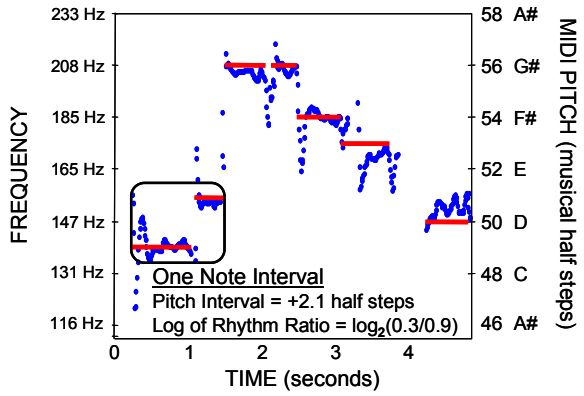

Figure 1. Encoding of a sung query

Figure 1 shows a transcription of a sung query by our system. Dots show the initial transcription. Horizontal lines show notes derived from the transcription. The rounded rectangle surrounding two notes indicates a single note interval derived from those two notes.

3. THE METRIC

A metric is a function $d(x,y)$ that defines the distance between elements x and y of a set. A metric must satisfy the following conditions: $d(x,y) \geq 0$ (non negativity); $d(x,y) = 0$ iff $x = y$; $d(x,y) = d(y,x)$ (symmetry); and $d(x,y) + d(y,z) \geq d(x,z)$ (triangle inequality).

While there are any number of functions that satisfy the requirements of a metric, most are unsuitable for meaningful melodic comparison. Since our focus is on speeding melodic search, a good metric for melodies must be quick to calculate and one where $d(X,Y)$ is small when a person would say melodies X and Y are similar and $d(X,Y)$ is large when a person would call X and Y dissimilar. We now describe a simple metric that has proven effective for melodic comparison.

We represent a melody X as a string of note intervals. We denote note intervals as lower case letters. Equation 1 defines a simple metric between note intervals x and y, with pitch intervals x_p and y_p and LIRs x_l and y_l.

$$d(x,y) = a|x_l - y_l| + b|x_p - y_p| \quad (1)$$

Here, a and b are non-negative values chosen to optimize performance on a set of example queries for a given database. This simple approach, when paired with the differential melodic encoding described in Section 2 (this encoding is crucial to the use of such a simple note metric), has been shown to produce comparable search performance to more complex distance measures, without the need to optimize large numbers of parameters [2].

The proof that this is a metric is straightforward: the absolute difference between two values is a metric. So is the weighted sum of two metrics. Thus, our distance measure is a metric on note intervals. Note this metric does not force quantization of the note values to a finite alphabet. Such quantization has been shown to introduce errors in melodic search using sung queries [10].

The distance between the query and each database element determines its ranking in search results. Of course, when searching in a melodic database, one is not comparing individual note intervals, but full melodies. To compare melodic strings, we use global edit distance. The edit distance between two strings is the cost of the least expensive way of transforming one string into the other. Here, transformation cost depends on a comparison function for the individual string elements. We have a fixed insertion/deletion cost of one, effectively forcing the other parameters to be in these units. If the comparison function for string elements is a metric (like Equation 1) then edit distance is also a metric. For a more in depth description and proof, see [12, 13].

4. VANTAGE POINT TREES

We now give an overview of the vantage point tree algorithm [9] for speeding search. We organize a database by choosing v elements at random from the database. These elements are the vantage points. Each element in the database has its distance measured to each vantage point using a metric. Around each vantage point, we partition the space into r concentric rings. We choose the size of these rings so that each contains nearly equal numbers of elements.

Given v vantage points with r rings per vantage point, this divides the metric space into r^v regions of intersection between rings, called branches. Each branch can be

uniquely identified by a *v*-tuple where the *i*th element specifies the ring for the *i*th vantage point. This tuple is the branchID. All database elements in the same branch are assigned the same branchID. BranchID values for database elements are computed prior to search.

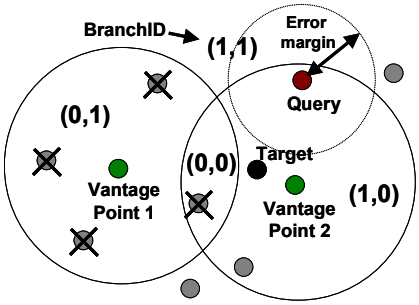

Figure 2. One level of a vantage point tree

Figure 2 shows one level of a vantage point tree with two vantage points and two rings per vantage points. Points in the graph indicate database elements (i.e. melodies). Each vantage point in this example has only two rings: a close inner region and a far outer region. This splits the space into four branches. In Figure 2, each branch ID is a binary duple.

To search the database, we find all branches that intersect the spherical region within an empirically-determined error margin around the query. This set of branches contains the only points we need to consider when searching the database. In Figure 2, this eliminates all melodies in branch (0,1) and (0,0), leaving the melodies in branches (1,0) and (1,1) as possible matches.

The error margin around the query represents our uncertainty about how well the metric corresponds to the choices a human would make in selecting the most similar melody. We are guaranteed to find all targets that are within the error margin of the query. However, the smaller the error margin, the fewer branches fall within it, speeding search. The larger the error margin, the less likely we are to accidentally remove the correct target from consideration.

Even a single-layer tree can eliminate a large number of database elements from consideration. The true power of this approach, however, lies in creating levels. Within any branch at level *l*, one can partition the space again by selecting vantage points within that branch and creating a set of branches at level *l*+1. This recursive structure is called a vantage point tree. We apply the algorithm at each level, until a maximum depth is reached or the leaves are too small to split further. At this depth, the remaining database elements are directly compared to the query to find the best match. The cost of searching the tree is negligible other than the comparisons to the vantage points; at worst, it requires a number of hash table lookups equal to the non-eliminated branches at each level. Given a poor metric (no branches are eliminated at any level) worst case performance is O(*n*). Typical performance is $O(\log_B(n))$. Log base *B* depends on the metric and tree structure. Different structures can have similar search quality, while varying in speed by an order of magnitude. This is shown in Section 8.

The vantage point tree used by Parker et al [5] is a special case with one vantage point per level (*v*=1) and two rings per vantage point (*r*=2). Similarly, the structure used by Typke et al [6, 7] can be realized with setting of *v* to the desired number of vantage points, *r* equal to the number of elements in the database, and only a single level (maximal tree depth *d*=1). Setting *v*=1, *r*=1 and *d*= 1 results in only one ring and all melodies must be directly compared to the query. This is linear search.

5. VANTAGE POINT TREE AS PREPROCESSOR

The metric in Section 3 is based on edit distance. Recent MIREX competitions show that recursive alignment (RA) generates a higher MRR than does the string-edit approach we can prove is a metric [14]. Unfortunately, RA is not a metric. Thus, we cannot directly apply vantage point trees to recursive alignment. We can, however, still use a vantage point tree preprocessor to speed search for a non-metric.

Let distance measure *M*(*a,b*) be a metric. Let *H*(*a,b*) be a slow non-metric we assume is ground-truth. If *M* and *H* agree within error bound *e*, we can build a vantage point tree with *M* and search with an error radius based on *e*. This precludes false negatives due to differences between *M* and *H*. The remaining database elements can be sorted using *H*. This lets us speed search by preprocessing with a vantage point tree built with *M*. We can progressively shrink the error bound to further speed search at the cost of increasing numbers of false negatives.

If we could find a vantage point tree able to put most queries in the same branches as their targets, it could be a useful preprocessor for any melodic search engine. Because each branch contains only a small portion of the database, the preprocessing could greatly speed up search, while still giving a high probability that the targets are within the search scope. Section 9 describes an experiment to find a good vantage point tree for this purpose.

6. DATA SET

Although there is nothing inherent in our approach that requires sung queries, we focus our experiments on the query-by-humming melodic search case, where queries are audio recordings of people humming or signing melodies. Our query set was drawn from the QBSH corpus [15] used in the 2006 MIREX comparison of query-by-humming melodic search systems [3]. We used 15 singers, each singing the same 15 songs from this dataset for a total of 225 queries. Our melody database

contained 2348 folk melodies from the Essen database (http://www.esac-data.org/) plus the 15 target melodies corresponding to the sung melodies in the query set, for a total of 2363 melodies. This database was chosen to emulate the database used in the 2006 MIREX competition.

The median melody length was 40 notes. Melodies were split into 10 note-interval subsequences with 3 note-intervals between the start of each subsequence and the search rank of the best subsequence was used as the rank of the melody. Melodies were broken into 11 subsequences, on average, resulting in 34,000 total database elements. While this is small compared to the millions of songs available in iTunes, the dataset used is a standard one used in the music search community. Copyright issues and the need to hand-vet melodies have limited the size of data sets used by this community.

7. TUNING THE METRIC

The metric from Section 3 may be tuned to favor pitch or rhythm similarity. Given the small number of parameters to be tuned, the model may be tuned using a relatively small set of examples. We learned parameters for a note segmenting preprocessor [10] and the note interval metric from Equation 1 using a simple genetic algorithm (GA) [16]. For tuning the metric, we selected five singers and used five of their songs for a total of 25 queries in our training set. Our testing set consisted of all 15 songs from each of the remaining 10 singers (150 queries). The GA used fitness proportional reproduction and single-point crossover. Each individual in the population was one set of segmentation and metric parameter values. To test the fitness of a set of parameter values, we ran a search engine using those values on the database from Section 6. Fitness for the GA was determined by MRR (see Section 8) on the training set and final results were validated on the testing set. Once good values for the metric were found, these values were used consistently in the experiments reported in this paper.

8. COMPARING TREES

We are interested in effect of tree architecture on the performance of a vantage point tree. To explore the space of the parameters for the vantage point tree, we generated 1736 random four-tuples (*v,r,d,e*). These represent the number of vantage points per level (*v*), rings per vantage point (*r*), maximal tree depth (*d*), and radius of the error margin around the query (*e*). Here *d* ranged from 1 to 20 levels, *v* ranged from 1 to 10, *r* ranged from 2 to 50, and *e* ranged from 0 to 10.

Trees were evaluated by the performance of a search-engine in finding correct targets for queries using the database from section 6. Given a vantage point tree, each query was processed. Once the final leaf branch was reached, all database melodies in the leaf were ranked for similarity to the query using the metric from Section 3. If the correct target was not in the same leaf as the query, we treated it as having been ranked 100th.

Call *c* the rank of the correct target for a query. Here, *c* = 1 indicates the correct target was ranked first. The *mean rank* for a trial is the average value for *c* over all queries in the trial. This measure can be sensitive to poorly ranking outliers. The *mean reciprocal rank* (MRR) is less sensitive to outliers. The reciprocal rank is 1/*c*. MRR (see Equation 2) is the mean of the reciprocal rank over all queries in a trial. Here, *Q* is the number of queries and *N* is the size of the database.

$$1 \geq MRR = \frac{\sum_{q=1}^{Q} \frac{1}{c_q}}{Q} \geq \frac{1}{N} \qquad (2)$$

If the system always ranks a correct answer first, the MRR will be 1. If the system gives random similarity values to targets, the MRR will be roughly log(N)/N. Chance MRR performance on our database is 0.03.

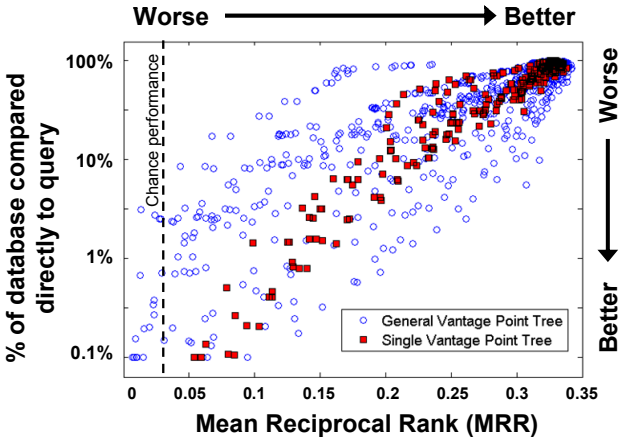

Figure 3. Performance of vantage point trees plus the metric as a stand-alone melodic search method.

Results for the 1736 vantage point trees are shown in Figure 3. Each point represents the performance of the search engine on the dataset from Section 6, given vantage tree parameter settings (*v,r,d,e*). Results show mean values for the 200 query melodies not used in training the metric. Parameter settings conforming to the simple vantage point tree method described in Parker et al [5] (Single Vantage Point Trees) are shown as solid squares. Trees with more than two rings per vantage point and/or more than one vantage point per level are shown as open circles (General Vantage Point Trees). The vertical dimension shows the proportion of the database compared directly to the query. Thus, a value of 1% means that 99% of the database was removed through use of the vantage

point tree, prior to the direct comparison of the final 1% to the query.

As Figure 3 shows, for every Single Vantage Point Tree, there is a General Vantage Point Tree that eliminates the same proportion of the database, while achieving a higher MRR. This indicates that using more complex vantage point tree structures can significantly improve search performance.

Figure 3 shows a general vantage point tree allows a string-alignment based method to achieve an MRR of 0.33 by comparing only 10% of the melodies in the database to the query melody. Full search of the database using the same method achieves an MRR of 0.346. This is an advance for edit-distance based search. Other search methods may also benefit from the metric and vantage point tree by using it as a front end. We now describe an experiment that explores this possibility.

9. FINDING A GOOD FRONT-END TREE

When a query is processed with a vantage point tree, we find which branches intersect the spherical region described by the error margin around the query. These branches are then divided into smaller branches at the next level. The tree is traversed until some depth limit is reached. At this point, we say the target song is still in our vantage point tree if at least one subsequence of the target melody is in one of the non-eliminated branches at the final level. The melodies in these branches may then be compared to the query by any desired search method.

By varying the value of error margin around the query (see Figure 2), we can change the number of the branches in the tree under consideration. Generally speaking, the larger the error margin is, the fewer branches will be eliminated from consideration. We wish to find an error margin that eliminates the largest portion of the database without removing the target melody.

To find a good tree architecture as a front-end for melodic search, we randomly generated 80 vantage point tree architectures by setting the number of vantage points and number of rings per point (v, r). For each pair of parameter settings, we then organized the database from Section 6 with a vantage point tree, recursively creating branches until reaching a level where the branches contain less than $2(v+r)$ melodies.

We then chose 150 queries and randomly selected 100 queries (the selecting set) from these 150 queries. We queried each tree with each query in the selecting set and counted the number times the correct target remained in the database after applying the vantage point tree. Call this t. We then calculated the percentage of database left after applying the tree, p. For each tree, we took the ratio t/p as a measure of the effectiveness of the tree. We then ranked the trees by this ratio and took the best one as the good vantage tree. This "good vantage tree" architecture was tested using the remaining 50 queries. We repeated this selecting and testing process three times using 3-fold cross validation. The best tree architecture from each of the three trials was *Trial1*: v=4 ,r=3, *Trial2:* v=3, r=5, and *Trial3:* v=3, r=5.

Figure 4 shows average performance over the three trials. Open circles with the dashed line show performance on the testing set. Solid circles with the solid line show performance on the selecting (training) set. The numeral by each point indicates the error margin around the query (see Section 4). The diagonal line shows the effect of shrinking the database by randomly removing a given percentage of the elements.

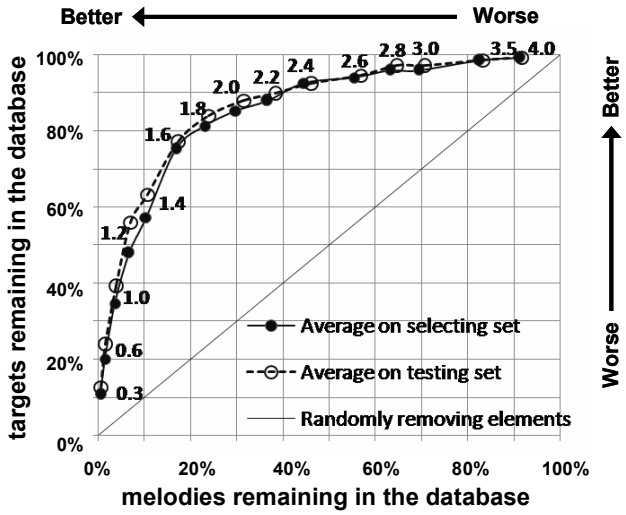

Figure 4. Performance of vantage point trees as a front-end for a melodic search system.

One may select the appropriate trade-off between search quality and search speed by choosing the appropriate error margin. For example, an error margin of 2.4 leaves 90% of the targets in the database, while eliminating 50% of the other database elements. An error margin of 1.7 retains 80% of the correct targets, but only 20% of the database, speeding search by a factor of five.

After narrowing the search scope by the vantage point tree preprocessor, one could use any melodic comparison approach for the remaining elements. This could make some time-consuming comparison methods practical in current computing-limited systems. If the errors made by the vantage point tree are uncorrelated with errors the subsequent search method makes, we could even improve MRR, since the tree could eliminate database elements that might confuse the other search method.

10. CONCLUSIONS

This paper describes a way to speed melodic database search. We apply a simple parameterized approach to building a metric for melodies. This simplicity lets us

learn good metric parameter values from a small set of queries. We use this metric to organize a database to speed search, employing a more general vantage-tree approach than existing work has used. This results in significantly improved search times while sacrificing very little search quality. Previous methods used to index a melodic database with vantage point trees are special cases in this framework. Any melody metric can be used with this approach to database organization. For example, one could choose one metric for skilled musicians and another for the general public.

One can also use a vantage point tree preprocessor to speed search for a non-metric. By choosing the correct parameters for the vantage point tree, one can balance search speed against the likelihood of eliminating the correct target from the database.

Optimal tree architecture for a particular application depends on the metric and the content of the database. In future work, we will explore the relationships between tree architecture, metric and dataset to improve our ability to select the right tree architecture.

11. ACKNOWLEDGEMENTS

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USING XQUERY ON MUSICXML DATABASES FOR MUSICOLOGICAL ANALYSIS

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ABSTRACT

MusicXML is a fairly recent XML-based file format for music scores, now supported by many score and audio editing software applications. Several online score library projects exist or are emerging, some of them using MusicXML as main format. When storing a large set of XML-encoded scores in an XML database, XQuery can be used to retrieve information from this database. We present some small practical examples of such large scale analysis, using the Wikifonia lead sheet database and the eXist XQuery engine. This shows the feasibility of automated musicological analysis on digital score libraries using the latest software tools. Bottom line: it's easy.

1 INTRODUCTION

MusicXML is an open file format developed by Recordare LLC [1]. Its development started around the year 2000 [2]. Some design decisions on the format are well explained in [3].

In 2002 a paper was presented on the XML conference [4] explaining how XQuery can be used to search for structures in a MusicXML document, but since then it has been very quiet on the front. In a 2007 poster, Viglianti [5] points out the possibilities of using XQuery for musicological analysis. Unfortunately, his text contains some errors - the most important one being the remark that XQuery would not support arrays. XQuery definitely does support arrays, it is even proved to be a Turing complete language [6]. Therefore has the same expressive power as languages like C++ or Java, and any computable function can be performed using XQuery.

In the meantime, XQuery 1.0 has made it to a W3C recommendation [7]. Galax [8] strives to be a reference implementation. eXist [9], Sedna [10], and Oracle's Berkeley DB XML [11] are native XML database management systems incorporating their own XQuery engines. All of those are open source and free. Both standards and software have thus matured.

Digital score libraries already exist, emanating from research (e.g. KernScores [12], based on HumDrum [13]) or the open source community (e.g. Mutopia [14], using Lilypond [15]). The Wikifonia project [16] uses MusicXML. It is a wiki-style collaborative environment for publishing lead sheets - i.e. reduced scores where arrangement details have been removed and only melody, lyrics and chord progression information is available. Wikifonia is at the moment of writing still relatively small. We worked with a database downloaded on April 1st 2008, which contained just over 200 songs.

To do automated musicological analysis, the HumDrum toolkit [13] is probably the most widely used today. It is very powerful, but it has its drawbacks. It works with input and output in its own plain text (ASCII) file format. Formatting output, converting to and from other formats, dynamically generating queries etc., require the creation of dedicated software or advanced scripts. To use it effectively, HumDrum requires a good deal of knowledge of UNIX style scripting, which is even more difficult to use on Windows platforms. On the other hand, XML formats are so generic that a huge set of software tools already exist to manipulate XML data in almost any way imaginable. Knowing that XQuery is Turing complete, we can state that it is theoretically possible to translate the whole HumDrum toolkit to XQuery.

There is not much software around that is capable of doing pattern analysis on MusicXML files: MelodicMatch [17] is at the moment of writing probably the most complete one. THoTH [18] also has some limited selection and extraction functionality based on the contents of a MusicXML file. Both software packages are closed source. Nevertheless, this kind of functionality is very desirable in music score software. For example, in regular PDF files, we are

not impressed any more by functionality that allows us to search for a text pattern, highlighting all occurrences in the text itself. But in score editors, searching for a music pattern and highlighting the results in the score, is a functionality that is only noticeable by its absence.

In this paper we give some practical examples that illustrate how XQuery can be used to easily and quickly search large collections of MusicXML scores for interesting data, and return the results formatted in any way desired. We will point out some interesting directions for future projects. By showing that only a few proven, stable, off-the-shelf software components are needed for advanced score analysis, we think that advanced search and information retrieval functionality can be incorporated in music score software very quickly.

2 USED TECHNOLOGIES

It is not our goal to provide an in-depth tutorial on MusicXML [1], XPath [19] or XQuery [7]. Numerous excellent books or online tutorials can be found on those subjects, and the websites of the respective projects contain valuable information. Nevertheless, to understand the examples in this paper, a small introduction is given in the next paragraphs.

MusicXML can encode scores in 2 ways: part-wise or time-wise. In a part-wise score, the file is roughly structured as follows: a score contains a number of parts, a part contains a number of measures, and a measure contains a number of notes. For a time-wise score, measure and part are switched in the hierarchy. An XSLT file (eXtensible Stylesheet Language Transformation) can be used to convert between the 2 formats, with a notable exception: multimetric music - where different parts have different time signatures - can only be properly encoded in part-wise scores. In practice, most MusicXML files today are part-wise encoded, and for the rest of this paper, we will assume part-wise file structure. Example files of MusicXML documents can be found on Recordare's website [1].

XPath is a language for retrieving information from an XML document. It is relatively easy to understand, and is used by XQuery to traverse XML documents. A complete reference of the latest W3C recommendation, version 2.0, can be found online [19]. Next to the document traversal functionality, XPath contains a wide range of operators for union and intersection, comparison, arithmetic, etc. A list of about 100 built-in functions completes the XPath language. Important to note is that an XPath query preserves document order.

XQuery extends XPath, making it more suitable for larger queries than XPath is. The most important extension is the so-called FLWOR (pronounce 'flower') construct - acronym for 'for, let, where, order by, return' - describing the typical form of an XQuery query. This is roughly analogous to the 'select, from, where' structure used in SQL (Structured Query Language). Results of XQuery FLWORs can be returned in any desired XML format, including XHTML, allowing immediate inclusion in a dynamic webpage. A second extension is the possibility to define functions yourself, functions which may be recursive, making XQuery Turing complete. Last but not least, *(: comments look like this :)*.

eXist [9] is an open source XML database management system, written in Java. It implements XPath 2.0, XQuery 1.0, and some other minor standards, recommendations and drafts. Several HTTP interfaces are provided, eXist can be easily deployed to run a whole database-driven website. In this paper, we used the latest stable release at the moment of writing, which is 1.2.0-rev7233. There was no specific reason to choose eXist over alternatives. It is a small project, easily set up on all kinds of platforms, with a very fast and complete XQuery engine and a simple graphical user interface. For a few queries to succeed, we needed to adapt the standard settings of eXist first: the maximum result size needed to be increased in the configuration file, and the maximum heap size of the Java virtual machine needed to be increased in the startup batch file.

3 QUERYING WIKIFONIA

In the next examples, we show some simple queries to retrieve basic information from the database. They can eventually be wrapped in a function to be used in more complex queries afterwards. To save the bandwidth of Wikifonia and speed up processing, we made a local copy of the database and worked from there. Wikifonia is at the moment of writing just a file repository, offering easy access to individual files with URLs that can be generated automatically. Iterating over all online documents and saving the results to a local database was done on April 1st, 2008, using the query in Listing 1, actually just a file dump.

Listing 1. Downloading the database. The URL of each document is composed using the *concat()* function that is built into XPath.

```
<wfdb> {
for $i in (0 to 1000)
let $s := concat("http://static.wikifonia.org/",
    $i,"/musicxml.xml")
return doc($s)
} </wfdb>
```

A better approach when working with large databases, is using 'collections', storing each file separately into the database. However, database manipulation functions are not part of the XQuery standard and can differ depending on the XQuery engine used. To keep things as uniform and clear as possible, we will keep working on a single file here, which we will call '*wfdb.xml*', that contains all songs. Since the database is relatively small, this is not problematic here: the file is only about 30 MB large and contains approximately 750000 lines of text.

3.1 Database contents and statistics

The number of note elements (this also includes rests, grace notes and cue notes) in the each song can be counted using the query in Listing 2. The largest song contains 1764 notes and the smallest only 5. Wrapping that code in the built-in *sum()* function and adapting it slightly informs us of a total of 50846 <note> tags in the database. Only the tag name needs to be adapted in the query to return the count of any tag in a MusicXML file, parts of a file, or the whole database.

Listing 2. Count the number of notes in each song. Element and attribute constructors are used in the return clause for mark-up.

```
for $i in doc("wfdb.xml")//score-partwise
let $j := count($i//note)
order by $j descending
return element{"song"}{attribute{"title"}
    {$i//movement-title/text()},
    attribute{"note-count"}{$j}}
```

Next, we can retrieve songs that meet certain criteria. The query in Listing 3 finds the song with the least number of notes and prints all notes that occur in that song. At the same time, it demonstrates the ability of XQuery to use nested queries. The query in Listing 4 returns a list of titles of those songs that have no rests in them.

Listing 3. Get the song with the least number of notes

```
for $i in doc("wfdb.xml")//score-partwise
let $j := $i//movement-title [ count($i//note) eq
  min( for $x in doc("wfdb.xml")//score-partwise
    return count($x//note)
  )]
return $j/..//note
```

Listing 4. Get the titles of songs with no rests

```
for $i in doc("wfdb.xml")//score-partwise
return $i[count($i//rest) eq 0]//movement-title
```

A music database will most likely contain some redundancies. The same song can be present in different editions, covered by other musicians, or there can be reductions for other instruments in separate files. These are just a few of the possibilities that cause the same song to be included in the database multiple times. Executing the query in Listing 5 checks only the titles, revealing that there are 3 songs called "Summertime" stored in the database and 9 other songs stored 2 times - as shown in Listing 6. Similarly, the query in Listing 7 allows us to find composers that have several compositions in the database. We found 12 "Traditional" songs in the database, 6 by Duke Ellington, 4 by A.L. Webber, etc.

Listing 5. Get the songtitles that are stored more than once in the database.

```
for $i in distinct-values(doc("wfdb.xml")//
    movement-title/text())
let $c := count(doc("wfdb.xml")//movement-title
    [text() = $i])
order by $c descending, $i
return ( element{"song"}{attribute{"count"}
    {$c}, $i} )[$c gt 1]
```

Listing 6. Part of the results of the query in Listing 5

```
<song count="3">Summertime</song>
<song count="2">Bernie's Tune</song>
<song count="2">Cherokee</song>
<song count="2">Could It Be Magic</song>
<song count="2">Everybody Hurts continued</song>
```

Listing 7. Sort authors that have more than 1 song in the database. The *lower-case()* function was used to iron out inconsistencies in capital letter usage.

```
let $multiset := doc("wfdb.xml")//creator
  [@type="composer"]/lower-case(text())
let $set := distinct-values($multiset)
for $i in $set
let $c := count(index-of($multiset, $i))
order by $c descending, $i
return ( element{"author"}{attribute{"count"}
  {$c}, $i} )[$c gt 1]
```

Eventually, we want to go looking for musically significant information. Using the same techniques as used above, Listing 8 returns the titles of all songs that are written in c minor. A more complex variant of this query, searching through all chords in a piece, is of special interest to people who are learning the guitar and only know a limited number of chords. They could then search for songs that only contain this limited set of chords, and thus find music they can already play completely despite a limited knowledge.

Listing 8. Find titles of all songs in c minor.

```
let $f := "-3"
let $m := "minor"
for $i in doc("wfdb.xml")//key
```

```
  [fifths/text() eq $f][mode/text() eq $m]
return $i/ancestor-or-self::score-partwise//
    movement-title
```

3.2 Database statistics

Using the built-in aggregation and arithmetic functions, one can generate very detailed statistics of the database contents. A first rudimentary example is presented in Listing 9. It iterates over the nominators and denominators of the time signatures that are present in the database, and generates all possible time signatures out of them. For each of these time signatures is calculated how many times it occurs in the database. This is converted to a percentage value and the results are ordered by that value. To keep the example simple, Listing 9 does not contain code to keep non-occurring time signatures from being generated. Instead, at the end the results having a count of 0 are removed from the list using the selection `[$c gt 0]`. A part of the results is presented in Listing 10. Note that time signatures can change in the middle of a song. Due to the query requesting all time signatures in the database wherever they occur, these time signature changes are also taken into account here.

Listing 9. Database statistics on time signatures.

```
let $t := count(doc("wfdb.xml")//time)
for $i in distinct-values(doc("wfdb.xml")//beats)
  for $j in distinct-values(doc("wfdb.xml")//
      beat-type)
  let $c := count(doc("wfdb.xml")//time
      [beats eq $i][beat-type eq $j])
  let $p := $c div $t * 100
order by $p descending
return element{"time"} {
  attribute{"beats"}{$i},
  attribute{"beat-type"}{$j},
  element{"count"}{$c},
  element{"percentage"}{$p}
} [$c gt 0]
```

Listing 10. Results of the query in Listing 9

```
<time beats="4" beat-type="4">
    <count>238</count>
    <percentage>49.6868475991649269</percentage>
</time>
<time beats="2" beat-type="4">
    <count>98</count>
    <percentage>20.4592901878914405</percentage>
</time>
<time beats="3" beat-type="4">
    <count>88</count>
    <percentage>18.37160751565762</percentage>
</time>
```

The same can be done for key signatures. In the code in Listing 11, all key signatures from 7 flats to 7 sharps are generated and both minor and major mode are considered. The first part of the result of this query is presented in Listing 12. Here too, note that key signatures can change in the middle of a song, and those key alterations are also present in the results. With some extra code this can be avoided if desired.

Listing 11. Database statistics on key signatures.

```
let $t := count(doc("wfdb.xml")//key)
for $i in (-7 to 7)
  for $j in ("minor","major")
  let $c := count(doc("wfdb.xml")//key
      [fifths eq string($i)][mode eq $j])
  let $p := $c div $t * 100
order by $p descending
return element{"key"} {
  attribute{"fifths"}{$i},
  attribute{"mode"}{$j},
  element{"count"}{$c},
  element{"percentage"}{$p}
} [$c gt 0]
```

Listing 12. Results of the query in Listing 11

```
<key fifths="0" mode="major">
    <count>64</count>
    <percentage>21.3333333333333333</percentage>
</key>
<key fifths="-1" mode="major">
    <count>39</count>
    <percentage>13</percentage>
</key>
<key fifths="-3" mode="major">
    <count>34</count>
    <percentage>11.3333333333333333</percentage>
</key>
```

A last query which is quite complex, is shown in Listing 13. It requests all lyrics of each song, sorts them by verse number, puts the syllables together if necessary so that they form words and sentences, and outputs them into what can be called a 'lyric library'. It can handle the special case that when there is only 1 verse present, it does not need to be numbered explicitly in MusicXML. Lyrics are only printed if there are any.

Listing 13. Extracting lyrics from each song and compiling them into a library.

```
<library> {
for $i in doc("wfdb.xml")//score-partwise
let $tit := $i//movement-title/text()
let $aut := $i//creator[@type="composer"]/text()
return
<song>{attribute{"title"}{$tit}}
  {attribute{"composer"}{$aut}}
{
  let $lyr := $i//lyric
  let $nrv := if(empty($lyr/@number)) then 1
    else xs:integer(max($lyr/@number))
  for $cur in (1 to $nrv)
  let $ver := $lyr[ if($nrv gt 1) then
```

```
    @number = $cur else true() ]/text
  let $s := string-join(
    for $syl in $ver return concat($syl/text(),
      if ($syl/../syllabic = ('begin','middle'))
      then '' else ' '), '')
  return
    <lyric>{attribute{"verse"}{$cur}}{$s}
      </lyric> [not(empty($lyr))]
} </song>
} </library>
```

3.3 Querying musical structure

Finding all occurrences of a single rhythmic or melodic motive, or a chord sequence, is a basic task when making a music theoretical analysis of a piece of music. In the next example, we work on the score of the 'Blue Danube' ('An der schönen blauen Donau'), the well-known waltz by Johann Strauss jr. The code in Listing 14 extracts all occurrences of the rhythmic pattern consisting of three notes of the same length followed by a note that is 3 times as long as the previous ones. This allows not only to find the pattern of 3 quarters followed by a dotted half note, but also faster or slower occurrences of the same rhythm - the comparison is based on relative duration values. This specific code example requires that all notes appear next to each other and in the same voice. The returned value indicates the position where the motive can be found. By simply removing the selection of a specific song, we can search the whole database for the specified motive.

Listing 14. Finding a rhytmic motive in a single song.

```
let $bd := doc("wfdb.xml")//score-partwise
    [movement-title = 'Blue Danube']
let $notes := $bd//note
for $i in (0 to count($notes))
let $s := subsequence($notes, $i, 4)
  (: now select those subsequences with the
    last note duration being 3 times that of
    the previous ones :)
let $durs := $s/duration
let $voic := distinct-values($s/voice)
where count($s/rest) = 0
and count($voic) = (0,1)
  (: '=' computes an intersection,
    whereas 'eq' compares single values :)
and $durs[1]/text() eq $durs[2]/text()
and $durs[2]/text() eq $durs[3]/text()
and number($durs[4]) eq number($durs[1]) * 3
return
<motive>
  <measure-start>{$s[1]/../@number/string()}
    </measure-start>
  <note-start>{index-of($s[1]/../note,$s[1])}
    </note-start>
</motive>
```

In [4] an XQuery function is presented that calculates MIDI pitch values of notes. The *where* clause in Listing 14 can be replaced by the code in Listing 15 which uses the aforementioned function. This example searches for the melodic motive of a large triad, followed by repetition of the last note. The triad can begin at any pitch, since only the intervals are taken into account.

Listing 15. Code Excerpt: Finding a melodic motive.

```
let $pits := $s/pitch      (: all pitch values :)
where count($s/rest) = 0      (: no rests :)
and count($voic) = (0,1)
and count($pits) = 4
and MidiNote($pits[1])+4 eq MidiNote($pits[2])
and MidiNote($pits[2])+3 eq MidiNote($pits[3])
and MidiNote($pits[3]) eq MidiNote($pits[4])
```

In [5] is pointed out that the generation of permutations of a sequence is of importance in certain music analysis tasks. A function can be written to generate all permutations of a sequence: XQuery allows defining recursive functions. But since permutation generation has a time complexity of order *O(n!)*, this is only usable on a very small scale. Also, when considering all possible permutations of a chord, transpositions need to be eliminated: this can be accomplished by calculating pitch values modulo 12.

4 FUTURE WORK

Musicological analysis is much more than just finding motives. The creation of an XQuery function database, a 'toolbox' for often used analysis tasks is one of the several practical projects that could (and, we think, needs) be undertaken to further automate large-scale analysis. An example of a practical tool that can be made available to the public is a web interface on top of the code presented here. This is not difficult to accomplish, since most XML database systems can be run as database back-end.

Of special interest are also regular expressions, the main power of the HumDrum toolkit [13]. The standard UNIX tools that it uses provide advanced regular expression processing on plain text files, but implementing this functionality in Xquery for XML files might be tricky - though it is theoretically possible.

Beyond merely statistical and structural analysis lies the domain of data mining. Here we try to isolate those patterns in the database that occur often, without having to define a specific pattern first. The Apriori algorithm [20] used to mine association rules in transactional databases has been translated to XQuery by Wan and Dobbie [21], and could be applied to a MusicXML database to search for structures that often appear together.

5 CONCLUSION

In this work we have shown the feasibility of using XQuery to mine a repository of music scores. We gave several examples of simple queries that can be run on a MusicXML database. They can form the basis for more complex queries. The strength of the XQuery language, the scalability of XML databases, the growing software support for the MusicXML format, combined with an increasing availability of digital scores, enable powerful musicological analysis with only a few lines of code. Awareness of the possibilities is an important factor in getting these XML-based technologies further developed. They allow musicologists, musicians, and the occasional music hobbyist to extract more significant information from an ever growing set of available scores in a very accessible way.

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TOWARDS QUANTITATIVE MEASURES OF EVALUATING SONG SEGMENTATION

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ABSTRACT

Automatic music structure analysis or song segmentation has immediate applications in the field of music information retrieval. Among these applications is active music navigation, automatic generation of audio summaries, automatic music analysis, etc. One of the important aspects of a song segmentation task is its evaluation. Commonly, that implies comparing the automatically estimated segmentation with a ground-truth, annotated by human experts. The automatic evaluation of segmentation algorithms provides the quantitative measure that reflects how well the estimated segmentation matches the annotated ground-truth. In this paper we present a novel evaluation measure based on information-theoretic conditional entropy. The principal advantage of the proposed approach lies in the applied normalization, which enables the comparison of the automatic evaluation results, obtained for songs with a different amount of states. We discuss and compare the evaluation scores commonly used for evaluating song segmentation at present. We provide several examples illustrating the behavior of different evaluation measures and weigh the benefits of the presented metric against the others.

1 INTRODUCTION

Automatic analysis of digital audio content has become an important research field in the last years. The rapid growth of music structure analysis also poses a question of sufficient evaluation of the proposed algorithms. Commonly the automatically estimated segmentation is compared to a ground-truth, provided by human experts. Music structure annotation is a challenging task even for the human experts. Its results might strongly vary depending on a particular application or even on the cultural background of the experts. For example, for popular western music the distinguishable and repeated parts could be "intro", "verse", "chorus", "bridge" and "outro".

There are two principal approaches to the music structure analysis, namely *sequence* and *state* representation [1]. In the present work we refer only to the state representation, i.e. we consider the music audio signal as a sequence of states. Each state is characterized by a consequent time region with a similar acoustical content and is assigned to a distinct label. Thus if the similar acoustical content appears once again in the music piece it is assigned to the same label. Especially for popular western music, the semantically distinguishable and repeated parts like "chorus" or "verse" generally have constant acoustical characteristics.

Evaluating song segmentation algorithms is not a trivial task. Possible solutions for the case of state representation have been already proposed by [2], [3] and [4]. According to the knowledge of the author, up to now there is no commonly established standard way of performing a segmentation evaluation. One of the key challenges is the unknown number of possible states which might vary depending on the song. Having just one state of interest (e.g. "chorus") [5] allows the use of *precision*, *recall* and *F-measure*, originally proposed in [6].

An ideal evaluation measure for song segmentation should possess the following properties:

- to provide the possibility to compare the results obtained by different authors andor for different algorithms;
- to be easily and intuitively understandable;
- to be insensitive to some particular properties varying for different songs, such as the number of states in the ground-truth segmentation.

It is not generally required that the estimated labels themselves should match the annotated ones, i.e. an additional stage of mapping the estimated labels into annotated ones is required. The architecture of the song segmentation system could imply that a given kind of mismatch between annotated and estimated segmentations is not considered to be a failure. For instance in [4] several annotated sequences are allowed to be mapped to a unique estimated one. Thus the evaluation measure should be able to treat this case correctly.

Representing a song as a sequence of possibly repeated states is in fact a classical clustering procedure. The same challenges appear while evaluating Speaker Clustering or Image Segmentation algorithms. As such similar evaluation measures can be used. Solomonoff et al. introduced the *purity* concept for Speaker Clustering evaluation [8], which was later extended by Ajmera et al. [9]. Another measure

was originally proposed by Huang and Dom [10] for evaluating Image Segmentation. Abdallah et al. [2] adapted it to the song segmentation task. The detailed explanation and the discussion of these evaluation measures are given in the next section.

In this paper we introduce a novel evaluation measure based on the information-theoretic conditional entropy. Generally, the number and the distribution of the states is different for each song in the test set. As such the evaluation scores, obtained for different songs are not directly comparable. The proposed evaluation measure is designed to overcome this challenge. Obviously, the more states there are in the song, the more "difficult" it is to get the true segmentation by chance, or randomly. If the song has just two states of equal duration, even a "random" segmentation with two states can lead to 50% of matching between the annotated and the estimated segmentations. This percentage decreases while the number of the states is increased . The absolute value of conditional entropy itself is strongly influenced by the number and by the distribution of the states in the annotated and the estimated segmentations. Therefore an additional normalization is required. We propose using two values of normalized conditional entropies as two independent scores corresponding to the over-segmentation (errors caused by false fragmentation) and to the under-segmentation (errors caused by false merge of the segments).

2 PREVIOUS WORK

2.1 Common notations

The usual form to represent the segmentation information is to provide the starting and the ending time of each segment and to associate each segment with a particular label of the state. The same information can also be represented as a sequence of numeric labels. Estimating the automatic segmentation usually includes feature extraction from the original acoustic signal which is accompanied by a time-domain windowing. On the later stages of the segmentation algorithm additional modeling or smoothing might be done leading to the change of time discretization unit. We can apply discretization with the same time unit to the annotated ground-truth segmentation. Thus, the segmentation information can also be represented via a discrete time sequence of numeric labels. In the present work A is a sequence of labels for each unit of time discretization (for each frame) for the annotated segmentation, and likewise E is a sequence of numerical labels for the estimated segmentation.

Additionally we denote:
N - total number of frames, equal for both annotated and estimated segmentations;
N_a - number of states in the annotated segmentation;
N_e - number of states in the estimated segmentation;
n_{ij} - number of frames that simultaneously belong to the state i in the annotated segmentation and to the state j in the estimated one;
n_i^a - total number of frames, that belong to the state i in the ground-truth segmentation;
n_j^e - total number of frames belonging to the state j in the automatic segmentation.

Note, that conventionally the subscript a and the running index i are assigned to the annotated segmentation. Correspondingly, the subscript e denotes the estimated segmentation and the running index j is associated to its states.

2.2 Pairwise precision, recall, and F-measure

One of the standard metrics for clustering evaluation is pairwise precision, recall and F-measure. This technique was used for Song Segmentation evaluation by Levy and Sandler [7]. Let M_a be a set of identically labeled pairs of frames in the ground-truth segmentation, i.e. pairs of frames that belong to the same state. Likewise let M_e be a set of identically labeled frames in the estimated segmenation. Then pairwise precision (P_p), pairwise recall (R_p), and pairwise F-measure (F_p) are defined as

$$P_p = \frac{|M_e \cap M_a|}{|M_e|} \tag{1}$$

$$R_p = \frac{|M_e \cap M_a|}{|M_a|} \tag{2}$$

$$F_p = \frac{2 \cdot P_p \cdot R_p}{P_p + R_p} \tag{3}$$

where $|\cdot|$ denotes the number of the corresponding pairs.

The pairwise precision shows the accuracy of the applied segmentation algorithm due to under-segmentation, while the pairwise recall indicates the over-segmentation accuracy.

2.3 Purity concept

The *purity concept* was first proposed in [8] for the evaluation of the Speaker Clustering. Solomonoff et al. introduced a quantity measure which describes "to what extent all the utterances from the cluster came from the same speaker" [8]. The score is designed to be maximal (equal to 1) if all utterances of the cluster come from the same speaker. It reaches the lowest level of $1/k$ if the utterances are evenly distributed between k speakers. Note, that the purity concept can be easily adapted to the case of song segmentation. In our case, the word 'speaker' corresponds to the states of the annotated segmentation, and 'cluster' is assigned to the states of the estimated one.

According to the above mentioned notations the cluster purity [8] is defined as

$$r_j^e = \sum_{i=1}^{N_a} n_{ij}^2 / (n_j^e)^2 \,. \tag{4}$$

Ajmera et al. [9] extended this evaluation measure using the *average cluster purity* (acp)

$$acp = \frac{1}{N}\sum_{j=1}^{N_e} r_j^e \cdot n_j^e \tag{5}$$

providing a weighted sum of single cluster purities. Furthermore they introduced the measures of *speaker purity* (r_i^a) and *average speaker purity* (asp)

$$r_i^a = \sum_{j=1}^{N_e} n_{ij}^2/(n_i^a)^2 \tag{6}$$

$$asp = \frac{1}{N}\sum_{i=1}^{N_a} r_i^a \cdot n_i^a \ . \tag{7}$$

The speaker purity estimates how well a speaker is limited to only one automatically estimated cluster. This measure is necessary to penalize a case of assigning each input utterance to a separate cluster. Ajmera et al. also suggested using a final score given as $K = \sqrt{asp \cdot acp}$.

In the case of song segmentation the above given asp measure corresponds to the over-segmentation (how well the segmentation is done due to the possible fragmentation mistakes) and likewise the acp depicts the accuracy due to the possible under-segmentation errors.

2.4 Concept of directional Hamming distance

An alternative approach to the segmentation evaluation was applied in [10] for the task of Image Segmentation. It was adapted and applied for the song segmentation task by Abdallah et al. [2].

Let T_a^i be a sequence of all frames forming a i-th state in the annotated ground-truth segmentation and T_e^j likewise a frame sequence belonging to the j-th state in the automatically estimated segmentation. The basic idea of the method is to establish the correspondence between the labels of two given segmentations by finding the sequence T_a^i with the maximum overlap for each sequence T_e^j of the estimated segmentation. The *directional Hamming distance* (d_{ae}) between annotated and estimated segmentations is given as

$$d_{ae} = \sum_{T_e^j} \sum_{T_a^k \neq T_a^i} |T_e^j \cap T_a^k| \tag{8}$$

where $|\cdot|$ denotes the duration of the overlapping parts. Normalizing the measure d_{ae} by the track length N gives a measure of the *missed boundaries*, $m = d_{ae}/N$. Accordingly the *inverse directional Hamming distance*

$$d_{ea} = \sum_{T_a^i} \sum_{T_e^l \neq T_e^j} |T_a^i \cap T_e^l| \tag{9}$$

and its normalization by the track length give a measure of the *segment fragmentation*, $f = d_{ea}/N$. The accuracy of the applied segmentation algorithm can be estimated using $1-f$ value for the under-segmentation and $1-m$ value for the over-segmentation.

We note that the evaluation measure identical to the $1-f$ score was independently proposed and applied in [4]. The author uses different mathematical expression leading to the identical numerical result. The over-segmentation mistakes were tolerated in [4], since several annotated states had been allowed to be mapped to an unique estimated one, due to an architecture of the segmentation system.

2.5 Mutual Information

In [2] Abdallah et al. proposed the information-theoretic measure (namely *mutual information*) for the segmentation evaluation. As we already mentioned in section 2.1 the segmentation information can also be represented via a discrete time sequence of numeric labels. In our notations, these are the sequences A and E for annotated and estimated segmentations correspondingly. The normalized 2D histogram of the mutual occurrence of the numeric labels in annotated and estimated segmentations can be treated as a *joint distribution* over the labels. Abdallah et al. suggested using the mutual information $I(A,E)$ as an evaluation score to measure the information in the class assignments. The mutual information is maximal when each state of the estimated segmentation maps to one and only one state of the ground-truth segmentation and it approaches zero value when the joint distribution over the labels is uniformly random. The biggest disadvantage of using the mutual information as an evaluation score is the unrestricted maximum value, which is dependent on the number of label classes and their distribution. As such the results, obtained for the songs with a different number of clusters or different distribution of the ground-truth labels, are incomparable.

3 PROPOSED METHOD

A possibility of using the conditional entropies as an evaluation score was first proposed in [2]. The authors noted that $H(A|E)$ measures the amount of ground-truth segmentation information that is *missing* in the estimated segmentation, while $H(E|A)$ measures the amount of the *spurious* information. Both $H(A|E)$ and $H(E|A)$ turn to zeros in the case of ideal segmentation. Some results of applying the conditional entropies as an evaluation score were presented in [11]. Likewise the mutual information score, discussed in section 2.5, the conditional entropies do not have a restricted maximum boundary. The higher the number of states in the segmentations (and the more uniformly these states are distributed) the higher conditional entropies.

In our method we overcome the disadvantages of using pure conditional entropy (namely, having the non-negative score with an unrestricted maximum value), and make the scores comparable for songs with a different number of clusters or different distribution of the state labels.

We denote the joint distribution (see section 2.5) of the labels (p_{ij}) and the marginal distributions for the annotated and estimated segmentations (p_i^a and p_j^e) correspondingly as

$$p_{ij} = \frac{n_{ij}}{\sum_{i=1}^{N_a}\sum_{j=1}^{N_e} n_{ij}} , \tag{10}$$

$$p_i^a = \frac{n_i^a}{\sum_{i=1}^{N_a}\sum_{j=1}^{N_e} n_{ij}} , \tag{11}$$

and

$$p_j^e = \frac{n_j^e}{\sum_{i=1}^{N_a}\sum_{j=1}^{N_e} n_{ij}} . \tag{12}$$

The conditional distributions are given respectively as

$$p_{ij}^{a|e} = \frac{n_{ij}}{n_j^e} \quad \text{and} \quad p_{ji}^{e|a} = \frac{n_{ij}}{n_i^a} .$$

Thus the conditional entropies can be written as

$$H(E|A) = -\sum_{i=1}^{N_a} p_i^a \sum_{j=1}^{N_e} p_{ji}^{e|a} \log_2 p_{ji}^{e|a} , \tag{13}$$

$$H(A|E) = -\sum_{j=1}^{N_e} p_j^e \sum_{i=1}^{N_a} p_{ij}^{a|e} \log_2 p_{ij}^{a|e} . \tag{14}$$

We propose to normalize the conditional entropies by the maximal conditional entropy for a given case. In the case of $H(E|A)$ we assume the annotated segmentation to be unchanged. The maximal conditional entropy $H(E|A)_{max}$ is achieved when the states of the automatically estimated segmentation are distributed uniformly through the states of the ground-truth segmentation. Note, that here we keep the original number of states in the estimated segmentation. In this case all the conditional distributions writes $p_{ij}^{a|e} = 1/N_e$ and the maximal conditional entropy writes

$$H(E|A)_{max} = -\sum_{i=1}^{N_a} p_i^a \sum_{j=1}^{N_e} \frac{1}{N_e} \log_2 \frac{1}{N_e} = \log_2 N_e .$$

Then we define the over-segmentation score (S_o) and the under-segmentation score (S_u) as

$$S_o = 1 - \frac{H(E|A)}{\log_2 N_e} \quad \text{and} \quad S_u = 1 - \frac{H(A|E)}{\log_2 N_a} .$$

Both scores are within the range of 0 and 1. They become maximal when the segmentations match each other perfectly, and approach a minimum value of 0 when the labels tend to be randomly chosen.

Score	Ex.1	Ex.2	Ex.3	Ex.4	Ex.5
P_p	1.00	0.53	1.00	0.53	0.55
R_p	1.00	1.00	0.23	0.62	0.55
F_p	1.00	0.70	0.38	0.57	0.55
asp	1.00	1.00	0.42	0.75	0.56
acp	1.00	0.50	1.00	0.54	0.56
K	1.00	0.71	0.65	0.64	0.56
$1-f$	1.00	1.00	0.42	0.75	0.67
$1-m$	1.00	0.58	1.00	0.58	0.67
$H(E\|A)$	0.00	0.00	1.69	0.50	0.92
$H(A\|E)$	0.00	1.09	0.00	0.94	0.92
$I(A,E)$	1.90	0.81	1.90	0.96	0.08
S_o	1.00	1.00	0.53	0.68	0.08
S_u	1.00	0.53	1.00	0.60	0.08

Table 1. Values of the different evaluation scores obtained for the examples 1-5

4 EXAMPLES

In this section we bring several examples to illustrate how the evaluation measures treat the typical errors appearing during song segmentation. For each of the examples we provide the schematic representation of the annotated and the estimated segmentations. The resulting evaluation scores for all examples are presented in Table 1.

Example 1

d	b	a	b	a	c	b	a	e	annot.
a	b	c	b	c	d	b	c	e	estim.

This example represents the ideal song segmentation for the typical song structure, consisting of the "chorus" (label a), "verse" (label b), "bridge" (label c), "intro" (d), and "outro" (e) in the annotated segmentation. In this case both errors of over-segmentation and under-segmentation are absent. As we see in the Table 1, all evaluation scores treat the case correctly. The mutual information reaches the maximal value of 1.90 bit, which is determined by the number and the distribution of the states for a given segmentation structure.

Example 2

d	b	a	b	a	c	b	a	e	annot.
a	b				a	b		a	estim.

Song segmentation algorithms are often realized by means of hierarchical agglomerative clustering. The challenging task is to terminate the agglomeration at the required level of the segmentation hierarchy, corresponding to the ground-truth reference segmentation. This example illustrates the

case, when the agglomeration algorithm is terminated too late, and the resulting states are too generalized. As such there is no over-segmentation errors, but under-segmentation scores are rather poor. Note, that the mutual information score (0.81 bit) is informative only in comparison to the maximal mutual information for a given ground-truth segmentation, obtained in Example 1.

Example 3

d	b	a		b	a		c	b	a		e	annot.
a	b	c	d	e	f	g	h	i	j	k	l	estim.

The opposite situation occurs if the automatically estimated segmentation tends to assign each input utterance to a separate state. That could happen if the agglomerative clustering is terminated at a very early stage. In this case there are no under-segmentation errors, but the over-segmentation scores are low.

Note, that the over-segmentation scores (asp, $1-f$ and S_o) and the under-segmentation scores (acp, $1-m$ and S_u) are in good correspondence for both Example 2 and Example 3. Contrastingly, the mutual information score for the Example 3 is close to the one obtained for the "ideal" segmentation in Example 1. The mutual information $I(A,E)$ can be written as

$$I(A,E) = H(E) - H(E|A) \tag{15}$$

where $H(E)$ is a marginal entropy of the automatically estimated segmentation. In this case a marginal entropy for 12 uniformly distributed states is really high (3.59 bit). As such even a high value of conditional entropy $H(E|A)$ (1.69 bit) cannot compensate it. The comparison of the Example 1 and Example 3 shows that a single score of mutual information is not directly applicable as an evaluation measure when the number of the states in the segmentations is changed.

Example 4

d	b	a		b	a		c	b	a		e	annot.
a	b		c	b		c		b		c	a	estim.

In most cases the estimated segmentation contains both over-segmentation and under-segmentation errors. For instance it happens when the borders of the segments are not determined correctly. In the presented example the second half of the "chorus" together with the "bridge" (label c in the estimated segmentation) is recognized as a distinct state, while the first half of the "chorus" is merged with the "verse" (label b in the estimated segmentation).

Clearly, the obtained estimated segmentation in Example 4 is not ideal, but it is obviously more appropriate than the estimated segmentation in Example 3. Note, that the measure K stays nearly unchanged, and even slightly decreases.

Example 5

b			a			b			a			annot.
a	b	a	b	a	b	a	b	a	b	a	b	estim.

The structure of the ground-truth segmentation in Examples 1-4 is typical (but not obligatory!) for Rock or Pop songs. This example depicts an annotated segmentation that could often appear e.g. for Jazz or Classical music pieces. As it was noted in section 1, the evaluation score should provide a possibility to compare the segmentation efficiency for different songs. In this case the estimated segmentation tends to be random. Obviously, this estimated segmentation is not of use in practical applications. In Table 1 we see, that the over-segmentation and the under-segmentation scores (S_o and S_u) are really low while the other evaluation measures (asp, acp, $1-f$, $1-m$) keep relatively high values. As such the values of $1-f$ and $1-m$ become comparable for the Examples 4 and 5, nevertheless that the estimated segmentation in Example 5 evidently carries less information about the corresponding ground-truth segmentation.

5 DISCUSSION

Analyzing the results listed in Table 1 we can summarize the following trends. First of all, the evaluation scores discussed in section 2 are strongly dependent on the amount and the distribution of the states in both annotated and estimated segmentations. Even within one musical genre these parameters differ for each song in the test set. Evaluating a song segmentation algorithm commonly implies calculating the evaluation scores for every entry of the test set and then providing the overall results. The latter becomes impossible if the evaluation score depends on the parameters that are different for every item of the test set.

We should mention that the higher the number of the states in the segmentations the more reliable the evaluation scores asp, acp, $1-f$ and $1-m$. Contrastingly, if the song consists only of a few states, even a "random" segmentation yields relatively high values.

The results show the benefits of treating the over-segmentation and the under-segmentation scores independently. If one of the scores is close to 1 and the other is relatively low, then we can assume that the automatic segmentation is performed on the other level of the segmentation hierarchy. In the worst, all states of the annotated segmentation are mapped into one state of the estimated segmentation, or vice versa each frame (time discretization unit) of the song forms a distinct state in the estimated segmentation. In the

latter cases the merging score K leads to the spurious results. For instance in the Examples 2 and 3 the score K indicates inconsistently high results. Therefore we believe using two independent scores S_o and S_u to be more appropriate for the case.

The comparison of the Examples 1 and 3 shows that the mutual information $I(A, E)$ is an unreliable evaluation score, especially if the number and the distribution of the states in both annotated and estimated segmentations are changed significantly.

The proposed evaluation score shows reliable results for all five presented examples. It is insensitive to changing the number of states in the segmentations and as such enables the comparison of automatic evaluation results, obtained from different songs. The scores can be loosely treated as an accuracy rate of the applied segmentation indicating the over-segmentation and under-segmentation errors. As a disadvantage of the S_o and S_u scores one can point out that in the strict sense these scores cannot be treated as the accuracy rates. The proposed normalization only restricts the boundaries of the scores and brings it within the range of 0 and 1. In point of fact the conditional entropies $H(E|A)$ and $H(A|E)$ are not a linear functions and thus a boundary restriction does not form the accuracy rate out of it. Therefore further investigations and a more complicated normalization scheme are needed.

6 CONCLUSION

In this paper we presented a novel approach to the song segmentation evaluation. The proposed evaluation score is based on the information-theoretic conditional entropies for comparing the estimated song segmentation with the one annotated by human experts (in the case of a state representation). Having applied the normalization scheme we formed the over-segmentation and the under-segmentation scores reflecting the accuracy rate of the obtained automatic segmentation given the annotated ground-truth. We compared our evaluation method to the commonly used approaches. By providing the illustrating examples we demonstrated the challenging points of the evaluation procedure and showed that the proposed over-segmentation and under-segmentation scores depicted more reliable results in comparison to the other evaluation measures.

By means of over-segmentation and under-segmentation scores one can compare the song segmentation efficiency obtained for the different songs, even when the number and the distribution of the states for these songs are various.

Since song segmentation is a particular case of classical clustering procedure, the proposed approach can be also applied for evaluating other clustering tasks like Image Segmentation or Speaker Clustering.

7 ACKNOWLEDGMENTS

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A TEXT RETRIEVAL APPROACH TO CONTENT-BASED AUDIO RETRIEVAL

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ABSTRACT

This paper presents a novel approach to robust, content-based retrieval of digital music. We formulate the hashing and retrieval problems analogously to that of text retrieval and leverage established results for this unique application. Accordingly, songs are represented as a "Bag-of-Audio-Words" and similarity calculations follow directly from the well-known Vector Space model [12]. We evaluate our system on a 4000 song data set to demonstrate its practical applicability, and evaluation shows our technique to be robust to a variety of signal distortions. Most interestingly, the system is capable of matching studio recordings to live recordings of the same song with high accuracy.

1 INTRODUCTION

Large digital music libraries are becoming commonplace on consumer computer systems, and with their growth our ability to automatically analyze and interpret their content has become increasingly important. The ability to find acoustically similar, or even duplicate, songs within a large audio database is a particularly important task with numerous potential applications. For example, an automated system similar to MusicBrainz [11] might organize a user's music collection by properly naming each file according to artist and song title. Another application could attempt to retrieve the artist and title of a song given a short clip recorded from a radio broadcast or perhaps even hummed into a microphone.

Due to the rich feature set of digital audio, a central task in this process is that of extracting a representative audio fingerprint that describes the acoustic content of each song. We hope to extract from each song a feature vector that is both highly discriminative between different songs and robust to common distortions that may be present in different copies of the same source song. With the multitude of compression formats and signal extraction processes, two copies of the same song can sound perceptually identical while having very different digital representations. Additionally, it is desirable for the audio fingerprint to compress the existing audio information into a much smaller representation, thus enabling efficient retrieval and requiring less storage than that of the initial data set.

In this paper, we present a novel hashing methodology that satisfies these constraints. We show that a technique based on methods for text retrieval performs well for the desired applications, and benefits from established research results in the area. Section 2 reviews existing work related to our application and section 3 details our application of the text retrieval techniques to content-based audio retrieval. Section 4 details our experimental evaluation of the proposed algorithm. Section 5 discusses practical implementation considerations and section 6 concludes with final remarks and suggestions for future work.

2 RELATED WORK

The problem of audio fingerprinting has been studied widely. In 2002, Haitsma and Kalker proposed a method for extracting audio fingerprints that they showed were robust to a variety of signal distortions. In addition, they outlined a database searching algorithm for locating a fingerprint most similar to a given target fingerprint [7]. One of the drawbacks of their system is the amount of memory required to store an audio fingerprint (approx. 100 KBytes for a 5 minute song). In addition, it was unclear whether or not their fingerprints could feasibly be used to match a studio recording to a live performance or a cover version (i.e. a performance of the original composition by another artist, possibly rearranged).

Existing work on cover song detection was presented for a competition at The Music Information Retrieval Evaluation eXchange (MIREX). In 2006, Dan Ellis' team from Columbia University won the competition by posting accuracy of about 60% using a method that computed similarities between songs by cross-correlating sequences of their so-called Chroma features [5]. Their similarity measure is equally applicable to the problem of matching a studio recording to a live performance. However, the high computational complexity of cross-correlating Chroma feature vector sequences does not make sense in an audio retrieval context.

We have not found previous research that directly applies text retrieval methods to the task of audio retrieval, but a similar approach has been taken for Object Recognition in images [13]. Further, Casey and Slaney [3] present a system

Figure 1. Block diagram for "Bag-of-Audio-Words" Representation

for discovering derivative audio works and describe a locality sensitive hashing procedure for matching similar songs that is derived from the "text shingles" method originally proposed in [2]. Finally, A vector quantization scheme similar to ours but using self-organizing maps is described in [14].

3 BAG-OF-AUDIO-WORDS REPRESENTATION

Our basic framework for extracting a song's Bag-of- Audio-Words representation is depicted in Figure 1. First, the song is converted from its original digital audio format into a 22.05 kHz, 16-bit, mono wav file. Next, the signal is divided into non-overlapping time segments and audio features are extracted from each segment. Then, a vector quantization (VQ) technique is used to map audio feature vectors to cluster numbers in the set $\{1, 2, \ldots, k\}$, where each cluster corresponds to what we refer to as an audio-word. Finally, each song is represented by a histogram of audio-word occurrences.

3.1 Audio Segmentation

In the segmentation process we extract non-overlapping 200 millisecond clips. We originally explored aligning audio segments to detected beats, which were extracted using the beat tracking algorithm proposed by Ellis and Poliner [5], but experimentally determined there to be little difference in system performance between the segmentation approaches. Additionally, uniform segmentation has the advantage of requiring less computational complexity.

3.2 Audio Feature Extraction

Several papers have characterized the suitability of numerous audio features for a variety of scenarios [5, 6, 10]. For our application, we chose the so-called normalized Chroma feature.

The Chroma feature is a 12-dimensional, real-valued vector that approximates an audio signal's strength at each musical note (e.g. A, A#, B, etc.), regardless of octave. Normalization of a Chroma is then performed by dividing by its vector norm.

We chose normalized Chroma features because it is very important to the performance of our audio-word histogram representations that the feature vectors for an audio segment and a distorted version are very similar. First, Chroma features are invariant to types of distortions that affect timbre because they only attempt to capture tonal information. Second, Chroma features are useful in detecting live/cover songs because they disregard information about octave, and are therefore somewhat invariant to certain differences between renditions or arrangements. Finally, the normalization of these Chroma features reduces the effects of a particular recording's loudness.

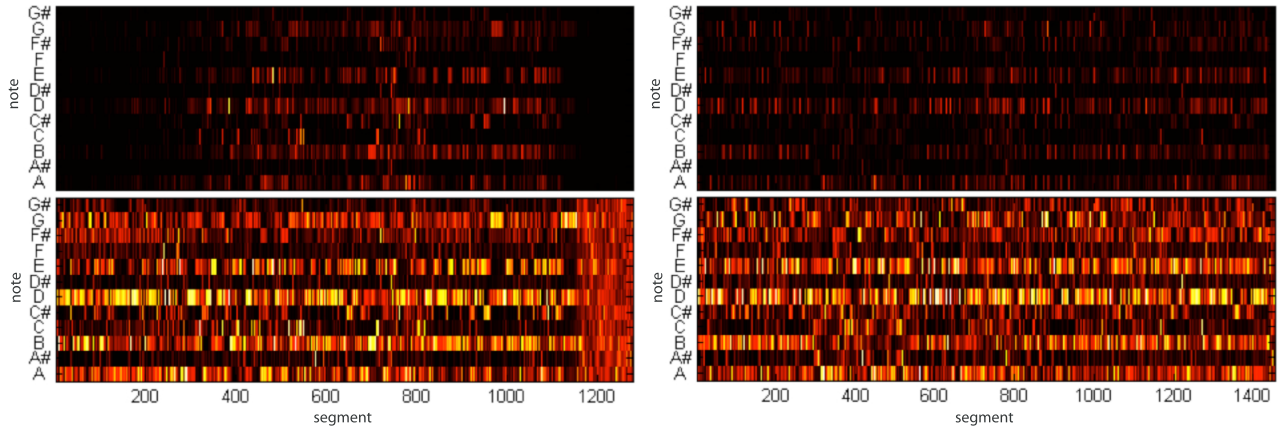
A song's chromagram is the sequence of the audio segment's Chroma features. Example chromagrams, with and without normalization, for the original and live performance recordings of the same song are depicted in Figure 2. We can see here the importance of normalization as the fading in volume from the original recording does not appear in the live performance. The normalization of Chroma vectors helps to eliminate the differences between the chromagrams of the two renditions.

3.3 Vector Quantization and Song-Level Histograms

Vector quantization primarily consists of performing clustering in the 12-dimensional Chroma space. The clustering process identifies k dense regions within a set of Chroma features extracted from our data set, which we collectively refer to as audio-words. Thereafter, when a Chroma feature is extracted from a song segment we calculate the nearest audio-word and consider the segment to be an occurrence of that audio-word. This quantization procedure forms the basis of matching distorted signals – song segments that sound very similar will have slightly different Chroma feature vectors but are expected to be assigned to the same audio-word.

For clustering we first collect a large sample of Chroma vectors from a variety of songs that are separate from our test set (approx. 100,000 vectors). We use K-Means to compute the k cluster centers, or audio-words.

The vector quantization procedure takes as input a song's sequence of Chroma vectors, and for each outputs one or more numbers in $\{1, 2, \ldots, k\}$ corresponding to the closest audio-word(s), as measured by Euclidean distance. Finally, the song x is mapped to a k-dimensional vector encoding the frequency of occurrence of each audio-word in the song: $\phi(x) = [f_1, f_2, \ldots, f_k]$ where f_i denotes the number of occurrences of the i-th audio-word in song x.



(a) Original Recording

(b) Live Recording

Figure 2. Chromagrams (with and without normalization) of the original and live performance of the same song.

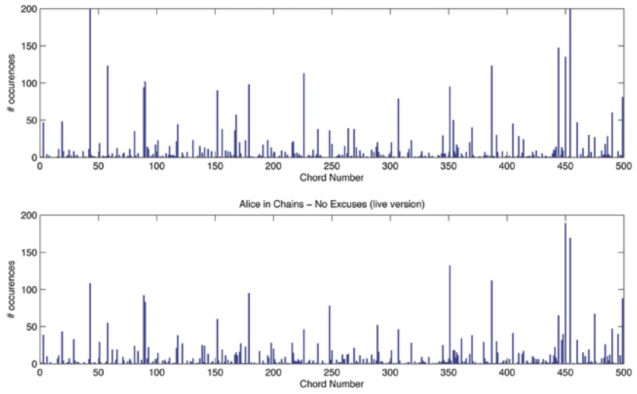


Figure 3. Audio-word histograms for the original and live recordings of Alice in Chains' "No Excuses"

By representing individual Chroma vectors by the cluster center to which they belong, we make equivalent in the histogram representation any segments that were musically very similar but may have had slightly different Chroma vectors. In terms of text retrieval, this is analogous to the stemming procedure often performed on the words of a text document [9].

3.4 Audio-Word Histogram Term Weighting

After forming the audio-word histograms we weight each of the k terms according to the term-frequency inverse document frequency (TF-IDF) scheme. We determine the term weightings $t_i, i \in \{1, 2, \ldots, k\}$, according to the following equation [1]:

$$t_i = \frac{n_{id}}{n_d} log \frac{N}{n_i} \tag{1}$$

Here n_{id} is the number of times the feature i occurs in song d, n_d is the total number of features in song d, N is the total number of songs in the database, and n_i is the number of songs in which feature i is present. The log term is larger for features that occur in few songs (rare features) and the leading term is large for features that occur many times in a given song. The n_d term serves to normalize the weights so that songs with many features can match songs with fewer features.

3.5 Calculating Song Similarity

Three different measures were considered and tested for computing the similarity between chord-histograms. We considered Cosine Similarity, Chi-Squared Similarity, and Euclidean distance (with normalization), given in equations (2), (3), and (4) respectively. In each of these equations, A and B represent the k-dimensional histogram vectors $\phi(a)$ and $\phi(b)$ of songs a and b.

$$sim_{ab} = \cos^{-1}\left(\frac{A \cdot B}{||A||\,||B||}\right) \tag{2}$$

$$sim_{ab} = \frac{1}{2}\sum_{i=1}^{k}\frac{(A_i - B_i)^2}{A_i + B_i} \tag{3}$$

$$sim_{ab} = \sqrt{\sum_{i=1}^{k}\left(\frac{A_i}{|A|} - \frac{B_i}{|B|}\right)^2} \tag{4}$$

For each of these similarity measures, a smaller value indicates a better song match. The cosine similarity measure helps when calculating the similarity between two songs without TF-IDF weightings because the dot product is normalized by the vector magnitudes, thus allowing songs of

different lengths to be similar. Analogously to a typical internet search engine query, given a query song or song clip, we use this similarity measure to return a ranked list of similar songs from the database. For our application we use the original WAV of the studio recording of a song as the query and expect the top results returned to be the distorted versions of the same song.

4 EXPERIMENTAL EVALUATION

We evaluated our system on a data set of 4000 songs drawn randomly from a variety of musical genres. A 4000 song data set under mp3 compression is roughly 40 gigabytes in size, which, if we use the size of current portable mp3 players as a guide, is a generous estimation of the size of a random user's music collection. In addition to the base 4000 tracks, we selected 60 additional tracks as query songs and quantify the system's ability to correctly retrieve distorted versions of the tracks from within the 4060 total song set. The idea here is to determine the discriminative power of the song-level feature vectors in a practical setting.

4.1 Vector Quantization

We experimented with several types of clustering algorithms besides K-Means, including hierarchical agglomerative clustering (HAC) and Gaussian mixture modeling (GMM) with expectation maximization. However, most of these other clustering algorithms involved greater computational complexity without improving the VQ and resulting song matching. We determined experimentally that $k = 500$ works well. Figure 4 depicts the Chroma vectors belonging to four different, and shows how some of the clusters resemble musical chords with two to three significant notes (lighter color indicates greater signal strength).

In addition, we determined experimentally that our matching performance could be improved by modifying the way in which we built our audio-word histograms. Instead of assigning a Chroma vector to the single closest cluster center, we assigned each one to the three closest centroids. We also tried some other soft assignment techniques. For example, instead of simply adding 1 to a Chroma's assigned histogram bins, we tried adding values weighted by how close the Chroma was to the cluster center. However, this approach actually hurt our results, especially in the case of identifying live/cover songs.

4.2 Robustness to Signal Distortion

The main experiment we performed in order to evaluate our Bag-of-Audio-Words song representation is as follows. First, we computed audio-word histograms for each of our 4060 songs. We then applied a signal distortion (overdrive, echo, etc.) using Adobe Audition to each of the 60 query songs,

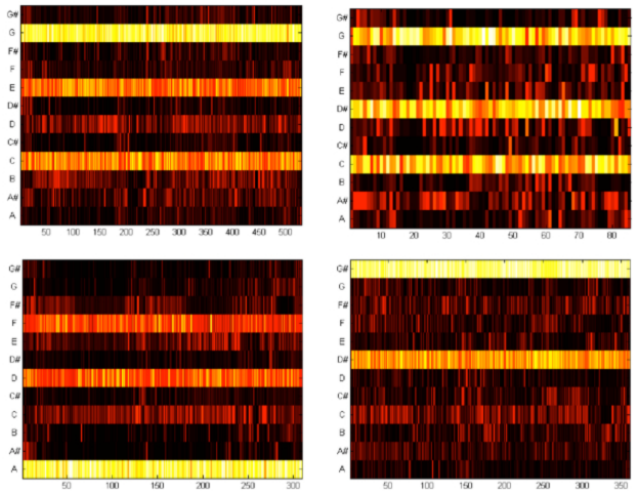


Figure 4. Four example clusters showing chord structure. (C major, C minor, D minor, D# power chord)

	Similarity Measure		
Distortion Type	Chi-Sq	Cosine	Euclidean
Overdrive	100%	100%	100%
Echo	98.3%	98.3%	98.3%
Reverb	100%	100%	100%
Speedup (1%)	98.3%	98.3%	98.3%
mp3 (32 Kbps)	100%	100%	100%
Live/Cover	67.8%	50.8%	50.8%

Table 1. Retrieval results

and computed the audio-word histograms for those distorted versions. Next, we computed our three similarity measures between each of the 4120 audio-word histograms. Finally, we calculated, for each similarity measure, the percentage at which the distorted query songs were most similar to the original query songs. Our results are outlined in Table 1.

4.3 Live/Cover Song Identification

In addition to facilitating matching between original and distorted recordings, our audio-word histogram representation was found to be useful in matching original studio recordings with live performances and cover songs to a lesser degree. To test this type of matching, we performed the same procedure as described in the previous section, but used a different set of 59 query songs for which we had live/cover versions (only a quarter of these were actually covers).

The results of this test are given in Table 1. It is somewhat surprising how good our live/cover song detection results were when compared to low-bitrate mp3 compression. In addition, it is interesting to note that our audio-word histograms were used to correctly match several cover songs.

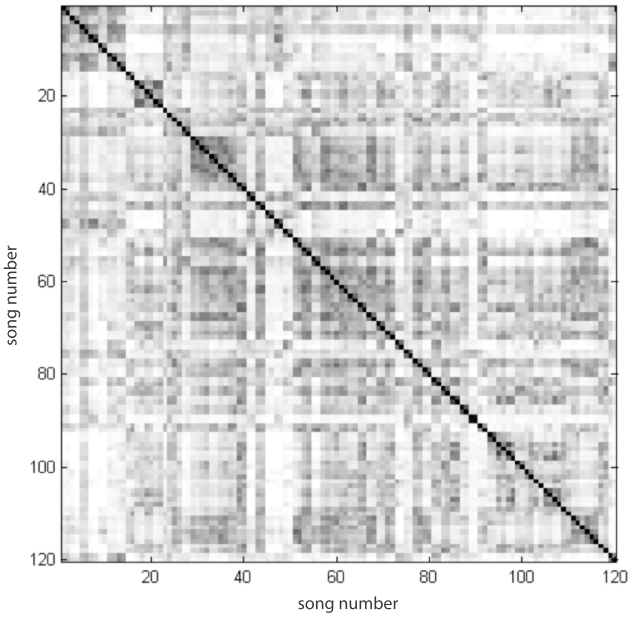


Figure 5. Similarity matrix for live performance retrieval.

For example, we correctly matched Stevie Ray Vaughn's cover version of "Little Wing" (6:49) to the original by Jimi Hendrix (2:24). The cover version is significantly longer and includes a much longer guitar solo.

Figure 5 shows a cosine similarity matrix for the query songs and their live/cover versions. In this matrix, a song at an odd index represents an original query song, and the song at the following even index is the live/cover version. If you look closely at this figure, you can see dark 2x2 squares along the diagonal, indicating that the original and live/cover versions have high similarity measures.

In addition to testing live/cover song retrieval from within a 4060 total song-set, we performed some other tests in the framework used by Ellis and Poliner [5]. Their procedure is as follows. First, they collected a set of approximately 80 original songs, and a separate set of cover versions of those 80. Then, their goal was to match a given song in the set of originals to its corresponding cover in the other set. For their 80 songs, they had a matching accuracy of around 60%. When we tested our matching algorithm on these same songs and in their framework, we only achieved an accuracy of 37.5% (using chi-squared similarity).

We then repeated the same procedure using our 59 songs, and the performance achieved by Ellis and Poliner's matching scheme was 70%, whereas our algorithm gave 90%. Our explanation for these surprising results is the following. Suppose we consider a cover song whose arrangement is such that every E minor chord (E, G, B) from the original version is replaced by an E power chord (E, B). In the case where song matching is based on chromagram cross-correlations, the matching between original and cover will be affected but not significantly. Under our scheme, these two chords would result in Chroma vectors that would map to different audio-word bins. Thus, our audio-word histogram representation can effectively be used to identify live song performances, but perform poorly on cover songs whose arrangements are significantly different from the original.

5 IMPLEMENTATION CONSIDERATIONS

Scalability was a central concern in the design of our proposed system. Here we discuss two implementation considerations – the Inverted Index and Locality Sensitive Hashing – that extend naturally to our application, and indicate that our algorithm will scale well to very large data sets.

5.1 Query Processing with an Inverted Index

The Inverted Index data structure is critical to the rapid processing speed of many text retrieval systems. This index contains all words in the text corpus and with each stores a list of every document in which that word is present. When performing a retrieval operation, the system looks at the inverted index to quickly retrieve a list of documents containing one or more of the words present in the query. When using the cosine similarity metric, only the words present in the query will affect the similarity measure, so documents not returned by a lookup in the inverted index can be safely ignored. In practice, this usually results a dramatic performance increase because the computationally expensive similarity metric must only be computed on a small subset of the entire database. Analogously, an Inverted Index for our application would contain each audio word and with each store a list of every song in which that word is present. This would be especially useful for an application in which only a short clip of a song is used to query the system.

5.2 Fast Search with Locality Senstive Hashing

The task of nearest-neighbor calculation in high-dimensional data can be efficiently implemented using Locality Sensitive Hashing (LSH). LSH is a hashing technique in which the probability that two objects are hashed to the same bin is proportional their similarity according to some metric. Hence, songs with very similar song-level histograms will likely be hashed to the same bin, allowing sub-linear determination of the nearest-neighbor of a given song within the data set, and therefore very rapid retrieval. LSH techniques exist for a number of similarity measures, including cosine similarity [4] and euclidean distance [8].

6 CONCLUSIONS

We have shown that a Bag-of-Audio-Words approach to audio retrieval can be both discriminative in a large data set

and robust to common signal distortions. We have also discussed numerous considerations for the practical application of our approach, addressing issues of scalability and efficiency. Excellent retrieval accuracy for a wide variety of distortions indicates that our approach will be useful for numerous applications. A natural extension of this work would be to add temporal information into the song-level feature vectors. Presently, the Bag-of-Audio-Words approach ignores all time-series information present in the initial song. Perhaps augmenting the song-level vectors to be a pyramid of audio word histograms, formed at different resolutions of song division, would lead to even better performance results.

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ENHANCED BLEEDTHROUGH CORRECTION FOR EARLY MUSIC DOCUMENTS WITH RECTO-VERSO REGISTRATION

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ABSTRACT

Ink bleedthrough is common problem in early music documents. Even when such bleedthrough does not pose problems for human perception, it can inhibit the performance of optical music recognition (OMR). One way to reduce the amount of bleedthrough is to take into account what is printed on the reverse of the page. In order to do so, the reverse of the page must be registered to match the front of the page on a pixel-by-pixel basis. This paper describes our approach to registering scanned early music scores as well as our modifications to two robust binarization approaches to take into account bleedthrough and the information available from the registration process. We determined that although the information from registration itself often makes little difference in recognition performance, other modifications to binarization algorithms for correcting bleedthrough can yield dramatic increases in OMR results.

1 MOTIVATION AND BACKGROUND

1.1 Fostering Interdisciplinary Research with OMR

"We stand at a moment of opportunity," opened Nicholas Cook at his invited talk for ISMIR 2005 in London. The opportunity is for historical musicologists and music information scientists to work together and revitalize the subdiscipline of computer-assisted empirical musicology [6]. This subdiscipline began in the 1960s [15], and although it has developed into a thriving discipline in several regions, it is largely moribund in North America, where a significant amount of other musicological research takes place. While Cook assigned musicologists a great deal of the responsibility for realizing this moment of interdisciplinary opportunity, he challenged researchers in music information retrieval to create large databases of the "highly reduced data"—e.g., scores—upon which musicological research relies. Such electronic databases are especially important for those older documents that are available in only a limited number of locations and for which archivists often restrict physical access, making it difficult to engage in large-scale comparative research.

Entering musical sources into such databases by hand is highly labor-intensive, which renders music digitization projects prohibitively costly for most institutions [3]. Optical music recognition (OMR), the musical analog to optical character recognition (OCR), is the most practical means by which to create such databases. The potential for optical recognition to transform research approaches has already been demonstrated by the recent explosion in the number of searchable electronic texts available for older books and journal articles, (e.g., JSTOR [1]). When treating historical documents, however, OMR and OCR systems struggle with various types of document degradation, including ink bleedthrough from the reverse side of the page [1]. Because these systems rely on the ability to distinguish foreground (ink) from background (paper), the darker the bleedthrough, the more likely it is that the bleedthrough will be classified as foreground and thereby corrupt the recognition process.

1.2 Binarization and Bleedthrough

Binarization is the common name for separating an image into its foreground and background. It is notoriously difficult to evaluate, leading most authors to resort to coarse subjective distinctions such as "better," "same," or "worse," e.g., [12]. As binarization is typically a preprocessing step in an automated image-processing pipeline with some other goal, e.g., OMR, one can use the evaluation metric for the ultimate goal, e.g., minimizing the amount of time necessary for a human editor to correct recognition errors, to evaluate the quality of the binarization algorithm. This paper uses a similar method to that described in our recently-published survey of binarization algorithms for music documents [4] to evaluate extensions to the algorithms we found to be best-performing.

Although there has been some previous work on binarization in the presence of bleedthrough that focuses specifically on historical documents [9, 10], a larger body of work

[1] http://www.jstor.org/

exists on the related problem of *showthrough* from the reverse side caused by the bright light used in photoreproduction (e.g., photocopying or scanning). Many approaches to showthrough (and bleedthrough) require that the recto and verso sides of each page be aligned, or *registered*, prior to the removal of showthrough [5, 8, 11, 20]. In addition to "blind" extensions to existing algorithms for the treatment of bleedthrough, i.e., those that do not take into account information from the reverse side of the page, we have developed our own technique for registering the recto and verso and extended the top-performing blind binarization algorithms for music to account for this information.

2 IMAGE PROCESSING

2.1 Binarization

2.1.1 KL Thresholding

In most cases, binarization algorithms seek to identify a global threshold for a given image. All pixels with gray levels below this threshold are classified as foreground and all pixels with gray levels above it are classified as background. The motivation for this simple classification technique, known as *thresholding*, is its computational speed.

Previous studies have demonstrated that thresholding approaches based on Shannon entropy are strong performers [4, 7]. For music documents in particular, [4] shows that a pair of related algorithms, [2] and [13], are optimal. These two algorithms consider images to be probability density functions, the so-called "monkey model," under which gray levels represent the probability that imaginary balls of luminance tossed by monkeys in a uniformly-distributed fashion would land on a particular pixel. Given a threshold T, the binarization of an image is represented not as zeros and ones but as the average gray levels of the foreground and background pixels in the original image: $\mu_0(T)$ and $\mu_1(T)$. Under this representation, the total luminance of the original image and the binarized image is identical. Thus, one can minimize the cross-entropy between the two images, or equivalently, minimize the Kullback-Leibler (KL) divergence, by minimizing

$$\theta(T) = \sum_{f(x,y)\leq T} f(x,y)\log\frac{f(x,y)}{\mu_0(T)} + \sum_{f(x,y)>T} f(x,y)\log\frac{f(x,y)}{\mu_1(T)} \quad (1)$$

where $f(x,y)$ represents the gray level of the image at pixel (x,y). This algorithm is the one proposed in [13] and is referred to in this paper as asymmetric KL thresholding. The algorithm proposed in [2] is very similar except that it minimizes the symmetrized form of the KL divergence:

$$\theta(T) = \sum_{f(x,y)\leq T} f(x,y)\log\frac{f(x,y)}{\mu_0(T)} + \sum_{f(x,y)\leq T} \mu_0(T)\log\frac{\mu_0(T)}{f(x,y)} + \sum_{f(x,y)>T} f(x,y)\log\frac{f(x,y)}{\mu_1(T)} + \sum_{f(x,y)>T} \mu_1(T)\log\frac{\mu_1(T)}{f(x,y)} \quad . \quad (2)$$

We call this technique symmetric KL thresholding. Both algorithms are analogous to Otsu's famous thresholding algorithm, [17], which also seeks to minimize the difference between the original image and a binarized representation of mean gray levels $\mu_0(T)$ and $\mu_1(T)$ but uses mean squared error as opposed to the KL divergence.

In an attempt to account for bleedthrough, which is usually lighter than foreground, we have added a third class to these algorithms by adding a second threshold U. Pixels with gray levels between T and U are considered to be bleedthrough and pixels with gray levels above U are considered to be true background. A new gray-level mean, $\mu_2(T,U)$ is computed for the bleedthrough pixels. For the asymmetric case, we minimize over both thresholds thus:

$$\theta(T,U) = \sum_{f(x,y)\leq T} f(x,y)\log\frac{f(x,y)}{\mu_0(T)} + \sum_{T<f(x,y)\leq U} f(x,y)\log\frac{f(x,y)}{\mu_2(T,U)} + \sum_{f(x,y)>U} f(x,y)\log\frac{f(x,y)}{\mu_1(U)} \quad . \quad (3)$$

We also implemented the analogous modification to (2). In this paper, these variants are referred to as three-class symmetric and three-class asymmetric KL thresholding. Samples of the output from these algorithms appear in Figure 1.

2.1.2 Gatos et al. 2004

Gatos et al. [10] is an adaptive technique for digital image binarization consisting of three stages: preprocessing, rough foreground estimation, background surface estimation, and final thresholding. In our earlier experiments [4], it was the only locally adaptive algorithm to perform well for music documents. Rough foreground estimation is achieved through the application of the adaptive thresholding method in [16], which tends to produce an over-complete (noisy) estimate of the foreground. A background surface is initially created from pixels classified as background during

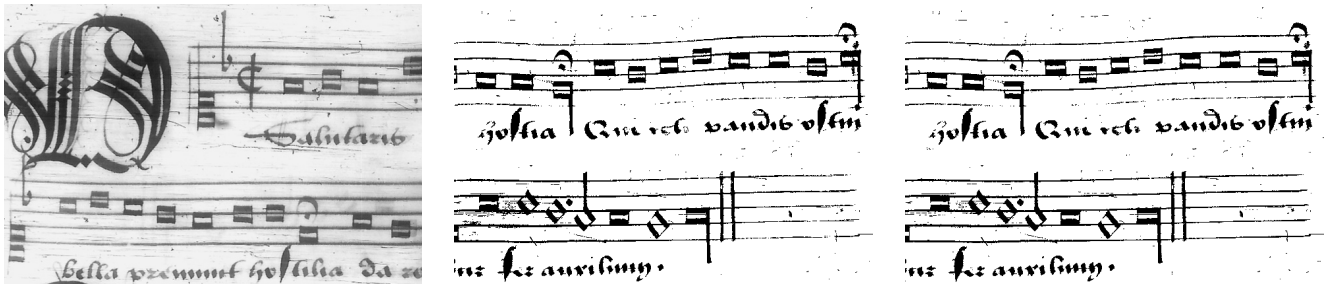
(a) Original recto image (b) Asymmetric KL: 2 classes (c) Symmetric KL: 2 classes

(d) Gatos et al. (21-px window) (e) Gatos et al. (31-px window) (f) Symmetric KL: 3 classes

Figure 1: Binarization output from selected algorithms on a page of the Occo Codex.

(a) Original recto image (b) Original verso image (c) Absolute difference between the original recto image and the registered, flipped verso image

Figure 2: Registration output for RISM M-0539



(a) Original verso image (b) Symmetric KL: recto-verso (c) Gatos et al. (31-px window): recto-verso

Figure 3: Binarization output from selected recto-verso algorithms on the same page of the Occo Codex.

this preliminary thresholding, and the values for the remaining pixels are derived by interpolating between the values of neighboring pixels. The final binarization is achieved by first calculating the distance between the background surface estimate with the original grayscale image: when an original pixel's gray level is greater than a threshold value derived from the corresponding background estimate, it is considered to be part of the foreground. Samples of the output from this algorithm also appear in Figure 1.

2.2 Registration

As the recto and verso side of a page are scanned separately, there is often some degree of misalignment and distortion between them. In order to take advantage of knowledge of the verso when correcting for bleedthrough, it is necessary to register the verso to the recto. We employ a feature-based registration technique, following from [8], that uses an affine transformation to minimize the total squared difference between pixels of recto and verso. In our experiments we constrained the affine transformation to avoid excessive translation or rotation. We also optimized the technique with the simulated-annealing approach described in [11]. Based on the success of the KL-based thresholding algorithms for music images, however, we replaced the sum-of-squares error metric used in [8] and [11] with symmetrized KL divergence.

Incorporating registration information into KL thresholding algorithms is similar to the three-class variant of the algorithms. Again, we minimize over two thresholds, T for the original image and U for the registered version of the reverse side. The bleedthrough class of pixels is restricted to those for which U classifies the pixel on the reverse side as foreground:

$$\begin{aligned}\theta(T,U) = &\sum_{f(x,y)\le T} f(x,y)\log\frac{f(x,y)}{\mu_0(T)} \\ &+ \sum_{f(x,y)>T\ \&\ g(x,y)\le U} f(x,y)\log\frac{f(x,y)}{\mu_2(T,U)} \\ &+ \sum_{f(x,y)>T\ \&\ g(x,y)>U} f(x,y)\log\frac{f(x,y)}{\mu_1(T,U)} \quad , \quad (4)\end{aligned}$$

where $g(x,y)$ represents the gray level of the image at the pixel from the registered reverse side corresponding to front-side pixel (x,y). Again, we also implemented the analogous modification to (2) for the symmetric form.

In order to take advantage of the registered verso in Gatos et al., we added an additional stage to the process between the foreground and background surface estimations. In this stage, both grayscale and binarized versions of the registered verso are used to guide removal of bleedthrough from the foreground estimation. If the pixel in the recto grayscale image is less than or equal to the corresponding pixel in the grayscale registered reverse, the corresponding pixel in the binary reverse is equal to 1, and the value of the pixel in the source grayscale image is less than a heuristically defined limit, then the value of the corresponding pixel in the foreground estimation is set to 0. Samples of this variant and a recto-verso variant from the KL thresholding family appear in Figure 1.

3 RESULTS

The data set we used for our experiments includes a total of 167 pages from three different books: two printed books of music by Luca Marenzio from 1581 and 1598, RISM M-0539 and RISM M-0582 [19], and one manuscript (handwritten book) compiled by the Alamire scriptorium in 1534, the Occo Codex. The 1581 Marenzio book contains significantly more bleedthrough than the 1598 book, and its landscape layout presented the registration process with additional challenges. The Occo Codex, where the amount of bleedthrough is comparable to the 1581 book but the layout is the same as the 1598 book, presented further challenges because most OMR systems, including our own (Aruspix [18]), are designed for printed scores rather than manuscripts.

The performance of recognition systems is traditionally presented in terms of recall and precision, but in the case of OMR, a more relevant statistic is the amount of time it will take a human to correct recognition errors. We present our results using a variant of our editing cost formula in [18], specifically

$$V = 1 - \left(\frac{{}^1\!/_4\, I + {}^1\!/_2\, S + D}{N}\right) , \quad (5)$$

where I is the number of musical symbols wrongly inserted by the OMR process, S is the number of musical symbols incorrectly classified by the OMR process, D is the number of musical symbols missed by the OMR process, and N is the total number of symbols on the page. The quantity V represents the savings in human editing time relative to transcribing a page by hand, i.e., $V = 0.75$ would mean that only a quarter of the human labor necessary for a music digitization project without OMR would be needed after introducing the OMR system. Perfect OMR would yield $V = 1.00$.

As the results in Table 1 show, all of these algorithms are good (as one would expect considering that top performers were selected for the experiments). Excepting the special cases of symmetric and asymmetric KL thresholding with three classes, which will be discussed below, the minimum average cost saving is 71 percent (asymmetric KL thresholding with two classes on the Occo Codex). For more accurate comparisons among algorithms, we also provide the odds multipliers (model coefficients transformed with the inverse link function) from a binomial regression model

Algorithm	RISM M-0539			RISM M-0582			Occo Codex		
	Rate	Odds multiplier		Rate	Odds multiplier		Rate	Odds multiplier	
Symmetric KL: 2 classes	0.88	0.56	(0.46, 0.67)	0.87	0.70	(0.65, 0.75)	0.75	0.81	(0.74, 0.87)
Symmetric KL: 3 classes	**0.93**	**1.00**	–	**0.90**	**1.00**	–	0.17	0.05	(0.05, 0.06)
Symmetric KL: recto-verso	0.89	0.66	(0.54, 0.81)	0.88	0.78	(0.71, 0.86))	0.76	0.86	(0.79, 0.93)
Asymmetric KL: 2 classes	0.88	0.57	(0.47, 0.69)	0.87	0.69	(0.64, 0.74))	0.71	0.65	(0.60, 0.70)
Asymmetric KL: 3 classes	0.93	0.99	(0.81, 1.22)	0.90	0.99	(0.92, 1.06)	0.11	0.03	(0.03, 0.04)
Asymmetric KL: recto-verso	0.89	0.63	(0.52, 0.77)	0.88	0.75	(0.69, 0.83)	0.76	0.85	(0.78, 0.92)
Gatos et al. (31-px window)	0.87	0.52	(0.43, 0.62)	0.89	0.86	(0.80, 0.92)	**0.79**	**1.00**	–
Gatos et al. (21-px window)	0.75	0.23	(0.19, 0.27)	0.88	0.81	(0.75, 0.87)	0.77	0.89	(0.82, 0.97)
Gatos et al. (31-px window): recto-verso	0.90	0.68	(0.56, 0.83)	0.89	0.88	(0.80, 0.97)	0.76	0.83	(0.77, 0.90)
Gatos et al. (21-px window): recto-verso	0.86	0.46	(0.38, 0.56)	0.88	0.81	(0.73, 0.89)	0.75	0.80	(0.74, 0.87)

Table 1: Average savings rates over manual transcription time for each algorithm and book. The corresponding odds multipliers (inverse-link-transformed regression coefficients), given here with their 95-percent-confidence intervals, take into account the variable difficulty of OMR for the different books and are scaled relative to a baseline of the best-performing algorithm for each book. Where the confidence intervals do not overlap, the differences are statistically significant.

over all pages in our experiment [4, 14], which take into account the fact that some books in the set are more difficult than others, a fact which is highly statistically significant ($p < 0+$). The baseline for the odds multipliers is set relative to the highest-performing algorithm for each book; thus they range from 0 to 1 and represent the proportion of the maximum recognition performance produced by any given algorithm. The 95-percent confidence intervals for the odds multipliers are also included. Where these intervals overlap, there is no statistically significant difference between the two algorithms.

The first point to note about the results is that although the symmetric KL thresholding family seems to perform slightly better than the asymmetric KL thresholding family on average, there is no statistically significant difference between them. For the printed music (RISM M-0539 and RISM M-0582), the three-class variants of KL thresholding are the clear winners. In the case of the Occo Codex, however, this pair of algorithms performs abysmally. The reason is clear from Figure 1f: in the presence of uneven lighting and variable ink density (e.g., from a quill pen rather than a printing press) these algorithms suffer a strong tendency to overclean. Curiously, our experiments showed that there was no statistically significant difference between the performance of the originally published version of the symmetric *two-class* KL algorithm [2], which contains an error in the normalization for mean foreground and background values, and the *three-class* variants of KL thresholding in our experiments. This unexpected similarity was true both for the Marenzio prints, for which these algorithms perform extremely well, and for the Occo Codex, for which these algorithms overclean and thus perform extremely poorly.

Although the global thresholds chosen by the two-class KL algorithms yield fair results, our adaptive algorithm set, based on [10], performs the best on the Occo Codex. In contrast to [4], for which the ideal window size was 21 pixels, a window size of 31 pixels works best on this manuscript. Looking at the performance of these algorithms for the print books, however, the preference for a 31-px window is not always statistically significant, which is in keeping with our earlier finding that choosing a single ideal window size for adaptive thresholding algorithms is impossible [4].

Incorporating information from the verso did not help the KL thresholding algorithms significantly. Although there are small improvements in the savings rate for all books in our data set, the confidence intervals for two-class KL thresholding and recto-verso KL thresholding all overlap considerably. For the Gatos-style algorithms, however, the registration does make a difference for RISM M-0539, the book with the most severe bleedthrough in our data set, although the improvement is only statistically significant for a window size of 21 pixels. Considering the computational resources necessary to register images, however, the improvements are unlikely to be worth the computational cost for all but the most bleedthrough-ridden OMR projects.

4 SUMMARY AND FUTURE WORK

On account of their computational speed and ease of implementation, we would recommend our new symmetric three-class KL thresholding algorithm for printed music and Gatos et al.'s algorithm for documents with uneven lighting or ink distribution, such as most manuscripts. Several techniques have also been presented in the literature for treating uneven lighting (e.g., [21]), and it is worth exploring whether correcting for uneven lighting before thresholding with a three-class KL algorithm would temper the overcleaning problem.

A very different technique for document binarization has also been published recently, based on independent component analysis (ICA) [22]. Early work with ICA for binarization was restricted to color images because multiple image channels are required for ICA to be meaningful. The more

recent work uses recto-verso registration to yield the two-channel minimum necessary for ICA. Because this algorithm is so markedly different from traditional approaches, we have not yet had an opportunity to compare it to our own algorithms. We expect that the presence of staves on both sides of the page will be too strong a violation of the independence assumption for ICA to work well for music images, but because the authors claim that the technique is more sensitive to the quality of registration than the content of the recto and verso pages, we are planning to explore their technique on our database of music images in the near future.

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HIGH-LEVEL AUDIO FEATURES: DISTRIBUTED EXTRACTION AND SIMILARITY SEARCH

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ABSTRACT

Today, automatic extraction of *high-level* audio features suffers from two main scalability issues. First, the extraction algorithms are very demanding in terms of memory and computation resources. Second, copyright laws prevent the audio files to be shared among computers, limiting the use of existing distributed computation frameworks and reducing the transparency of the methods evaluation process. The *iSound Music Warehouse* (iSoundMW), presented in this paper, is a framework to collect and query high-level audio features. It performs the feature extraction in a two-step process that allows distributed computations while respecting copyright laws. Using public computers, the extraction can be performed on large scale music collections. However, to be truly valuable, data management tools to search among the extracted features are needed. The iSoundMW enables similarity search among the collected high-level features and demonstrates its flexibility and efficiency by using a weighted combination of high-level features and constraints while showing good search performance results.

1 INTRODUCTION

Due to the proliferation of music on the Internet, many web portals proposing music recommendations have appeared. As of today, the recommendations they offer remain very limited: manual tagging has proved to be time consuming and often results in incompleteness, inaccuracy and inconsistency; automatic tagging systems based on web scanning or relying on millions of users are troubled, e.g., by mindless tag copying practices, thus blowing bag tags. Automatic extraction of music information is a very active topic addressed by the Music Information Retrieval (MIR) research community. Each year, the Music Information Retrieval Evaluation eXchange (MIREX) gives to researchers an opportunity to evaluate and compare new music extraction methods [9]. However, the MIREX evaluation process has proved to be resource consuming and slow despites attempts to address these scalability issues [4, 11]. So far, concerns with copyright issues have refrained the community to distribute the extraction among public computers as most algorithms require the audio material to be available in order to perform the feature extraction [1] . The features are therefore extracted from a relatively small music collection that narrows their generality and usefulness. Additionally, the feature extraction, being run by private computers on a private music collection, limits the transparency of the evaluation process. These limitations call for the development of a system able to extract meaningful, high-level audio features over large music collections. Such a system faces data management challenges. Noteworthily, the impressive amount of information generated requires an adapted search infrastructure to become truly valuable.

Our intention is to create a system able to cater for different types of features. Present literature mainly focuses on features that have either absolute or relative values, thus motivating the handling of both kinds of features. In this paper, the exact selection of the features is actually not as important as it is to demonstrate how extraction can be handled on public computers and enabling researchers to compare results obtained by using different algorithms and features.

The contributions of this paper are two-fold. First, we propose a framework for collecting high-level audio features (that were recently proposed by [5, 6, 7, 13]) over a large music collection of 41,446 songs. This is done by outsourcing the data extraction to remote client in a two-step feature extraction process: (1) dividing the audio information into short term segments of equal length and distributing them to various clients; and (2) sending the segment-based features gathered during step one to various clients to compute high-level features for the whole piece of music. Second, we propose a flexible and efficient similarity search approach, which uses a weighted combination of high-level features, to enable high-level queries (such as finding songs with a similar happy mood, or finding songs with a similar fast tempo). Additionally, to support the practical benefits of these contributions, we propose a short scenario illustrating the feature extraction and search abilities of the iSoundMW.

For the general public, the iSoundMW music offers recommendation without suffering from a "cold start", i.e., new artists avoid the penalties of not being well known, and new listeners obtain good music recommendation before being profiled. For researchers, the iSoundMW (1) offers flexi-

[1] https://mail.lis.uiuc.edu/pipermail/evalfest/2008-May/000765.html

ble search abilities; (2) enables visual comparison of both segment-based and aggregated high-level features; (3) provides a framework for large scale computations of features; and (4) gives good search performances.

The remainder of the paper is organized as follows. Related work is presented in Section 2. Section 3 offers an overview of the system, explains the process of collecting the high-level audio features, describes how similarity search with weighted coefficients is performed, and how the search can be further optimized by using range searches. Section 4 illustrates the similarity search on a concrete example. Section 5 concludes and presents future system improvements and research directions.

2 RELATED WORK

Research on distributed computing has received a lot of attention in diverse research communities. The Berkeley Open Infrastructure for Network Computing framework (BOINC) is a well-known middleware system in which the general public volunteers processing and storage resources to computing projects [1, 2] such as SETI@home [3]. However, in its current state, BOINC does not address copyright issues, does not feature flexible similarity search, and does not enable multiple steps processes, i.e., acquired results serve as input for other tasks. Closer to the MIR community, the On-demand Metadata Extraction Network system (OMEN) [15] distributes the feature extraction among *trusted nodes* rather than public computers. Furthermore, OMEN does not store the computed results, and, like BOINC, does not allow similarity search to be performed on the extracted features.

Audio similarity search is often supported by creating indexes [10]. Existing indexing techniques can be applied to index high dimensional musical feature representations. However, as a consequence of the subjective nature of musical perception, the triangular inequality property of the metric space is typically not preserved for similarity measures [14, 16]. Work on indexes for non-metric space is presented in the literature [12, 17]. Although the similarity function is non-metric, it remains confined in a pair of lower and upper bounds specifically constructed. Therefore, using these indexes would impose restrictions on the similarity values that would limit the flexibility of the iSoundMW.

3 SYSTEM DESCRIPTION

In this section, we present an overview of the system followed by a more detailed description of a two-step process for extracting high-level features. Later, we describe how flexible similarity searches are performed and how they are further optimized using user-specified constraints.

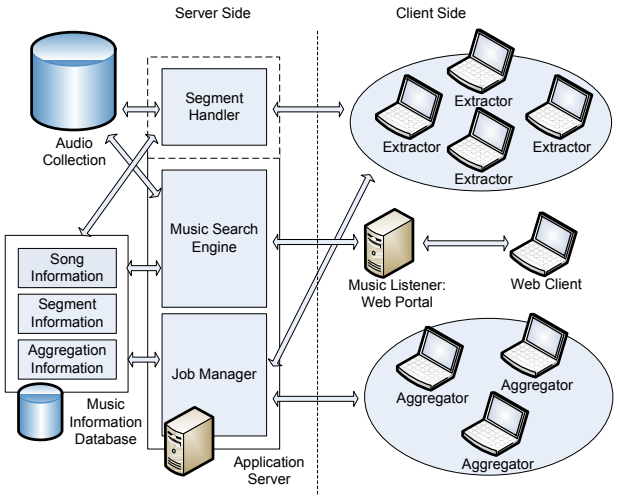

Figure 1. System architecture

3.1 iSoundMW Overview

The iSoundMW system has a client-server architecture, shown in Figure 1. The server side is composed of a central data repository and an application server that controls the extraction and similarity searches. The data repository stores all the audio collection in MP3 format and uniquely identifies each by a number. It also contains all the editorial information, e.g., the artist name, the album name, the band name, the song title, the year, the genre, and copyright license, and the physical information, e.g., the file size, the format, the bit rate, and the song length, that are stored in the music information database. Additionally, the Music Information Database holds all the extracted feature information.

The application server is composed of three distinct components. First, the *job manager* assigns extraction or aggregation tasks to the clients, collects the results, and prepares progress reports about each extraction process. Second, the *segment handler* splits the audio information into short term segments overlapping or non-overlapping of equal length, and makes them available to the clients they have been assigned. Future versions of the system will have multiple segment handlers, enabling multiple music collections to be analyzed without violating copyright for any of them. Third, the *music search engine* serves requests such as finding the most similar song to a given seed song with respect to a combination of features.

The client side is composed of three different types of clients. First, the *segment extractors* are receiving short term segments, and extracting their high-level features, e.g., the pitch, or the tempo. Second, the *segment aggregators* are performing computations based on the high-level features of the segments to produce other high-level features requiring features over different segments, e.g., the mood. Third, the *music listeners* perform similarity search queries on the extracted high-level features.

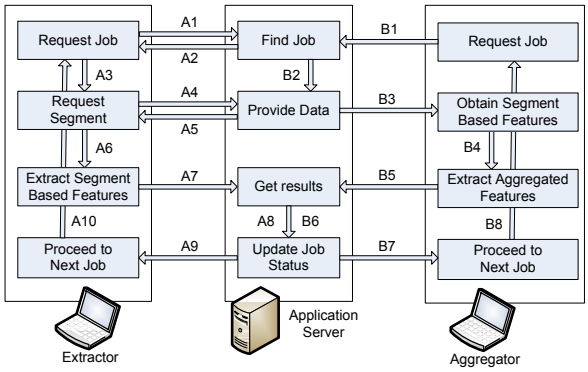

Figure 2. Two-step extraction process

3.2 Collecting the High-Level Features

The increasingly fast growth of music collections motivates the adoption of a distributed approach to perform feature extraction. Such approach, however, brings forward copyright issues, i.e., the copyrighted material in the music collection prevents the audio content to be freely distributed. We propose to address this issue by performing the feature extraction in a two-step process. In the proposed two-step extraction process, the full songs are not available to, or reconstructible by, the computers performing the extraction of the audio features.

Step 1, as illustrated by the A arrows in Figure 2, consists of dividing the audio information into short term segments of equal length and distributing them to the clients. Dividing the songs into segments offers the following advantages. First, it limits and normalizes the memory size and processor time needed on the client side to perform the feature extraction, as most extraction methods require resources proportional to the length of the audio source. Second, it avoids copyright issues as the audio segments are very short and distributing them falls under "fair use" since the segments are only temporally stored in memory on the client, and the full songs cannot be reconstructed by the clients.

Step 2, as illustrated by the B arrows in Figure 2, consists of sending the segment-based features gathered during step one to the clients. Using the features of step 1, the clients are computing high-level features for the whole piece of music. The computation of the aggregated features can be performed over the features obtained from multiple segments, as they are not subject to copyright issues. While having the segment based and the aggregated high-level features computed separately represents a potential overhead, it remains insignificant compared to the resources needed to perform each of the two feature extraction steps.

A *job* is the smallest atomic extraction or aggregation task that a remote client has to perform. Each job is uniquely identified for each extraction method and is composed of the following attributes: a song reference, a segment number, a starting point, an ending point, a lease time, a client reference, a counter of assignments, and a result holder. The assignment of jobs to clients is a critical part of the segment extraction process. Some randomness in the jobs assignment prevents clients to reconstruct the full song from the distributed segments as the segments assigned to a client will belong to different songs. However, since all the segments of a song have to be processed by step 1 before moving to step 2, assigning the segments randomly to clients delays the obtainment of results. Some locality constraints are therefore enforced in order to quickly acquire preliminary results and proceed to step 2.

In order to control which job should be assigned next, jobs are assigned in sequence. To avoid all the clients trying to obtain the same job, the assignment of a job is relaxed from being the minimal sequence number to being one of the lowest numbers in the sequence. Assigning jobs "nearly" in sequence still allows the application server to control which jobs should be prioritized for segment extraction and feature aggregation, e.g., when a similarity search is requested for a new song without any features extracted.

The current configuration of the system has shown its usability on a music collection of 41,446 songs composed of 2,283,595 non-overlapping 5 seconds segments. In terms of scalability, the job manager, supported by a PostgreSQL 8.3 database running on an Intel Core2 CPU 6700 @ 2.66GHz, 4GB of RAM under FreeBSD 7.0, was able to handle 1,766 job requests and submissions per second. Running on the same computer, the job manager and the segment handler were able to serve 87 clients per second; the bottleneck being the CPU consumption mostly due to the on-the-fly segmentation of the MP3 files. At this rate, an average bandwidth of 13,282 KB/s was consumed to transfer the segmented files of the data collection. Given that, by experience, the average processing of a segment by a modern desktop computer takes 5 seconds, the presented configuration would be able to handle over 400 clients. In a setup where the segment handler is run separately and not considering network limitations, the job manager could serve close to 9000 clients.

3.3 Similarity Search

Search among the collected features is a valuable tool to compare and evaluate extraction results. Similarity search raises two main challenges. First, similarities are of two types: similarities that can be computed rapidly on-the-fly and similarities that have to be pre-computed. Second, each similarity is tweaked dynamically with different user defined weight coefficients in order to adjust the final similarity value. In the following, we propose to find the 10 songs the most similar, with respect to a user defined weighted combination of features, to a given seed song.

Similarities that can be computed on the fly are stored in a single table, *abssim*: ("songID", "abs1", "abs2", ...), where

Abssim Table

SongID	Tempo	Pitch
1	1	0
2	2	3
3	3	2
...	...	...

Relsim Table

SeedID	SongID	Timbre
1	1	0
1	2	6
1	3	2
...	...	...
2	1	6
2	2	0
...	...	...

Figure 3. The absolute and relative similarity tables

the songID is the primary key identifying the song and abs1, abs2, ..., are the different attributes. The table is composed of 41,446 songs and 18 attributes, such as the tempo, the motion, the articulation, the pitch, and the harmonic. Similarities that cannot be computed on the fly have to be precomputed and stored for each pair of songs as some similarities are not symmetric and do not fulfill the triangular inequality. They are stored in a second table, *relsim*: ("seedID","songID","rel1"), where the "seedID" and "songID" refers to a pair of songs and "rel1" is the similarities value between the two songs. The relsim table has 1.7 billion rows and a timbre attribute. The abssim and relsim tables are illustrated in Figure 3.

The distance function, the similarity search is based on, is as follows:

$$\begin{aligned}\text{dist}(x,y) = a \times |\ \text{pitch}(x,y)| + b \times |\ \text{tempo}(x,y)| \\ + c \times |\ \text{timbre}(x,y)| + \ldots\end{aligned}$$

where x and y are two songs, tempo, pitch, and timbre are the computed differences between x and y for each feature, and a, b, and c are their respective coefficients. The pitch difference, like the tempo, can be computed on the fly: $\text{pitch}(x,y) = x.\text{pitch} - y.\text{pitch}$. The timbre difference is pre-computed and requires a lookup in the *relsim* table.

Assume the following query: find the 10 songs with the lowest distance from a given seed song. First, we compute the difference between the values of the seed song and the values of all the other songs in the table "abssim". This requires a random disk access using an index on the "songID" to read the values of the seed song, followed by a sequential scan to compute on the fly all the similarity values from the "abssim" table. Second, using an index on the "seedID", we select all the pre-computed similarities that correspond to the query songs. The song pairs with an identical "seedID" are contiguous on disk to minimize the number of disk accesses. Third, the 41,446 similarities from both resulting sets have to be joined. Fourth, the final distance function is computed and the 10 closest songs are returned.

Table 1 shows the average query processing time of the most costly operations involved in queries run on the MW described in Section 3. The performance is acceptable; most of the query processing time is cause by the join (20%) and the ordering (75%) operations. Hash joins have shown slightly better performance than, merge sort and nested loop joins. Merge joins should provide the performance results if the sort operation can be avoided by respecting the data organization on disk.

Hash join	123 ms
Merge join	141 ms
Nested loop	163 ms
Top K	405 ms
Total runtime	575 ms

Table 1. Time Cost of Similarity Search

3.4 Similarity Search within a Range

The query cost is mainly due to the join and the Top-K operations over the whole music set. We introduce similarity searches within ranges to reduce the size of the set on which the join and the Top-K are performed, thus decreasing the search time. We propose to search, within a user specified range for each feature, for the most similar songs to a given seed song.

The abssim table is used for each similarity search. Its small size allows sequential scans to be performed fast. Therefore, filtering out the values outside the query range effectively reduces the search time. Similarly, performing a filtering of the selected rows from relsim contributes to reducing the search time. However, several additional improvements can be made, they are illustrated in Figure 4.

First, a partial index on the seedID for the similarity values that are below a given threshold can be created. If the partial index has a good selectivity, speed is gained by trading the sequential scan for a few random disk access. This is of critical importance as the database grows larger. Using a threshold with a too high selectivity reduces the chances of the partial index being used.

Second, to further improve the search, one can create arrays containing pairs of songID and similarity value for each seedID. Arrays offer the following advantages. As the par-

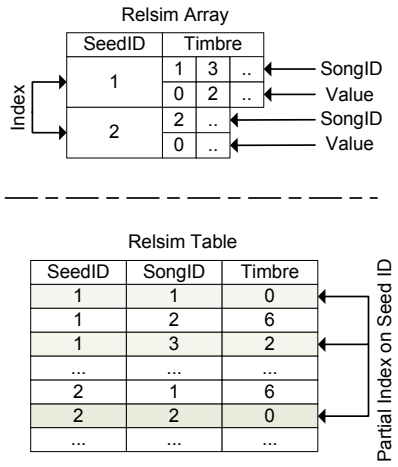

Figure 4. Range queries with arrays or a partial index

	Table	Array	Table + PI
Query time (ms)	130	15	54
Query time (%)	100	12	42
Size (MB)	70000	71550	70200
Size (%)	100	112	100

Table 2. Cost of Similarity Search within a Range

tial index, only the similar values are accessed, thus greatly reducing the cost of filtering. The similar values are clustered on disk for each seed song, similarity values are stored in the array in ascending order, and the array itself is compressed, therefore allowing a complete array to be retrieved in a single disk access. Accessing similarity values is done in an efficient way: one index lookup for locating the array, and one disk access to read the pairs of songID and value.

Table 2 presents the costs in search time and in disk space of each approach for the same set of constraints. As expected, specifying ranges for the similarity search significantly improves the response time compared to searching in the complete music collection; a simple filtering can improve the response time by a factor of 4. Using a partial index or arrays, further decreases the search time but come with a storage cost [8]. The search time improvements are dependent on the selected range as well as the selectivity of the threshold chosen for both the array and the partial index.

Similarity searches using the partial index are slower than arrays due to the random disk accesses that are required to read the values. However the partial index offers two major advantages. First, it does not require any backstage processing; the partial index is updated as new data is entered in the relsim table. Updating the arrays requires some additional processing that could be performed automatically with triggers. Second, the choice of using the partial index or not is delegated to the query planner, all queries are treated transparently regardless of the range chosen, i.e., no extra manipulation is needed to handle the array organization of the data when a threshold is reached.

4 THE ISOUNDMW

A web interface is connected to the iSoundMW. It offers a visual understanding of the functioning of the system and gives an opportunity to observe the information extracted from different songs and compare the result with the music being played simultaneously.

The setup is as follows. The music collection has a size of 41,446 songs in MP3 format, segments are non-overlapping and have a 5 seconds length. The Music Information Database is partially loaded with some segments information and some aggregated feature information, but some segments still have to be extracted and aggregated. Some clients are connected to the application server and are processing some segment extraction and feature aggregation jobs. If all the segments have not been extracted, the corresponding jobs will be prioritized. to ensure that the extracted features are available. The user interface, as shown in Figure 5, consists of a music player and a graph showing the content of the extraction as the music is being played.

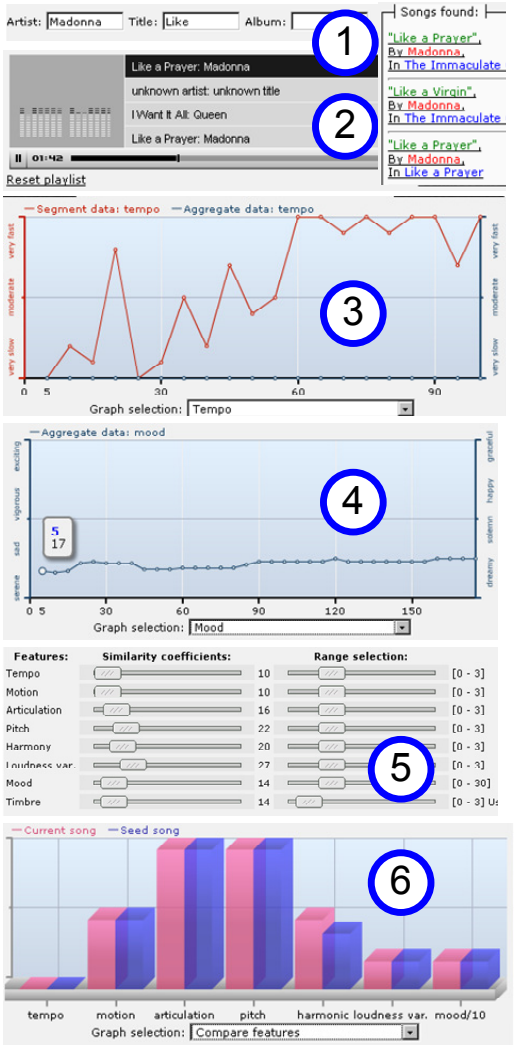

Figure 5. Web interface

The main steps of a short scenario are presented below.
Step 1: A user searches for a famous song and provides its title, artist name, or album name, e.g., the user enters "Madonna" for the artist and "Like" for the title. The system retrieves a list of maximum 20 candidates present in the database based on the ID3 tags of the MP3 in the music collection, 3 songs in this scenario. The song "Like a Prayer" is listed twice, as it belongs to two different albums. The user selects one of the two "Like a Prayer" songs.
Step 2: The system searches for the 10 most similar songs to the song selected, places them in the playlist, and starts playing the songs. The most similar song to the song selected is generally the song itself, therefore the song appears first in the generated playlist. At this stage, the search is based on

default coefficients and one the complete database. The second version of the song "Like a Prayer" appears further in the generated playlist.
Step 3: As the song is being played, the tempo analysis based on the extracted segments and the aggregated features is displayed in the graph. For each 5 seconds (corresponding to a segment length), new points are placed on the graph. If the segments have not been extracted, a request is sent to the application server to prioritize the corresponding jobs. The graph updates as new extraction results are arriving from the extraction and aggregation clients.
Step 4: Any extracted features given on an absolute scale can be displayed on the graph, e.g., the user has selected to display the mood features.
Step 5: Moving to the playlist configuration panel, the user can select, for each of the extracted features, the weight to be used as a coefficient for the similarity search, e.g., we choose to put a high coefficient on the pitch and the mood. Once the tuning of the weights is done, when the user selects a new song, the system searches for the songs that, with the given coefficients, are the most similar to the song currently being played. Additionally, the user can select a range of values in which the search should be performed.
Step 6: When similar songs are being played, the user can select to compare the currently played song with the song originally selected to generate the playlist, e.g., the main difference between the two versions of "Like a Prayer" rely in the harmonic feature.

5 CONCLUSION AND FUTURE WORK

The automatic extraction of high-level features over a large music collection suffers from two main scalability issues: high computation needs to extract the features, and copyright restrictions limiting the distribution of the audio content. This paper introduces the iSoundMW, a framework for extracting high-level audio features that addresses these scalability issues by decomposing the extraction into a two-step process. The iSoundMW has successfully demonstrated its ability to efficiently extract high-level features on a music collection of 41,446 songs. Furthermore, the iSoundMW proves to be efficient and flexible for performing similarity searches using the extracted features. This is done by allowing users to constrain the search within a range and specify a weighted combination of high-level features. To further optimize the search, three different approaches are compared in terms of query time and storage. The threshold for building the partial index and arrays are decisive parameters to obtain good search performance.

Future work encompasses integrating different similarity functions in the features search, providing comparison between them, and enabling user feedback and its reuse for user specific recommendation.

6 ACKNOWLEDGMENTS

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A COMPARISON OF STATISTICAL AND RULE-BASED MODELS OF MELODIC SEGMENTATION

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ABSTRACT

We introduce a new model for melodic segmentation based on information-dynamic analysis of melodic structure. The performance of the model is compared to several existing algorithms in predicting the annotated phrase boundaries in a large corpus of folk music.

1 INTRODUCTION

The segmentation of melodies into phrases is a fundamental (pre-)processing step for many MIR applications including melodic feature computation, melody indexing, and retrieval of melodic excerpts. In fact, the melodic phrase is often considered one of the most important basic units of musical content [16] and many large electronic corpora of music are structured or organised by phrases, for example, the Dictionary of Musical Themes by Barlow and Morgenstern [2], the Essen Folksong Collection (EFSC) [33] or the RISM collection [28].

At the same time, melodic grouping is thought to be an important part of the perceptual processing of music [11, 14, 27]. It is also fundamental to the phrasing of a melody when sung or played. Melodic segmentation is a task that musicians and musical listeners perform regularly in their everyday musical practice.

Several algorithms have been proposed for the automated segmentation of melodies. These algorithms differ in their modelling approach (supervised learning, unsupervised learning, music-theoretic rules), and in the type of information they use (global or local).

In this paper, we introduce a new statistical model of melodic segmentation and compare its performance to several existing algorithms on a melody segmentation task. The motivation for this model comparison is two-fold: first, we are interested in the performance differences between different types of model; and second, we aim to build a hybrid model that achieves superior performance by combining boundary predictions from different models.

2 BACKGROUND

2.1 Evaluation Measures

In modern information retrieval, *Precision*, *Recall*, and *F1* have become standard measures for assessing model performance. These measures are usually defined in terms of *True Positives, TP* (i.e. the number of times a model predicts a specific outcome correctly), *False Positives, FP* (i.e. the number of times a model predicts a specific outcome incorrectly), and *False Negatives, FN* (i.e. the number of times a model incorrectly does not predict a specific outcome):

$$precision = \frac{TP}{TP + FP} \quad recall = \frac{TP}{TP+FN}$$

$$F1 = \frac{2 \cdot precision \cdot recall}{precision + recall}$$

2.2 Models of Melodic Segmentation

GTTM: Melodic grouping has traditionally been modelled through the identification of local discontinuities or changes between events in terms of temporal proximity, pitch, duration and dynamics [6, 16, 36]. Perhaps the best known examples are the Grouping Preference Rules (GPRs) of the Generative Theory of Tonal Music (GTTM) [16]. The most widely studied of these GPRs predict that phrase boundaries will be perceived between two melodic events whose temporal proximity is less than that of the immediately neighbouring events due to a slur, a rest (GPR 2a) or a relatively long inter-onset interval or IOI (GPR 2b) or when the transition between two events involves a greater change in register (GPR 3a), dynamics (GPR 3b), articulation (GPR 3c) or duration (GPR 3d) than the immediately neighbouring transitions. Some of these GPRs have been quantified [14] and studied in psychological experiments [11, 14].

LBDM: Cambouropoulos [6] proposes a related model in which boundaries are associated with any local change in interval magnitudes. The *Local Boundary Detection Model* (LBDM) consists of a *change* rule, which assigns boundary strengths in proportion to the degree of change between consecutive intervals, and a *proximity* rule, which scales the boundary strength according to the size of the intervals involved. The LBDM operates over several independent parametric melodic profiles $P_k = [x_1, x_2, \ldots, x_n]$ where $k \in \{\text{pitch, ioi, rest}\}, x_i > 0, i \in \{1, 2, \ldots, n\}$ and the boundary strength at interval x_i is given by:

$$s_i = x_i \times (r_{i-1,i} + r_{i,i+1})$$

where the degree of change between two successive intervals:

$$r_{i,i+1} = \begin{cases} \frac{|x_i - x_{i+1}|}{x_i + x_{i+1}} & \text{if } x_i + x_{i+1} \neq 0 \wedge x_i, x_{i+1} \geq 0 \\ 0 & \text{if } x_i = x_{i+1} = 0. \end{cases}$$

For each parameter k, the boundary strength profile $S_k = [s_1, s_2, \ldots, s_n]$ is calculated and normalised in the range $[0, 1]$. A weighted sum of the boundary strength profiles is computed using weights derived by trial and error (0.25 for $pitch$ and $rest$, and 0.5 for ioi), and boundaries are predicted where the combined profile exceeds a predefined threshold.

Grouper: Temperley [36] introduces a model called *Grouper* which accepts a melody, in which each note is represented by its onset time, off time, chromatic pitch and level in a metrical hierarchy, and returns a single, exhaustive partitioning of the melody into non-overlapping groups. The model operates through the application of three *Phrase Structure Preference Rules* (PSPRs):

PSPR 1 (Gap Rule): prefer to locate phrase boundaries at (a) large IOIs and (b) large offset-to-onset intervals (OOI); PSPR 1 is calculated as the sum of the IOI and OOI divided by the mean IOI of all previous notes;

PSPR 2 (Phrase Length Rule): prefer phrases with about 10 notes, achieved by penalising predicted phrases by $|(\log_2 N) - 3|$ where N is the number of notes in the predicted phrase;

PSPR 3 (Metrical Parallelism Rule): prefer to begin successive groups at parallel points in the metrical hierarchy.

The first rule is another example of the Gestalt principle of temporal proximity (cf. GPR 2 above); the second was determined through an empirical investigation of the typical phrase lengths in a collection of folk songs. The best analysis of a given piece is computed offline using a dynamic programming approach where candidate phrases are evaluated according to a weighted combination of the three rules. The weights were determined through trial and error. By way of evaluation, Temperley used Grouper to predict the phrase boundaries marked in 65 melodies from the EFSC achieving a recall of 0.76 and a precision 0.74.

Other Models: Tenney and Polansky [37] were perhaps the first to propose models of melodic segmentation based on Gestalt-like rules. Other authors have combined Gestalt-like rules with higher-level principles based on parallelism and music structure [1, 7]. Ferrand et al. [13] introduce an approach based on the idea of 'melodic density' (i.e., segment at points of low cohesion between notes) and compare the methods performance to the LBDM. In contrast, Bod [3] argues for a supervised learning approach to modelling melodic grouping structure. A model based on data-oriented parsing (DOP) yielded $F1 = 0.81$ in predicting unseen phrase boundaries in the EFSC. A qualitative examination of the data revealed that 15% of the phrase boundaries predicted by the Markov-DOP parser cannot be accounted for by Gestalt principles. These models are mentioned for completeness, but are not included in our comparison.

2.3 The IDyOM Model

We present a new model of melodic grouping (the Information Dynamics Of Music model) that is inspired by previous research in musicology, music perception, computational linguistics and machine learning.

From a musicological perspective, it has been proposed that perceptual groups are associated with points of closure where the ongoing cognitive process of expectation is disrupted either because the context fails to stimulate strong expectations for any particular continuation or because the actual continuation is unexpected [21, 22]. These proposals may be given precise definitions in an information-theoretic framework which we define by reference to a model of unsupervised inductive learning of melodic structure. Briefly, the models we propose output conditional probabilities of an event e, given a preceding sequential context c. Given such a model, the degree to which an event appearing in a given context in a melody is unexpected can be defined as the *information content*, $h(e|c)$, of the event given the context:

$$h(e|c) = \log_2 \frac{1}{p(e|c)}.$$

The information content can be interpreted as the contextual unexpectedness or surprisal associated with an event. Given an alphabet $\mathcal{E}$ of events which have appeared in the prior experience of the model, the uncertainty of the model's expectations in a given melodic context can be defined as the *entropy* or average information content of the events in $\mathcal{E}$:

$$H(c) = \sum_{e \in \mathcal{E}} p(e|c) h(e|c).$$

We propose that boundaries will occur before events for which unexpectedness (h) and uncertainty (H) are high.

In addition to the musicological basis, there is a precedent for these ideas in experimental psychology. Empirical research has demonstrated that infants and adults use the implicitly learnt statistical properties of pitch [32], pitch interval [30] and scale degree [29] sequences to identify segment boundaries on the basis of higher digram ($n = 2$) transition probabilities within than between groups.

There is also evidence that related information-theoretic quantities are important in cognitive processing of language. For example, it has recently been demonstrated that the difficulty of processing words is related both to their information content [17] and the induced changes in entropy of grammatical continuations [15]. More specifically, experimental work has demonstrated that infants and adults reliably identify grouping boundaries in sequences of synthetic syllables [31] on the basis of higher transition probabilities within than between groups.

Furthermore, research in machine learning and computational linguistics has demonstrated that algorithms that segment before unexpected events can successfully identify word boundaries in infant-directed speech [4]. Similar strategies for identifying word boundaries have been implemented using recurrent neural networks [12]. Recently, Cohen et al. [8] proposed a general method for segmenting

sequences based on two principles: first, so as to maximise n-gram frequencies to the left and right of the boundary; and second, so as to maximise the entropy of the conditional distribution across the boundary. The algorithm was able to successfully identify word boundaries in text from four languages and episode boundaries in the activities of a mobile robot.

IDyOM itself is based on n-gram models commonly used in statistical language modelling [18]. An n-gram is a sequence of n symbols and an n-gram model is simply a collection of such sequences each of which is associated with a frequency count. During the *training* of the statistical model, these counts are acquired through an analysis of some corpus of sequences (the training set) in the target domain. When the trained model is exposed to a sequence drawn from the target domain, it uses the frequency counts associated with n-grams to estimate a probability distribution governing the identity of the next symbol in the sequence given the $n-1$ preceding symbols. The quantity $n-1$ is known as the *order* of the model and represents the number of symbols making up the context within which a prediction is made.

However, n-gram models suffer from several problems, both in general and specifically when applied to music. The first difficulties arise from the use of a fixed-order. Low-order models fail to provide an adequate account of the structural influence of the context. However, increasing the order can prevent the model from capturing much of the statistical regularity present in the training set (an extreme case occurring when the model encounters an n-gram that does not appear in the training set and returns an estimated probability of zero). In order to address these problems, the IDyOM model maintains frequency counts during training for n-grams of all possible values of n in any given context. During prediction, distributions are estimated using a weighted sum of all models below a variable order bound. This bound is determined in each predictive context using simple heuristics designed to minimise uncertainty. The combination is designed such that higher-order predictions (which are more specific to the context) receive greater weighting than lower-order predictions (which are more general).

Another problem with n-gram models is that a trained model will fail to make use of local statistical structure of the music it is currently analysing. To address this problem, IDyOM includes two kinds of model: first, the *long-term* model that was trained over the entire training set in the previous step; and second, a *short-term* model that is trained incrementally for each individual melody being predicted. The distributions returned by these models are combined using an entropy weighted multiplicative combination scheme [26] in which greater weights are assigned to models whose predictions are associated with lower entropy (or uncertainty) at that point in the melody.

A final issue regards the fact that music is an inherently multi-dimensional phenomenon. Musical events have many attributes including pitch, onset time, duration, timbre and so on. In addition, sequences of these attributes may have multiple relevant dimensions. For example, pitch interval, pitch class, scale degree, contour and many other derived features are important in the perception and analysis of pitch structure. In order to accommodate these properties, the modelling process begins by choosing a set of basic features that we are interested in predicting. As these basic features are treated as independent attributes, their probabilities are computed separately and in turn, and the probability of a note is simply the product of the probabilities of its attributes. Each basic feature (e.g., pitch) may then be predicted by any number of models for different derived features (e.g., pitch interval, scale degree) whose distributions are combined using the same entropy-weighted scheme.

The use of long- and short-term models, incorporating models of derived features, the entropy-based weighting method and the use of a multiplicative as opposed to a additive combination scheme all improve the performance of IDyOM in predicting the pitches of unseen melodies [24, 26]. Full details of the model and its evaluation can be found elsewhere [9, 23, 24, 26].

The conditional probabilities output by IDyOM in a given melodic context may be interpreted as contextual expectations about the nature of the forthcoming note. Pearce and Wiggins [25] compare the melodic pitch expectations of the model with those of listeners in the context of single intervals [10], at particular points in British folk songs [34] and throughout two chorale melodies [19]. The results demonstrate that the statistical system predicts the expectations of listeners as least as well as the two-factor model of Schellenberg [35] and significantly better in the case of more complex melodic contexts.

In this work, we use the model to predict the pitch, IOI and OOI associated with melodic events, multiplying the probabilities of these attributes together to yield the overall probability of the event. For simplicity, we don't use any derived features. We then focus on the unexpectedness of events (information content, h) using this as a boundary strength profile from which we compute boundary locations (as described below). The role of entropy (H) will be considered in future work. The IDyOM model differs from the GPRs, the LBDM and Grouper in that it is based on statistical learning rather than symbolic rules and it differs from DOP in that it uses unsupervised rather than supervised learning.

2.4 Comparative evaluation of melody segmentation algorithms

Most of the models described above were evaluated to some extent by their authors and, in some cases, compared quantitatively to other models. In addition, however, there exist a small number of studies that empirically compare the performance of different models of melodic grouping. These studies differ in the algorithms compared, the type of ground truth data used, and the evaluation metrics applied. Melucci and Orio [20], for example, collected the boundary indications of 17 music scholars on melodic excerpts from 20 works by Bach, Mozart, Beethoven and Chopin. Having combined the boundary indications into a ground truth, they evaluated the performance of the LBDM against three base-

line models that created groups containing fixed (8 and 15) or random (between 10 and 20) numbers of notes. Melucci and Orio report false positives, false negatives, and a measure of disagreement which show that the LBDM outperforms the other models.

Bruderer [5] presents a more comprehensive study of the grouping structure of melodic excerpts from six Western pop songs. The ground truth segmentation was obtained from 21 adults with different degrees of musical training; the boundary indications were summed within consecutive time windows to yield a quasi-continuous boundary strength profile for each melody. Bruderer examines the performance of three algorithms: Grouper, LBDM and the summed GPRs quantified in [14] (GPR 2a, 2b, 3a and 3d). The output of each algorithm is convolved with a Gaussian window to produce a boundary strength profile that is then correlated with the ground truth. Bruderer reports that the LBDM achieved the best and the GPRs the worst performance.

Another study [38] compared the predictions of the LBDM and Grouper to segmentations at the phrase and subphrase level provided by 19 musical experts for 10 melodies in a range of styles. The performance of each model on each melody was estimated by averaging the F1 scores over the 19 experts. Each model was examined with parameters optimised for each individual melody. The results indicated that Grouper tended to outperform the LBDM. Large IOIs were an important factor in the success of both models. In another experiment, the predictions of each model were compared with the transcribed boundaries in several datasets from the EFSC. The model parameters were optimised over each dataset and the results indicated that Grouper (with mean F1 between 0.6 and 0.7) outperformed the LBDM (mean F1 between 0.49 and 0.56).

All these comparative studies used ground truth segmentations derived from manual annotations by human judges. However, only a limited number of melodies can be tested in this way (ranging from 6 in the case of [5] to 20 by [20]). Apart from Thom et al. [38], Experiment D, there has been no thorough comparative evaluation over a large corpus of melodies annotated with phrase boundaries.

3 METHOD

3.1 The Ground Truth Data

We concentrate here on the results obtained for a subset of the EFSC, database `Erk`, containing 1705 Germanic folk melodies encoded in symbolic form with annotated phrase boundaries which were inserted during the encoding process by folk song experts. The dataset contains 78,995 sounding events at an average of about 46 events per melody and overall about 12% of notes fall before boundaries. There is only one hierarchical level of phrasing and the phrase structure exhaustively subsumes all the events in a melody.

3.2 Making Model Outputs Comparable

The outputs of the algorithms tested vary considerably. While Grouper marks each note with a binary indicator (1 = boundary, 0 = no boundary), the other models output a positive real number for each note which can be interpreted as a boundary strength. In contrast to Bruderer [5] we chose to make all segmentation algorithms comparable by picking binary boundary indications from the boundary strength profiles.

To do so, we devised a method called *Simple Picker* that uses three principles. First, the note following a boundary should have a greater or equal boundary strength than the note following it: $S_n \geq S_{n+1}$. Second, the note following a boundary should have a greater boundary strength than the note preceding it: $S_n > S_{n-1}$. Whilst these principles, simply ensure that a point is a local peak in the profile, the third specifies how high the point must be, relative to earlier points in the profile, to be considered a peak. Thus the note following a boundary should have a boundary strength greater than a threshold based on the linearly weighted mean and standard deviation of all notes preceding it:

$$S_n > k\sqrt{\frac{\sum_{i=1}^{n-1}(w_i S_i - \overline{S}_{w,1\ldots n-1})^2}{\sum_1^{n-1} w_i}} + \frac{\sum_{i=1}^{n-1} w_i S_i}{\sum_1^{n-1} w_i}$$

The third principle makes use of the parameter k which determines how many standard deviations higher than the mean of the preceding values a peak must be to be picked. In practice, the optimal value of k varies between algorithms depending on the nature of the boundary strength profiles they produce.

In addition, we modified the output of all models to predict an implicit phrase boundary on the last note of a melody.

3.3 The Models

The models included in the comparison are as follows:

Grouper: as implemented by [36]; [1]

LBDM: as specified by [6] with $k = 0.5$;

IDyOM: with $k = 2$;

GPR2a: as quantified by [14] with $k = 0.5$;

GPR2b: as quantified by [14] with $k = 0.5$;

GPR3a: as quantified by [14] with $k = 0.5$;

GPR3d: as quantified by [14] with $k = 2.5$;

Always: every note falls on a boundary;

Never: no note falls on a boundary.

Grouper outputs binary boundary predictions but the output of every other model was processed by Simple Picker using a value of k was chosen from the set $\{0.5, 1, 1.5, 2, 2.5, 3, 3.5, 4\}$ so as to maximise F1 (and secondarily Recall).

4 RESULTS

The results of the model comparison are shown in Table 1. The four models achieving mean F1 values of over 0.5

[1] Adapted for use with Melconv 2 by Klaus Frieler.

Model	Precision	Recall	F1
Hybrid	0.87	0.56	0.66
Grouper	0.71	0.62	0.66
LBDM	0.70	0.60	0.63
GPR2a	0.99	0.45	0.58
IDyOM	0.76	0.50	0.58
GPR2b	0.47	0.42	0.39
GPR3a	0.29	0.46	0.35
GPR3d	0.66	0.22	0.31
Always	0.13	1.00	0.22
Never	0.00	0.00	0.00

Table 1. The model comparison results in order of mean F1 scores.

(Grouper, LBDM, GPR2a, IDyOM) were chosen for further analysis. Sign tests between the F1 scores on each melody indicate that all differences between these models are significant at an alpha level of 0.01, with the exception of that between GPR2a and LBDM. In order to see whether further performance improvements could be achieved by a combined model, we constructed a logistic regression model including Grouper, LBDM, IDyOM and GPR2a as predictors. Backwards stepwise elimination using the Bayes Information Criterion (BIC) failed to remove any of the predictors from the overall model. The performance of the resulting model is shown in the top row of Table 1. Sign tests demonstrated that the Hybrid model achieved better F1 scores on significantly more melodies than each of the other models.

5 DISCUSSION

We would like to highlight four results of this evaluation study. First, we were surprised by the strong performance of one of the GTTM preference rule, GPR2a. This points to the conclusion that rests, perhaps above all other melodic parameters, have a large influence on boundaries in this melodic style. Consequently, all of the high-performing rule-based models (Grouper, LBDM, GPR2a) make use of a rest or temporal gap rule while IDyOM includes rests in its probability estimation. Future research should undertake a more detailed qualitative comparison of the kinds of musical context in which each model succeeds or fails to predict boundaries.

Second, it is interesting to compare the results to those reported in other studies. In general, the performance of Grouper and LBDM are comparable to their performance on a different subset of the EFSC reported by Thom et al. [38]. The performance of Grouper is somewhat lower than that reported by Temperley [36] on 65 melodies from the EFSC. The performance of all models is lower than that of the supervised learning model reported by Bod [3].

Third, the hybrid model which combines Grouper, LBDM, GPR2a and IDyOM generated better performance values than any of its components. The fact that the *F1* value seems to be only slightly better than Grouper is due to the fact that logistic regression optimises the log-likelihood function for whether or not a note is a boundary given the boundary indications of the predictor variables (models). It therefore uses information about positive boundary indications (*P*) and negative boundary indications (*N*) to an equal degree, in contrast to *F1*. This suggests options, in future research, for assigning different weights to *P* and *N* instances or including the raw boundary profiles of LBDM and IDyOM in the logistic regression procedure. Another possibility is to use boosting to combine the different models which may lead to better performance enhancements than logistic regression.

Finally, it is interesting to note that an unsupervised learning model (IDyOM) that makes no use of music-theoretic rules about melodic phrases performed as well as it does, in comparison to sophisticated rule-based models. In comparison to supervised learning methods such as DOP, IDyOM does not require pre-segmented data as a training corpus. This may not be an issue for folk-song data where we have large corpora with annotated phrase boundaries but is a significant factor for other musical styles such as pop. IDyOM learns regularities in the melodic data it is trained on and outputs probabilities of note events which are ultimately used to derive an information content (unexpectedness) for each note event in a melody. In turn, this information-theoretic quantity (in comparison to that of previous notes) is used to decide whether or not the note falls on a boundary.

We argue that the present results provide preliminary evidence that the notion of expectedness is strongly related to boundary detection in melodies. In future research, we hope to achieve better segmentation performance by providing the statistical model with more sophisticated melodic representations and examining the role of entropy (uncertainty) in melodic boundary detection.

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A FRAMEWORK FOR AUTOMATED SCHENKERIAN ANALYSIS

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ABSTRACT

In Schenkerian analysis, one seeks to find structural dependences among the notes of a composition and organize these dependences into a coherent hierarchy that illustrates the function of every note. This type of analysis reveals multiple levels of structure in a composition by constructing a series of simplifications of a piece showing various elaborations and prolongations. We present a framework for solving this problem, called IVI, that uses a state-space search formalism. IVI includes multiple interacting components, including modules for various preliminary analyses (harmonic, melodic, rhythmic, and cadential), identifying and performing reductions, and locating pieces of the *Ursatz*. We describe a number of the algorithms by which IVI forms, stores, and updates its hierarchy of notes, along with details of the *Ursatz*-finding algorithm. We illustrate IVI's functionality on an excerpt from a Schubert piano composition, and also discuss the issues of subproblem interactions and the multiple parsings problem.

1 SCHENKERIAN ANALYSIS

A number of types of music analysis are concerned with "labeling" individual objects in a musical score. In harmonic analysis, labels are assigned to chords and notes corresponding to harmonic function; in contrapuntal voice segregation, labels are assigned to notes indicating voice assignment. A rhythmic analysis may assign different levels of metrical importance to notes. These styles of analysis are similar in that they often describe musical components in isolation, or only in relation to their immediate neighbors on the musical surface.

Structural analysis, on the other hand, emphasizes discovering relationships among notes and chords in a composition, rather than studying individual tones in a vacuum. The word "structure" here refers to "the complete fabric of the composition as established by melody, counterpoint, and harmony in combination" [1].

Schenkerian analysis is the most well-developed type of structural analysis. This type of analysis examines the "interrelationships among melody, counterpoint, and harmony" [1] in a hierarchical manner. Schenker's theory of music allows one to determine which notes in a passage of music are more structurally significant than others. It is important not to confuse "structural significance" with "musical importance;" a musically important note (e.g., crucial for articulating correctly in a performance) can be a very insignificant tone from a structural standpoint. Judgments regarding structural importance result from finding dependences between notes or sets of notes: if a note X derives its musical function or meaning from the presence of another note Y, then X is dependent on Y and Y is deemed more structural than X.

The process of completing a Schenkerian analysis proceeds in a recursive manner. Starting from the musical score of a composition, one may locate any number of structural dependences. After no more can be found, an abstracted score is produced by rearranging or removing the less structural notes. The new abstracted score will reveal new dependences: when their subservient neighbors are moved or eliminated, various structurally important notes in the original score will be deemed less significant to their new neighbors.

This iterative process illustrates Schenker's conception that tonal compositions consist of a "continuum of interrelated structural levels," [1] where each structural level of the music represents that composition at a different level of abstraction. Each successively abstract level expands or prolongs various aspects of the previous one. The most abstract level of the piece is the *background* level. At the other end of the spectrum is the *foreground* level: an analysis at this level usually still contains most of the notes of the score and most closely represents the musical surface. Between these levels is the *middleground* level. While Schenker's own analyses usually only contain these three levels, there can be many levels in between the surface level music and the ultimate background level; rarely are the levels clearly delineated.

Schenker theorized that at the background level, all tonal works could be represented by one of three basic outlines, consisting of a three chord harmonic progression (the *Bassbrechung* or *bass arpeggiation*) supporting a three-, five-, or eight-note descending melodic line (the *Urlinie* or *fundamental line*). Together, these form the *Ursatz* or *fundamen-*

tal structure. The *Ursatz* is a basic structure that appears over most of the length of a single composition, though it often manifests itself at other levels in the structural hierarchy as well. While the *Ursatz* is an important facet of Schenkerian analysis, the idea of structural levels is more encompassing, as the *Ursatz* can be seen as another musical device that may appear at multiple levels of abstraction [2].

2 PREVIOUS WORK AND STATE OF THE ART

The most widely known formalization of musical structure in a hierarchical fashion similar to Schenker's is that of Lerdahl and Jackendoff [8]. The authors attempt to describe the structure of music from a linguistic perspective by providing preference-rule systems that govern various aspects of musical structure. They present a set of rules for grouping of notes in a voice, and another for deducing metrical structure in term of strong beats and weak beats. Their most intriguing contributions here, however, are the preference-rule systems for two recursive reductive systems, one based on "time-span analysis" that is governed by metrical and grouping structure, and another based on "prolongational analysis" that is controlled by the ideas of rising and falling musical tension in a piece. These reductions can be illustrated by trees, where the leaves are the surface-level notes of the piece and higher-level branches represent the more structural tones, or by an equivalent musical depiction as a sequence of staves showing successive levels of the resultant hierarchy of pitches.

Though Lerdahl and Jackendoff frame their discussion in terms of discovering a grammar of music, they acknowledge that their system is incomplete and probably cannot be directly turned into a formal algorithm. This is mainly due to the lack of weightings for the preference rules and that the authors' representations are based primarily on vertical segmentation of the input music as chords, and do not give enough weight to voice-leading considerations. Nevertheless, a number of people have undertaken formalization of Lerdahl and Jackendoff's system, though the results usually require tweaking of parameters. For example, Hirata and Aoyagi [6] present a framework for the representation of musical knowledge in the spirit of Lerdahl and Jackendoff, while Hamanaka et al. [4, 5] describe an implementation that can automatically make reductions according to a set of preference rules, but the weights on the rules must be adjusted by hand for each piece.

Temperley [15] presents models for meter, phrase structure, voice segregation, pitch spelling, harmonic structure, and key structure using a preference rule system very similar to that of Lerdahl and Jackendoff. Temperley, however, went further and was able to use his models in algorithms that "implement Lerdahl and Jackendoff's initial conception." The models and algorithms vary in level of success and sophistication [13]. All of the models developed could be useful in automating Schenkerian analysis, though the concept of hierarchical reductions is not discussed. Temperley later re-developed some of these modules and a number of new ones with a Bayesian statistics slant [16]; here he does present a high-level discussion on how to test or verify Schenker's ideas from a Bayesian viewpoint.

Schenkerian analysis has largely resisted formalization because it was not presented in any sort of formal manner by Schenker himself. Indeed, many textbooks illustrate concepts largely through examples and long paragraphs of prose [1, 3] rather than by giving a recipe for constructing an analysis step-by-step. There have only been three large-scale initiatives in developing a computational procedure for Schenkerian analysis; other relevant work (e.g., Meehan [12]) was undertaken on a smaller scale and was restricted to models, not algorithms. The first large project, the work of Smoliar [14], resulted in a tool used to assist a human in performing an analysis, largely by confirming the validity of reductions; the system did not perform any analysis on its own. The second, by Kassler [7], described a model that took the analysis from the middleground level to the background, but did not work directly with the surface-level musical notes (what one sees in the actual score). The third project in Schenkerian analysis is more current: an undertaking by Marsden [9, 10, 11], which seeks to derive an analysis directly from a MIDI-style music representation. This project, however, is still in its infancy and is focused on calculating all possible reductions using a dynamic programming algorithm and scoring each analysis. For example, it makes no mention of locating the *Ursatz* (a critical step), does not take advantage of other notational information that would be given by a higher-level representation of a score such as MusicXML, and the published research makes no mention of harnessing previous research in voice segregation, key finding, or harmonic analysis to bootstrap the process.

3 THE IVI FRAMEWORK

We now explain the functionality of the IVI framework, a system to perform automated Schenkerian analysis. The IVI name (pronounced like ivy, the plant) was chosen for the I-V-I (tonic-dominant-tonic) harmonic structure that Schenker theorized was at the background level of all tonal compositions. The framework consists of numerous components, which we detail in the following section.

We have chosen MusicXML as the input format for IVI. While the MIDI file format is more widely used for music file storage than MusicXML, the richer score format of MusicXML facilitates the analysis process. A MIDI file gives little information to the user other than pitch and timing information; even such vital clues as the spelling of accidentals is lost. Though hierarchical music analysis is designed to reflect processes in tonal music that are aurally

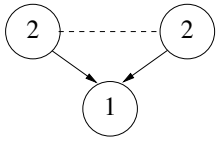

Figure 1. The DAG structure illustrating how the middle note in a neighbor tone musical figure is dependent on the outer tones.

perceived by music theorists and average listeners alike (and so the process could conceivably use MIDI as input), having as much of the printed score as possible available to IVI removes many of the preprocessing steps that otherwise would be inevitable. MusicXML gives us key and time signatures, clefs, beaming information, stem directions, slurs, ornaments, barlines, and written repeats, all of which are potentially useful for Schenkerian analysis, and most of which are not included in a MIDI file.

3.1 Data Structures

The primary data structure that IVI uses to store an analysis (in any state of completeness) is the directed acyclic graph, or DAG. A DAG is an abstract data type consisting of a set of *nodes* (or vertices) and *arcs* (or edges). An arc is a connection between two nodes.

We augment the basic DAG definition to support two different kinds of arcs:

- Regular (directed) arcs: An arc directed from node x to node y specifies that the note(s) represented by y are dependent on those represented by x. That is, a node's children are its dependents.
- Equivalence arcs: These arcs are non-directional. An equivalence arc connecting nodes x and y represents the claim that x and y must be at the same structural level. These arcs are inserted to force nodes to be at the same level.

Note that with equivalence arcs as well as regular arcs between nodes, we can enforce both dependences and structural level equivalences. We show the DAG for a neighbor note idiom in Figure 1, where regular arcs are shown as solid lines, and equivalence arcs are shown as dashed lines.

Initially, the DAG contains one node for each note in the input composition and no arcs; this corresponds to the surface-level information before any analysis has occurred.

3.2 The Informed Search Paradigm

We can formalize Schenkerian analysis as a computational search problem. The search space consists of all possible empty, intermediary, and completed Schenkerian analyses of a piece; since we are using DAGs, this is the space of all possible DAGs over n nodes. This space is prohibitively large for an exhaustive search; Marsden [11] showed that even if we restrict ourselves to DAGs that resemble binary trees, the size of the search space is still factorial in n, the number of notes in the input piece. Therefore, we employ various informed (heuristic) search methods.

A state in the search space consists of an individual DAG, corresponding to a partial Schenkerian analysis. Goal states are completed analyses, where there are only a small number of nodes at the top-most level of the DAG; these nodes will correspond to the tones of the *Ursatz*. The various operators that IVI can apply correspond to either individual Schenkerian reductions, which transform the DAG by adding new nodes and edges, or adding *properties* to the search state. A property is an attribute-value pair that stores additional information about the the state of the analysis not captured by the DAG; the attributes are name-note pairings. The most important property is called `level` and may be attached to any note. This is an integer that represents the importance of that note in the structural hierarchy. Higher numbers represent higher levels of structure. Initially, all notes are assigned a `level` of 1. There is a procedure (detailed later) that updates these values as the analysis progresses. Additional properties correspond to marks an analyst would make on a physical musical score, such as harmonic analysis chord symbols. Since there are myriad property assignments and reductions possible at numerous locations in the analysis, we are investigating using various ranking algorithms to evaluate possible reductions and determine the musical plausibility of each.

Two metrics are needed to perform an intelligent search. If n is a state in the search tree, we need a metric $\hat{g}(n)$ to tell us the distance from the start state (a "blank" analysis) to n, and $\hat{h}(n)$ to tell us the estimated distance from n to a goal state. $\hat{g}$ depends primarily on the number of reductions applied and the aggregate musical plausibility level of those reductions. $\hat{h}$ takes into account more global measurements, such as the plausibility of the current analysis as a whole (rather than individual reductions), the `level`s of each of the most structural notes in the DAG, and the probability of further reductions.

Before the intelligent search begins, however, IVI needs to perform a number of preliminary analyses: a harmonic analysis to determine the initial chord structure, a melodic analysis to determine voice leading, a rhythmic analysis to locate strong beats and weak beats, and a cadential analysis to locate cadences. While we already have basic algorithms in place for these first two analyses, we plan on harnessing the wealth of existing published algorithms for harmonic, melodic, and rhythmic analyses to improve IVI. We have found no existing systems for explicitly finding cadences, however, and so we are currently implementing our own cadence-locating algorithm that will use the results from the other three analysis modules and measurements of harmonic

rhythm to identify cadences.

After the preprocessing steps, the search control algorithm begins its work. Once a goal state is found, IVI can save the resulting analysis and backtrack to locate other plausible analyses if desired by the user. We embrace the idea of multiple musically-plausible analyses. Because we are operating with numerical parameters, there will be a single analysis that is deemed the best (or most-likely). Still, much musical knowledge and insight can be gleaned from seeing the "second tier" of analyses.

4 IMPLEMENTATION DETAILS

4.1 Memory Issues

One of the major problems facing a naïve implementation of IVI's search control is that of memory management. Storing a complete DAG and all the properties of all the notes at every search state in the search tree would be infeasible, as the computer would quickly run out of RAM. While we are investigating using beam search to ameliorate this situation, there is a fundamental data structure change we use to lower our memory requirements.

First, we choose to implement the storage of the DAG itself as a collection of properties in a search state. We use properties named "children" and "parents" to store a set of regular arcs indicating structural dependences, and an "equivalent-to" property to store equivalence arcs. By doing this, all local information about a search state is combined into a collection of properties. We cannot store a global set of properties in a central location as we need to support multiple (possibly conflicting) analyses that may interpret the same musical structures in different fashions, leading to different values for the same name-note property attribute.

Because properties are local to a search state, one may be tempted to copy the property set (currently implemented as a hash table) when creating successor states of a particular search state. However, much memory can be saved (with a small sacrifice in time) if we allow properties to be implicitly inherited from a search state's predecessor. When IVI needs to look up a property, it checks the property set of the current search state, and if the property attribute in question is not found, it walks up the search state's chain of preceding states, looking for the appropriate attribute. This inheritance model prevents us from copying properties many times over and wasting memory. We plan on investigating the loss in efficiency due to time spent traversing the search tree, and possibly implementing a compromising solution where properties *are* duplicated, but only at certain states in the search tree.

4.2 Updating the Structural Levels

Updating the `level` of nodes in the DAG must take place after reductions are performed that modify the DAG in some fashion. If one temporarily coalesces nodes connected by equivalence edges into "meta-nodes," then the pure DAG structure is restored while preserving equivalence relations. It is then possible to assign new `level` values by using a dynamic programming algorithm to find the longest path from each node to a node with no parents; these path lengths become the new `level` values.

4.3 Voice Segregation and Searching for the Ursatz

IVI's current voice segregation algorithm works by finding voice-leading connections between individual notes. It prefers stepwise connections between notes, and therefore does very well at identifying cases of implicit polyphony, where voice changes are often signaled by leaps. The procedure also can detect voices being doubled at the octave; instead of creating separate voices for each musical line in the doubling, properties are stored in the current search state indicating the notes are part of a doubling.

Our computational procedure for finding the *Ursatz* in a piece requires we already have the soprano line extracted, which IVI's voice segregation algorithm does for us. IVI first identifies the final most structural cadence (the one with the highest values for `level`) and locates the $\hat{2}$-$\hat{1}$ and V-I pieces. (We use an integer with a circumflex accent to indicate a scale degree.) Next, one must locate possible locations for the primary tone of the *Urlinie*. A method for accomplishing this is to search for candidate primary tones by locating all notes on scale degrees $\hat{3}$, $\hat{5}$, and $\hat{8}$, and selecting the most structural of each scale degree. Now that potential beginning and ending points are determined, the last step is to locate the intermediary tones of the *Urlinie* (not needed for a $\hat{3}$-line). This can be viewed as an optimization procedure: we have a notescape of candidate tones with various levels of structure indicated by the `level` property, and we must pick the descending sequence that maximizes, say, the sum of the levels of structure of each individual tone. This is a variant of the *longest decreasing subsequence problem*, where the items are weighted, and can be solved with dynamic programming. After the *Urlinie* is found, the last remaining note to locate is the initial tone of the bass arpeggiation, which is contained in the initial tonic chord of the composition. Note that as the analysis proceeds, levels of structure may change, so it is possible to run this procedure whenever a new level is created in the analysis.

5 A SHORT EXAMPLE

We illustrate running a few of IVI's implemented components on the first eight measures of Schubert's *Impromptu in B♭ major*, Opus 142, Number 3, depicted in Figure 2. IVI's extraction of the soprano line is shown in Figure 3. Notice how IVI correctly separated the voices in the implied polyphony, leaving only the upper voice in the right

hand part. Though it may appear that in measures 5–7, IVI simply took the top note of the right-hand octave doubling, IVI actually detected this doubling (as described earlier) and transfered the register of the soprano voice up an octave.

IVI also correctly locates the manifestation of the *Ursatz* at this level of the *Impromptu*, as displayed in Figure 4. For the primary tone, IVI chose the D6 in measure 5 over the D5 in measure 1 as it prefers later-occurring notes to earlier ones when structural levels cannot be distinguished, as was the case here. (We use scientific pitch notation; C4 is middle C.) We told IVI to look for a $\hat{3}$-line as the *Ursatz*-finding algorithm currently requires specifying the scale degree of the primary tone.

6 FUTURE WORK

6.1 Multiple Parsings

There is no single correct Schenkerian analysis for any given musical composition. Because this type of analysis is based in part on how a listener perceives a piece, different analysts may produce (sometimes significantly) different analyses. For example, a single piece may admit two different types of *Urlinie* (e.g., a $\hat{3}$-line and a $\hat{5}$-line).

IVI must detect when there are multiple musically-valid interpretations of some collection of notes in a piece and consider possible analyses continuing from each of the viewpoints. At the crux of this issue is the problem of conflicting reductions, where two or more reductions can be applied but not independently; applying one precludes the later possibility of applying another. Furthermore, there is also the issue of backtracking if either reduction should lead to a non-musically-valid analysis later on in the process. This is inevitable even when humans perform Schenkerian analysis; reductional decisions can have far-reaching consequences and may occasionally lead to dead ends.

We are considering a number of approaches to the multiple parsings problem. Weighted preference rules are a standard option, having been used with success by Lerdahl and Jackendoff [8], Temperley and Sleator [17], and Temperley [15] in their music analysis formalisms. Our current model for preference rules uses a weighted system for measuring the amount of "evidence" supporting various reductions. The intelligent search framework around which IVI is based can use this evidence metric in its heuristics to choose which analysis paths to follow. The backtracking component of intelligent search gives us a method for handling multiple interpretations and also any dead ends encountered: because the search proceeds down multiple possible paths at once by saving intermediate states in a queue, the algorithm is always currently investigating the state most likely to lead to the most musically-probable analysis.

6.2 Subproblem Interactions and Levels of Influence

Music analysis involves multiple interacting subproblems of various levels of influence. It is impossible to isolate and solve each problem in a vacuum; they are intertwined and cannot (and should not) be separated. This, however, creates a situation of seemingly circular dependences among the problems. For example, doing a complete harmonic analysis without knowing anything about the melodic and contrapuntal aspects of a composition is challenging at best, whereas an analysis of the voice leading of the piece is informed greatly by knowing the underlying chord structure. Music theorists analyze compositions opportunistically, solving the easier cases within a subproblem before tackling the harder ones (within the same subproblem or a different one), as often the easier instances will shed new light on how to solve the harder cases.

Every reduction in IVI has some amount of evidence associated with it. We have already discussed using this evidence measurement to handle the multiple parsings problem; it gives us a method for choosing between multiple non-independent reductions. The evidence for various possible reductions and property assignments, however, is an ever-fluctuating quantity. Each reduction performed has the potential to guide new ones, and so these evidence variables must be updated after every reduction. This suggests a method for improving the performance of the preliminary analyses (melodic, harmonic, rhythmic, and cadential), as one can execute an iterative process by which the next most likely property attribute-value pair is added at every step, and evidence values are updated, possibly causing changes to other property values. This contrasts with other methods for, e.g., harmonic analysis, where often all of the chord labels are assigned in an order that is independent of the actual music being examined.

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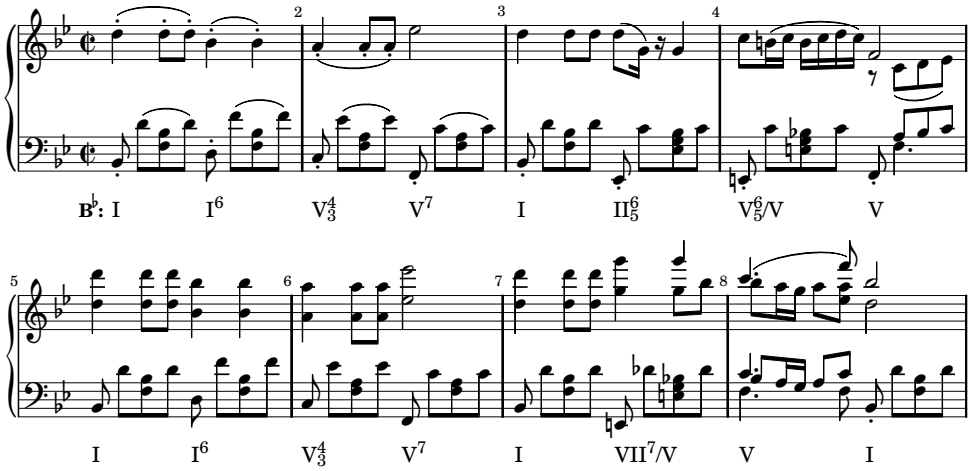

Figure 2. The input score to IVI, with harmonic labels.

Figure 3. IVI's extraction of the soprano line. (Numbers above notes indicate corresponding measures from Figure 2.)

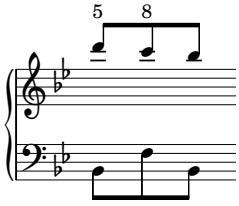

Figure 4. IVI's extraction of the *Ursatz*. (Numbers above notes indicate corresponding measures from Figure 2.)

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A MUSIC IDENTIFICATION SYSTEM BASED ON CHROMA INDEXING AND STATISTICAL MODELING

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ABSTRACT

A methodology is described for the automatic identification of classical music works. It can be considered an extension of fingerprinting techniques because the identification is carried out also when the query is a different performance of the work stored in the database, possibly played by different instruments and with background noise. The proposed methodology integrates an already existing approach based on hidden Markov models with an additional component that aims at improving scalability. The general idea is to carry out a clustering of the collection to highlight a limited number of candidates to be used for the HMM-based identification. Clustering is computed using the chroma features of the music works, hashed in a single value and retrieved using a bag of terms approach. Evaluation results are provided to show the validity of the combined approaches.

1 INTRODUCTION

This paper presents a methodology and its Web-based implementation for the automatic identification of music works. The approach is an extension of *audio fingerprinting* techniques. As it is well known, an audio fingerprint is a unique set of features that allows to identify digital copies of a given audio file because it is robust to the presence of noise, distortion, lossy compression and re-sampling. Identification is carried out by comparing the fingerprint of the unknown audio file with the fingerprints stored in a database, normally using indexing techniques. A comprehensive tutorial on methods and techniques for audio fingerprinting can be found in [3].

The main difference between the proposed approach and typical fingerprinting techniques is that the goal is to identify any possible recording of a given music work through statistical modeling. This representation is automatically computed from audio recordings, which are stored in a database. To this end, the methodology is strictly related to research work on *cover identification*. Yet, the envisaged application scenario is the automatic identification of baroque, classical, and romantic music or of any music genre where a written score is assumed to be the starting point of performances. In all these cases, the term "cover" is misleading because it assumes the existence of a prototype recording from which all the others derive. For this reason, the more general term *music identification* is used in this paper; for simplicity, the term "classical music" is used to address all the aforementioned repertories.

An automatic system able to identify performances of classical music may be helpful in a variety of situations. For instance, most theaters and concert halls routinely record all the rehearsals. This valuable material may totally lack metadata, with the possible exception of the date of the rehearsals, and the reconstruction of how a particular production has been created may become a difficult task. Similar considerations apply to the recordings of concerts that have been broadcasted and archived by radio and television companies. Most of the times, metadata are uninformative about the content of each single audio track, if not totally missing. From the end user point of view, automatic labeling of classical music can be helpful because many compositions have been recorded in a number of different CDs that, apart from the recordings made by well known artists, may not be listed in online services such as Gracenote [9], in particular for the typical case of compilations containing selected movements of recordings already released.

Automatic identification of classical music can be considered also a viable alternative to fingerprinting. In order to effectively identify a recording, fingerprinting techniques need to access the original audio files. For classical music this means that all the different recordings of a given work should be present, with an increase in computational costs, not mentioning the economical aspects in creating such a large reference collection.

The proposed methodology integrates an approach for the identification of orchestral live music [14] based on hidden Markov models (HMMs). A HMM is generated for each music work, starting from an audio recording. The approach showed to be effective but not scalable, because the recording has to be compared to all the recordings in the database. The identification becomes clearly unfeasible when the database contains a large number of music works.

To this end, an additional component has been designed with the aim of improving scalability. The idea is to carry out a clustering of the collection to highlight a limited number of candidates to be used for the HMM based identifi-

cation. The main requirement of clustering is efficiency, while effectiveness can be modulated by carefully selecting the size of the cluster.

2 CHROMA-BASED DESCRIPTION

Chroma features have been extensively used in a number of music retrieval applications. The concept behind chroma is that octaves play a fundamental role in music perception and composition [1]. For instance, the perceived quality – e.g. major, minor, diminished – of a given chord depends only marginally on the actual octaves where it spans, while is strictly related by the pitch classes of its notes. Following this assumption, a number of techniques have been proposed based on chroma features, for chord estimation [7, 18], detection of harmonic changes [11], and the extraction of repeating patterns in pop songs [8]. The application of chroma features to an identification task has been already been proposed for classical [15] and for pop [12] music.

2.1 Indexing

We propose to use chroma features as *indexes* for a collection of classical music recordings. These recordings are stored in a database system, and are used to create statistical models aimed at identification as described in Section 3. The basic idea is that information retrieval techniques can be generally employed outside the textual domain, because the underlying models are likely to be shared by different media [13]. Chroma features are considered as pointers to the recordings they belong to, playing the same role of words in textual documents. The information on the time position of chroma features is used to directly access to relevant audio excerpts.

One major advantage of indexing in text retrieval is that the list of index terms can be accessed in logarithmic, or even constant, time. The same cannot apply to feature vectors, because exact match has to be replaced by similarity search, which is less efficient. A number of research works has been carried out to effectively carry out similarity search in high dimensional spaces, achieving good results in different tasks such as K-nearest neighbors search, clustering, and range searches [20]. In particular, a promising technique is Locality Sensitive Hashing [6]. This techniques aims at applying to feature vectors a carefully chosen hashing function with the aim of creating collisions between vectors that are close in the high dimensional space. The hashing function itself becomes then an effective tool to measure the similarity between two vectors.

It can be noted that fingerprint techniques, such as the one described in [10], can be considered variants of Locality Sensitive Hashing. Following this idea we propose to represent the 12-dimensional chroma vector with a single integer value, obtained through an hashing function. In particular, the actual hash depends on the ranks of the chroma pitch classes, and on on their absolute values.

More formally, a chroma vector $c(i)$ is an array of pitch classes, where $i \in \{1 \dots 12\}$, which are computed from the Fourier transform $S(f)$ of a windowed signal according to equation

$$c(i) = \sum_f B_i(f) \cdot S(f) \tag{1}$$

where $B_i(f)$ is a bank of bandpass filters, each filter centered on the semitones belonging to pitch class i and with a bandwidth of a semitone. The filterbanks may range over all the audible octaves, even if it is common practise to limit its support. In our case the minimum and maximum frequencies are respectively 20 and 5000 hertz.

Once vector $c(i)$ is computed for a window of the signal, we proposed to compute its rank-based quantization $q(i)$ through the following rule

$$q(i) = \begin{cases} rank[c(i)] & if\ \ rank[c(i)] \leq k \\ 0 & elsewhere \end{cases} \tag{2}$$

where k is the number of quantization levels that are taken into account and $rank[c(i)]$ is a function that outputs the rank of the value of the energy in pitch class i over the values of the whole array, that is $rank[c(j)] = 1$ if pitch class j has the highest energy and so on.

Finally, a non negative integer hash h is computed from $q(i)$ according to equation

$$h = \sum_{i=1}^{12} k^{i-1} q(i) \tag{3}$$

where common hashing techniques can be additionally applied to store the values h in one array, which can be accessed in constant time. In particular, we simply compute the reminder of h divided by a carefully chosen prime number.

Indexing is applied both to the recordings in the database and to the queries. One of the problems that may affect retrieval effectiveness is that chroma-based representation is sensible to transpositions. If the recording used as a query and the recording stored in the database are played in different tonalities, they have totally different sets of chroma. We addressed this problem by considering that a transposition of s semitones will result in a rotation of vector $c(i)$ of s steps (the direction of the rotation depending on the direction of the transposition). The value of h^s of quantized chroma h transposed by s ascending semitones, where $s \in \{1 \dots 11\}$ can be computed through equation

$$h^s = k^{12-s}(h \ \% \ k^s) + \lfloor \frac{h}{k^s} \rfloor \tag{4}$$

where $\%$ gives the reminder of a division and $\lfloor \cdot \rfloor$ is the floor function. Thus a query is represented by twelve sets

of hashes H^s, that is the representation H^0 in the original tonality and eleven additional ones that take into account all the possibile transpositions. This representation can be considered as an application of query extension techniques.

2.2 Retrieval

With the main goal of efficiency, continuing the parallel with textual information retrieval techniques, retrieval is carried out using the *bag of words* paradigm. Thus the similarity between the query and the recordings in the collection is carried out by simply counting the number of hashes of quantized chroma they have in common. No attempt to align, not even to consider the relative ordering of the quantized chroma, is made. The only additional processing is to highlight the excerpts of each recordings that potentially correspond to the query. This is simply made by using the timing information contained in the indexes to divide the recordings in overlapping frames.

The similarity M_q between the query and the recordings is computed by simply counting the number of hashes they have in common, that is by performing an intersection between the set H^0 representing the query – or one of its transposed versions H^s with $s \in \{1 \dots 11\}$ – with the set H_{jf} representing the frame f of the recording j in the database. This can be expressed through equation

$$M_q(s, j, f) = \|H^s \cap H_{jf}\| \tag{5}$$

where $||\cdot||$ returns the cardinality of a set, that is the number of its elements. The approach allows us to sort the recordings in the dataset according to the value of M and, at the same time, to highlight the frames f that more likely correspond to the query and the difference in semitones between the two tonalities.

The proposed similarity value is not very common in information retrieval, where more complex measures are exploited, such as the cosine of the angle between the vectors representing the query and the items in the database. Moreover, the similarity does not take into account the number of occurrences of a given hash value within the query and the recordings, which is normally exploited through the $tf \cdot idf$ weighting scheme. The choice of this particular similarity measure has been motivated by a number of tests showing that this simple measure outperformed more complex ones. It has to be considered that the distribution of a given hash value inside the database, which is taken into account by the *inverse document frequency* element idf, does not have a clear effect on the similarity between two recordings of the same music work. At the same time, the distribution of a given hash within a recording, the *term frequency* element tf, can be affected by the rate at which chroma vectors are computed and give different similarity values depending on the difference in tempo between two recording.

The top A retrieved recordings are considered as the cluster of candidates that are used in the second step, as discussed in the next section. It is important to note that quantized chroma are used only in the clustering step, where recall is much more important than precision. It is likely that more sophisticated techniques will increase precision, by assigning a higher rank to the recording that corresponds to a query, but will have a minor impact on recall.

2.3 Chroma Evaluation

The indexing methodology has been evaluated with a testbed composed by a database of 1000 recordings and by a query set of 50 different performances of a subset of the works in the database. The audio files were all polyphonic recordings of orchestral music, with a sampling rate of 44.1 kHz. We segmented the recordings in frames, as discussed in the previous section, with a length of 10 seconds. Subsequent frames had an overlap of 50%. Queries were made of audio excerpts of about the half of the length of the frames, taken at random within the audio query files. Testing was carried out by extracting random frames of the 50 queries, matching them with all the frames of the recordings in the database and measuring the rank of the recording that corresponded to the query. This experimental setup aims at simulating a real situation, where the user can query the system using any part of an unknown recording. The purpose of the evaluation was to prove the validity of hashes of the quantized chroma features as index terms, to highlight the best configuration of the parameters, and to select the size of the cluster.

We tested the effectiveness of the approach depending on the size of the window used to compute the chroma features. According to this value, the precision of chroma features would increase or decrease, affecting the clustering rates. We tested the system with different window sizes, while the hopsizes between windows consistently set to $1/4$ of the window size. Results showed that the optimal window length, with this experimental setup, is 1024 points (about 23 milliseconds). Moreover, we tested different sizes of the hashing map by varying the quantization level k, that is the number of pitch classes that are used to compute the hashing function, as explained in Section 2.1. The results highlighted the optimal number of levels used for quantization, which is $k = 6$. With this value, and with this experimental setup, a cluster of the first 100 recordings is enough to assure that the correct document will be processed in the following HMM-based identification step. A cluster size of only 50 elements contains the correct document in 96% of the cases.

These results are particularly encouraging, because they are obtained with a compact representation of audio recordings – which allows us to compute an exact match between features – and a retrieval based on a simple bag of terms approach – which does not require any alignment technique.

Table 1 reports the percentage of times the correct recording has been ranked within predefined thresholds.

method	$=$ **1**	$\leq$ **10**	$\leq$ **50**	$\leq$ **100**	$\leq$ **200**
chroma	69	88	96	100	100

Table 1. Identification rates using the quantized chroma

3 A COMPLETE IDENTIFICATION SYSTEM

The first system architecture was proposed in [14], where the results of a set of experimental tests have been presented. Identification is based on a *audio to audio* matching process, which goal is to retrieve all the audio recordings that represents the same music content as the audio query. The audio recordings used for the identification are stored in a relational database, together with a set of relevant metadata. Audio to audio matching has been also proposed in [15], where a chroma representation was used to identify classical music. Another application of chroma features to an identification task, this time on pop music, has been proposed in [12] where chroma features of the query song and the songs in the database are aligned using Dynamic Time Warping (DTW). Global similarity, obtained through DTW, between audio excerpts has been used in [5] to align in real time two different performances of the same music work.

The current retrieval architecture, which is depicted in Figure 1, is based on two main components: a clustering block (on the left) that selects efficiently a number of candidate compositions and an identification block (on the right) that computes the probability that the audio query correspond to each of the candidates.

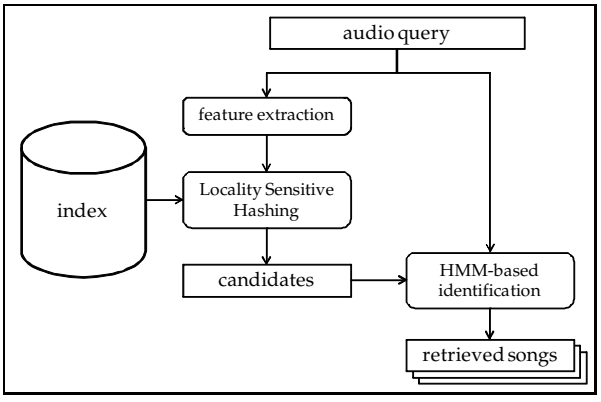

Figure 1. Structure for a music identification system

In this section, for the sake of completeness, the main elements of the identification block are briefly summarized. A more detailed description can be found in [14]. In a first step, the segmentation process extracts audio subsequences that have a coherent acoustic content. Segmentation exploits a technique that has been proposed for the segmentation of textual documents in coherent parts, a task also called topic segmentation [4]. The aim of segmentation is to divide the music signal into subsequences that are bounded by the presence of music events, where an event occurs whenever the current pattern of a music piece is modified, typically one or more new notes being played or stopped.

Coherent segments of audio are analyzed through a second step in order to compute a set of acoustic parameters that are general enough to match different performances of the same music work. To this aim each segment needs to be described by a compact set of features that are automatically extracted. Considering pitch as the most relevant parameter for music identification and because pitch is related to the presence of peaks in the frequency representation of an audio frame, the parameter extraction step is based on the computation of local maxima in the Fourier transform of each segment, averaged over all the frames in the segment.

In a final step a HMM is automatically built from segmentation and parameterization to model music production as a stochastic process. HMMs are stochastic finite-state automata, where transitions between states are ruled by probability functions. At each transition the new state emits a random vector with a given probability density function [19]. The basic idea is that music recordings can be modeled with HMMs providing that states are labeled with events in the audio recording and their number is proportional to the number of segments, transitions model the temporal evolution of the audio recording and observations are related to the audio features previously extracted that help distinguishing different events.

At identification time, an unknown recording of a performance is pre-processed in order to extract the features modeled by the HMMs. All the models are ranked according to the probability of having generated the acoustic features of the performance to be identified.

3.1 Evaluation of the Complete System

We evaluate the effectiveness and the efficiency of a system implementing the methodology by using the same experimental collection described in Section 2.3. Clustering has been carried out by selecting, for each query, 100 frames of the recordings in the database according to their similarity with the query or its transposed versions. Hashes have been computed using an analysis window of 1024 points and using $k = 6$ levels of quantization. The queries had a length of 7 seconds, while recordings in the database have been segmented in overlapping frames of 10 seconds.

Results are shown in Table 2, which compares the effectiveness of the complete approach with the one based on HMM. As it can be seen, clustering improves effectiveness because the complete method gave a Mean Reciprocal Rank

(MRR) of 90.2, which is significantly higher than MRR for the pure HMM-based approach. Moreover, for the complete approach, every query gives the correct work ranked among the first 10 positions, which is an important result for many of the applications of supervised music identification that have been described in Section 1.

method	$=$ **1**	$\leq$ **3**	$\leq$ **5**	$\leq$ **10**	$\leq$ **20**	**MRR**
only HMM	84	88	90	94	94	86.7
complete	86	94	94	100	100	90.2

Table 2. Identification rates using the HMM-based approach and the complete methodology

The complete approach gives clear improvements also on efficiency. Because chroma-based clustering can be carried out efficiently, its computational cost is negligible compared to HMM-based identification. Thus the reduction of the number of candidates gives an improvement of orders of magnitude, because the cluster contains a fixed number of excerpts of recordings while, without clustering, the query should compared against all the experts of all the recordings. In our experiments, identification of an audio excerpts of about 7 seconds is carried out in 2.5 seconds.

It is interesting to note that most of the queries have a different tempo than the corresponding recordings in the database. The approach seems to be robust to tempo differences, even if commercial recordings, especially of orchestral music, do not present large deviations from the tempo written in the score. In order to test robustness to tempo, queries have been stretched from 0.8 to 1.2 of their original bpms (larger deviations were totally unrealistic with this particular dataset). Results with this artificial dataset did not showed changes in the identification rate, yet more experiments should be carried out with real data to effectively measure the robustness to tempo deviations

3.2 A Prototype for Music Identification

The system in Section 3 has been implemented in a Web-oriented prototype available on-line [16], which works as a demonstrator of the proposed approach. The core engine of the application, which include indexing, modeling and identification steps, has been implemented in C++. The music collection is composed by the indexed music recordings and is stored in a PostgreSQL database, including both the metadata and the music content information. A number of queries are available for the user to test the system, including some recordings where different noisy sources have been added to the original queries, such as environmental noises, and typical noises of analog equipments (i.e., clicks, hisses). Additional queries are available with piano solo version of orchestral music and dramatic changes in tempo, both faster and slower than the original, in order to show the effect of changes in orchestration and in tempo, although the approach has not been tested extensively with changes in orchestration.

All the examples are coded with lossy compression, in order to be easily listened also through the Web. The user can access to the metadata associated to the recordings in the database and listen to the rank list of music works returned by the prototype. For this reason, only copyright free recordings are stored in the database, which thus has a limited size of about 1000 music works.

At the current state the prototype allows to test the system with the supplied examples, by visualizing the first 20 results of the whole rank list. In order to find the right match, the user can listen to the resulting elements to find the most similar one. Considering all the experimental results reported in the previous sections, the user has good probability of listening just the first retrieved music works to find the relevant one. The approach is thus to perform a supervised identification, because the final choice of which is the correct music work is left to the user.

4 CONCLUSIONS

This paper describes a methodology for the identification of works of classical music. It is based on two steps. At first it carries out a clustering of a database of recordings in order to select a set of candidates music works. Clustering is based on the introduction of an hashing function that maps the chroma features on integer numbers. The second step computes the similarity between the audio query and the recordings in the database by exploiting an application of HMMs. Thus identification is carried out using two different strategies, the first to achieve efficiency, the second to refine effectiveness.

The methodology has been implemented in a Web-based prototype that allows the user to identify even short audio excerpts of classical music. The identification rate, which has been computed using a database of 1000 recordings of orchestral music, is encouraging. Already using the first step, the system was able to correctly identify the query in 69% of the times. The second step introduces an additional refinement of the identification rate that reaches, in our experiments, 86%.

It is important to note that the proposed approach assumes that different performances are based on a similar music score, even if played with different instrumentation, tempo, or key signature. The system has not been tested on music genres where improvisation is a predominant feature, such as jazz music, or where the music score can be substantially modified through arrangements. Yet, some results on the effectiveness of the individual techniques show that the general approach could be extended also, at least, to pop and rock music. For instance, many works on cover song

identification are based on the use of chroma vectors [18] while HMM-based identification has been applied also to pop music with encouraging results [17].

It could be argued that the proposed methodology can hardly be extended to the identification of recordings of all the music genres. One of the lessons learnt by the information retrieval community is that some of the techniques for text retrieval (i.e., stemming), are not equally effective for all the languages and all the retrieval tasks. Nevertheless, we aim at extending the proposed approach to other music genres, by introducing different descriptors that are related to the main features of each music genre. For instance, the main melody in the case of pop and rock or the chord progression in the case of jazz.

5 ACKNOWLEDGMENTS

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RHYME AND STYLE FEATURES FOR MUSICAL GENRE CLASSIFICATION BY SONG LYRICS

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ABSTRACT

How individuals perceive music is influenced by many different factors. The audible part of a piece of music, its sound, does for sure contribute, but is only one aspect to be taken into account. Cultural information influences how we experience music, as does the songs' text and its sound. Next to symbolic and audio based music information retrieval, which focus on the sound of music, song lyrics, may thus be used to improve classification or similarity ranking of music. Song lyrics exhibit specific properties different from traditional text documents – many lyrics are for example composed in rhyming verses, and may have different frequencies for certain parts-of-speech when compared to other text documents. Further, lyrics may use 'slang' language or differ greatly in the length and complexity of the language used, which can be measured by some statistical features such as word / verse length, and the amount of repetative text. In this paper, we present a novel set of features developed for textual analysis of song lyrics, and combine them with and compare them to classical bag-of-words indexing approaches. We present results for musical genre classification on a test collection in order to demonstrate our analysis.

1 INTRODUCTION

The prevalent approach in music information retrieval is to analyse music on the symbolic or audio level. Songs are therein represented by low level features computed from the audio waveform or by transcriptions of the music. Additionally, all songs but instrumental tracks can be treated as textual content by means of song lyrics. This information can be exploited by methods of classical text information retrieval to provide an alternative to music processing based on audio alone. On the one hand, lyrics provide the means to identify specific genres such as 'love songs', or 'Christmas carols', which are not acoustic genres per se, but, to a large degree, defined by song lyrics [10]. Christmas songs, for instance may appear in a wide range of genres such as 'Punk Rock' or 'Pop' in addition to the classic 'Christmas carol'. On the other hand, song lyrics might sometimes be the only available option for extracting features, for example, when audio files are not available in an appropriate format, or only via a stream when bandwith is a limiting factor. Further, processing of audio tracks might be too time consuming compared to lyrics processing.

Song lyrics may differ to a great extent from the documents often dealt with in traditional text retrieval tasks such as searching the web or office documents. In addition to its plain text content, song lyrics exhibit a certain structure, as they are organised in blocks of choruses and verses. Also, lyrics might feature other specific properties, such as slang language in 'Hip-Hop' or 'Rap' music, or other statistical information such as (average) line lengths, or words per minute. Also special characters, e.g. the number of exclamation marks used, might be of interest.

In this paper, we present a set of features composed of these various textual properties. We then use them for genre classification, and compare and combine them with features resulting from standard bag-of-words indexing of the song texts. We aim to show the following: a) rhyme, part-of-speech, and simple text statistic features alone can be used for genre classification, and b) the combination of bag-of-words features and our feature sets is worthwile.

The remainder of this paper is structured as follows. We first give an overview of related work on music information retrieval focusing on lyrics processing in Section 2. Then, we give an introduction to lyrics analysis and explain our feature sets in detail in Section 3. In Section 4, we report from experiments performed on a manually compiled collection of song lyrics. Finally, we draw conclusions and give an overview of future work in Section 5.

2 RELATED WORK

In general, music information retrieval (MIR) is a broad and diverse area, including research on a magnitude of topics such as classic similarity retrieval, genre classification, visualisation of music collections, or user interfaces for accessing (digital) audio collections. Many of the sub-domains of MIR – particularly driven by the large-scale use of digital

audio over the last years – have been heavily researched.

Experiments on content based audio retrieval, i.e. based on signal processing techniques, were reported in [3] as well as [14], focusing on automatic genre classification. Several feature sets have since been devised to capture the acoustic characteristics of audio material, e.g. [12].

An investigation of the merits for musical genre classification, placing emphasis on the usefulness of both the concept of genre itself as well as the applicability and importance of genre classification, is conducted in [8].

A system integrating multi modal data sources, e.g. data in the form of artist or album reviews was presented in [1]. Cultural data is used to organise collections hierarchically on the artist level in [11]. The system describes artists by terms gathered from web search engine results. A promising technique for automatic lyrics alignment from web resources is given in [4].

A study focusing solely on the semantic and structural analysis of song lyrics including language identification of songs based on lyrics is conducted in [6]. Song lyrics can also be feasible input for artist similarity computations as shown in [5]. It is pointed out that similarity retrieval using lyrics, i.e. finding similar songs to a given query song, is inferior to acoustic similarity. However, it is also suggested that a combination of lyrics and acoustic similarity could improve results, which motivates future research. Genre classification based on lyrics data as well as its combination with audio features is presented in [9].

Further, the combination of a range of feature sets for audio clustering based on the Self-Organising Map (SOM) is presented in [7]. It is shown that the combination of heterogeneous features improves clustering quality. Visualisation techniques for multi-modal clustering based on Self-Organising maps are given in [10], demonstrating the potential of lyrics analysis for clustering collections of digital audio. Similarity of songs is defined according to both modalities to compute quality measures with respect to the differences in distributions across clusterings in order to identify interesting genres and artists.

3 LYRICS FEATURES

In this section we present the feature set computed from the song lyrics, namely bag-of-words, rhyme, part-of-speech, and statistical features.

3.1 Bag-of-Words Features

A common approach in text retrieval is to index documents with the bag-of-words method. Here, each unique term occurring in any of the documents of the collection is regarded a feature. To populate the feature vectors, information about the frequency of occurrences of the terms in the collection is gathered. A simple approach is the Boolean Model, which only considers whether a term is present in a document or not. More sophisticated, one can apply a term weighting scheme based on the importance of a term to describe and discrimante between documents, such as the popular $tf \times idf$ (term frequency $\times$ inverse document frequency) weighting scheme [13]. In this model, a document is denoted by d, a term (token) by t, and the number of documents in a corpus by N. The *term frequency* $tf(t,d)$ denotes the number of times term t appears in document d. The number of documents in the collection that term t occurs in is denoted as *document frequency* $df(d)$. The process of assigning weights to terms according to their importance for the classification is called 'term-weighing', the $tf \times idf$ weight of a term in a document is computed as:

$$tf \times idf(t,d) = tf(t,d) \cdot ln(N/df(t)) \qquad (1)$$

This weighting scheme is based on the assumption that terms are of importance when they occur more frequently in one document, and at the same time less frequently in the rest of the document collection. The bag-of-words approach focuses on grasping the topical content of documents, and does not consider any structural information about the texts. This method tends to, already with a low number of documents, result in high-dimensional feature vectors. Thus, often a feature space reduction is required for further processing, which is often achieved by cutting of words which occur either too often, or too rarely.

3.2 Rhyme Features

A rhyme is a linguistic style, based on consonance or similar sound of two or more syllables or whole words at the end of one line; rhymes are most commonly used in poetry and songs. We assume that different genres of music will exhibit different styles of lyrics, which will also be characterised by the degree and form of the rhymes used. 'Hip-Hop' or 'Rap' music, for instance, makes heavy use of rhymes, which (along with a dominant bass) leads to its characteristic sound. We thus extract several descriptors from the song lyrics that shall represent different types of rhymes.

One important notion is that consonance or similarity of sound of syllables or words is not necessarily bound to the lexical word endings, but rather to identical or similar phonemes. For example, the words 'sky' and 'lie' both end with the same phoneme /ai/. Before detecting rhyme information, we therefore first transcribe our lyrics to a phonetic representation. Phonetic transcription is language dependent, thus the language of song lyrics would first need to be identified, using e.g. TextCat [2] to determine the correct transcriptor. However, as our test collection presented in this paper features only English songs and we therefore use English phonemes only, we omit details on this step.

We distinguish two patterns of subsequent lines in a song text: *AA* and *AB*. The former represents two rhyming lines,

Feature Name	Description
Rhymes-AA	A sequence of two (or more) rhyming lines ('Couplet')
Rhymes-AABB	A block of two rhyming sequences of two lines ('Clerihew)
Rhymes-ABAB	A block of alternating rhymes
Rhymes-ABBA	A sequence of rhymes with a nested sequence ('Enclosing rhyme')
RhymePercent	The percentage of blocks that rhyme
UniqueRhymeWords	The fraction of unique terms used to build the rhymes

Table 1. Rhyme features for lyrics analysis

Feature Name	Description
exclamation_mark, colon, single_quote, comma, question_mark, dot, hyphen, semicolon	simple count
d0 - d9	counts of digits
WordsPerLine	words / number of lines
UniqueWordsPerLine	unique words / number of lines
UniqueWordsRatio	unique words / words
CharsPerWord	number of chars / number of words
WordsPerMinute	the number of words / length of the song

Table 2. Overview of text statistic features

while the latter denotes non-rhyming. Based on these basic patterns, we extract the features described in Table 1.

A 'Couplet' *AA* describes the rhyming of two or more subsequent pairs of lines. It usually occurs in the form of a 'Clerihew', i.e. several blocks of Couplets *AABBCC... ABBA*, or *enclosing rhyme* denotes the rhyming of the first and fourth, as well as the second and third lines (out of four lines). We further measure 'RhymePercent', the percentage of rhyming blocks, and define the unique rhyme words as the fraction of unique terms used to build rhymes 'UniqueRhymeWords'.

Of course, more elaborate rhyming patterns, especially less obvious forms of rhymes, could be taken into account. In order to initially investigate the usefulness of rhyming at all, we do not take into account rhyming schemes based on assonance, semirhymes, or alliterations, amongst others. Initial experimental results lead to the conclusions that some of these patterns may well be worth studying. However, the frequency of such patterns in popular music is doubtful.

3.3 Part-of-Speech Features

Part-of-speech tagging is a lexical categorisation or grammatical tagging of words according to their definition and the textual context they appear in. Different part-of-speech categories are for example nouns, verbs, articles or adjectives. We presume that different genres will differ also in the category of words they are using, and therefore we additionally extract several part of speech descriptors from the lyrics. We count the numbers of: *nouns*, *verbs*, *pronouns*, *relational pronouns* (such as 'that' or 'which'), *prepositions*, *adverbs*, *articles*, *modals*, and *adjectives*. To account for different document lengths, all of these values are normalised by the number of words of the respective lyrics document.

3.4 Text Statistic Features

Text documents can also be described by simple statistical measures based on word or character frequencies. Measures such as the average length of words or the ratio of unique words in the vocabulary might give an indication of the complexity of the texts, and are expected to vary over different genres. The usage of punctuation marks such as exclamation or question marks may be specific for some genres. We further expect some genres to make increased use of apostrophes when omitting the correct spelling of word endings. The list of extracted features is given in Table 2.

All features that simply count character occurrences are normalised by the number of words of the song text to accommodate for different lyrics lengths. 'WordsPerLine' and 'UniqueWordsPerLine' describe the words per line and the unique number of words per line. The 'UniqueWordsRatio' is the ratio of the number of unique words and the total number of words. 'CharsPerWord' denotes the simple average number of characters per word. The last feature, 'WordsPerMinute' (WPM), is computed analogously to the well-known beats-per-minute (BPM) value [1] .

4 EXPERIMENTS

In this section we report on experiments performed for a test collection of 397 song lyrics. We used classical bag-of-words indexing as well as the feature sets introduced in Section 3. Genre classification was done for a range of setups with Naïve Bayes, k-Nearest Neighbour with different values for k, SVMs with linear and polynomial kernels (more complex kernels did not improve accuracy), and a Decision

[1] Actually we use the ratio of the number of words and the song length in seconds to keep feature values in the same range. Hence, the correct name would be 'WordsPerSecond', or WPS.

Genre	Songs	Genre	Songs
Country	29	Pop	42
Folk	44	Punk Rock	40
Grunge	40	R&B	40
Hip-Hop	41	Reggae	33
Metal	46	Slow Rock	42

Table 3. Composition of the test collection

Tree (J48). We used a slightyly modified version of the Weka toolset for running and evaluating our experiments [2] .

4.1 Test Collection

The task of compiling a feasible test collection for genre classification by lyrics is tedious for the preprocessing task must provide correct lyrics both in terms of structure and content. Therefore, all lyrics were manually preprocessed in order to remove additional markup like '[2x]' or '[chorus]', and to include the unabridged lyrics for all songs. We paid special attention to completeness in terms of the resultant text documents being an as adequate and proper transcription of the songs' lyrics as possible.

Starting from a private collection of about 12.000 songs, we selected a random sample of 30-45 songs from each of the ten genres listed below. We removed songs that seemed not feasible because of their length (extremely short or overlength tracks) and removed songs in languages other than English to avoid the complexity that comes with phoneme transcription to other languages. This resulted in a collection of 397 remaining songs.

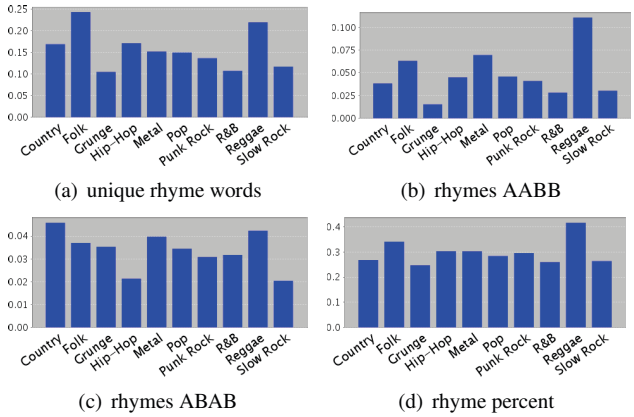
We aimed at having a high number of different artists in order to prevent biased results by too many songs from the same artist; our collection comprises 143 different artists. Table 3 shows the numbers of songs per genre.

The lyrics for all songs were automatically retrieved from the Internet. All songs were manually filtered, e.g. we removed instrumental songs. Annotations like [chorus] or [verse] were substituted by the respective parts of the text, i.e. the full texts of all choruses and verses were placed correctly.

4.2 Feature Analysis

To illustrate the discriminative power of the new feature set, we present how values vary across different genres. Due to space limitations, we selected a subset of four exemplary features from each type of rhyme, part-of-speech, and text statistic features.

In Figure 1, a selected set of Rhyme features is illustrated. The first subplot shows the number of unique words



(a) unique rhyme words (b) rhymes AABB

(c) rhymes ABAB (d) rhyme percent

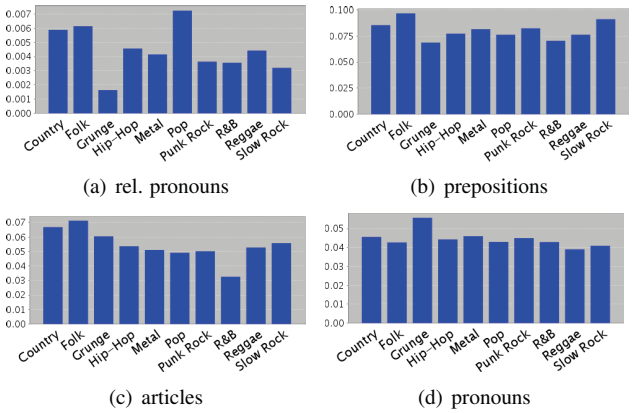
Figure 1. Average values for rhyme features

used to build the rhyme patterns. It can be observed that 'Folk' and 'Reggae' have by far the highest value, while other genres like 'R&B', 'Slow Rock' and 'Grunge' are obviously using less rhyming language. 'Reggae' and 'Grunge' can also be well distinguished by the *AABB* pattern, as well as 'R&B' and 'Slow Rock'. The latter can also be well discriminated against other genres regarding the usage of the *ABAB* pattern, while 'Reggae' is particularly intriguing when it comes to the total percentage of rhyme patterns in its song lyrics. Other rhyme features have similar characteristics, with *AA* patterns having being the least discriminative between the genres.

Figure 2 gives an overview of part-of-speech features, namely relational pronouns, prepositions, articles, and pronouns. Relational pronouns show a pretty strong fluctuation acrros all genres, with 'Grunge' notably having the lowest value, while 'Pop', 'Country' and 'Folk' exhibit the highest values. Articles help to distinguish especially 'R&B' from the other features by having a much lower average than the other genres. To some extent, also 'Folk' can be discriminated by having the highest value. Pronoun usage is mostly equally distributed, but 'Grunge' stands out with a much higher value. Preposition also do not show a huge variation across the different classes. For the other features, nouns, verbs and adjectives have very similar values, while both adverbs and modals do show some fluctuations but do not significantly differ across genres.

An overview of the most interesting statistic features is presented in Figure 3. Genres like 'Reggae', 'Punk Rock' and 'Hip-hop' use an expressive language that employs a lot of exclamation marks. Similar, 'Hip-Hop', 'Punk Rock' and 'Metal' seem to use a lot of digits in their lyrics. This can be explained partly with the names of other artists mentioned in the songs, as in many 'Hip-Hop' pieces of '50 Cent' or '2 Pac'. Also, 911 is an often mentioned number. 'Folk', 'Hip-Hop', and to a lesser extent also 'Country' seem to be characterised by more creative lyrics, which

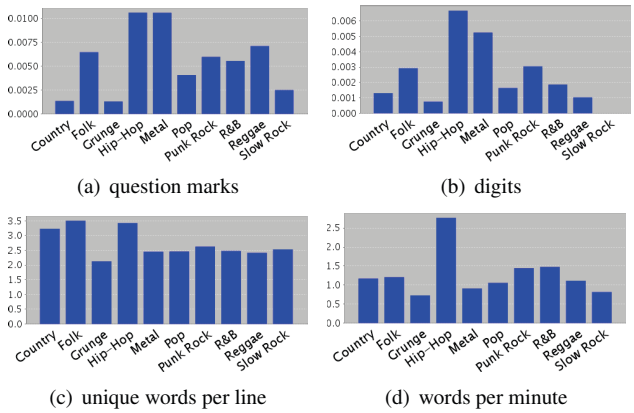
[2] http://http://www.cs.waikato.ac.nz/ml/weka/



(a) rel. pronouns (b) prepositions

(c) articles (d) pronouns

Figure 2. Average values for part-of-speech features



(a) question marks (b) digits

(c) unique words per line (d) words per minute

Figure 3. Average values for statistic features

manifests in a bigger word pool used in these genres. Last, 'Words per Minute' is the most discriminative feature for 'Hip-Hop', which indeed is characterised by fast language, and in most of the cases has very short lead-in and fade-out sequences. Also 'R&B' and 'Punk Rock' have these characteristics, while 'Grunge', 'Slow Rock' and 'Metal' feature long instrumental parts or a more moderate singing tempo.

4.3 To Stem or Not to Stem

We used Porter's stemming algorithm to remove word endings. Its impact on classification accuracies varies; in some cases, it lowers classification accuracy. We performed experiments for a standard Naïve Bayes classifier, and Support Vector Machines (SVM) with linear kernel and Decision Trees (J48) for three different dimensionalities computed by document frequency thresholding feature selection. For the Bayes classifier and SVMs, stemming improved the results for all three dimensionalities (full set, 3000, 300). For the Decision Tree setup, it did not yield improvements. Overall, we did not find significant differences in accuracies, yet, we had expected setups without stemming to be superior caused by word endings in 'slang' language.

4.4 Classification Results

We performed a set of classification experiments, a summarisation of the results is given in Table 4. It shows the experiments we performed on the test collection, omitting results on different values for k for the k-Nearest Neighbour and kernels other than the linear one for SVMs, as they showed the same trends, but had a worse overall performance. For a set of ten genres, we assume the absolute baseline for classification experiments to be the relative number of tracks in the largest class, i.e. $46/397 = 11.59\%$. However, we rather consider the results obtained with the bag-of-words approach as the baseline to compare to. The values given are macro-averaged classification accuracies for the correctly classified tracks computed with ten-fold cross validation. We want to point out that classification accuracies are lower than comparable results based on audio content, yet we show that genre classification by lyrics features can be complementary for they might classify different genres better than audio only.

For all three classifiers, feature sets including our proposed features achieved the best results (experiments 6, 4, and 3). The addition to the bag-of-words approach (with stemming) always yields better results, this becomes even clearer in the case of the SVM (experiments 6, 8, and 10). The best result of all experiments was achieved with a dimensionality of 38, which is sa significantly lower dimensionality than the one of the best performing bag-of-words approach (experiment 6, dimensionality: 3336). We want to point out the improvement in performance of about 7.5% absolute between 24.83 and 33.47 (relative increase of about a third). We also performed the same set of experiments with non-stemmed bag-of-words features and combinations thereof; they revealed the same trends but are omitted due to space limitations.

5 CONCLUSIONS

In this paper, we introduced style and rhyme features for lyrics processing. We showed that all our feature sets outperform the baseline in terms of classification accuracies in a set of experiments. Further, we showed that the combination of style features and classical bag-of-words features outperforms the bag-of-words only approach and in some cases reduces the needed dimensionality.

These results indicate that style and rhyme features can contribute to better music classification or a reduction in dimensionality for similar classification accuracies. Hence, we plan to investigate more sophisticated rhyme patterns. As the classification experiments reported in this paper were based on a simple concatenation approach, more elaborate

Exp.	Feature Combination	Dim	5-NN		Naïve Bayes	SVM		Decision Tree	
1	rhyme	6	13.17		13.67	− 12.83	−	13.82	-
2	part-of-speech (pos)	9	15.99		20.79	16.04	−	15.89	-
3	text statistic	23	**28.53**	+	20.6	− 30.12		**26.72**	
4	rhyme / pos / text statistic	38	25.33	+	23.37	**33.47**	+	24.60	
5	bag-of-words #1	3298	13.63		25.89	24.83		23.63	
6	bag-of-words / rhyme / pos / text statistic #1	3336	13.88		**27.58**	29.44	+	24.39	
7	bag-of-words #2	992	12.34		25.13	21.96		23.11	
8	bag-of-words / rhyme / pos / text statistic #2	1030	16.56		25.78	27.37		23.77	
9	bag-of-words #3	382	14.06		22.74	22.27		22.17	
10	bag-of-words / rhyme / pos / text statistic #3	420	15.06		24.50	29.36		24.05	

Table 4. Classification results for different feature combinations. The highest accuracies per column are printed in bold face; statistically significant improvement or degradation over the base line experiment (5, column-wise) is indicated by (+) or (−), respectively

methods such as ensemble learning will be taken into consideration. Finally, we plan on improve lyrics preprocessing by heuristics for rhyme detection and automatic alignment thereof.

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AUTOMATIC CHORD RECOGNITION BASED ON PROBABILISTIC INTEGRATION OF CHORD TRANSITION AND BASS PITCH ESTIMATION

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ABSTRACT

This paper presents a method that identifies musical chords in polyphonic musical signals. As musical chords mainly represent the harmony of music and are related to other musical elements such as melody and rhythm, the performance of chord recognition should improve if this interrelationship is taken into consideration. Nevertheless, this interrelationship has not been utilized in the literature as far as the authors are aware. In this paper, bass lines are utilized as clues for improving chord recognition because they can be regarded as an element of the melody. A probabilistic framework is devised to uniformly integrate bass lines extracted by using bass pitch estimation into a hypothesis-search-based chord recognition. To prune the hypothesis space of the search, the hypothesis reliability is defined as the weighted sum of three reliabilities: the likelihood of Gaussian Mixture Models for the observed features, the joint probability of chord and bass pitch, and the chord transition N-gram probability. Experimental results show that our method recognized the chord sequences of 150 songs in twelve Beatles albums; the average frame-rate accuracy of the results was 73.4%.

Keyword: chord recognition, bass line, hypothesis search, probabilistic integration

1 INTRODUCTION

In recent years, automatic recognition of musical elements such as melody, harmony, and rhythm (Figure 1) from polyphonic musical signals has become a subject of great interest. The spread of high-capacity portable digital audio players and online music distribution has allowed a diverse user base to store a large number of musical pieces on these players. Information on the content of musical pieces such as their musical structure, mood and genre can be used together with text-based information to make music information retrieval (MIR) more efficient and effective. Manual annotation requires an immense amount of effort, and maintaining a consistent level of quality is not easy. Thus, techniques for extracting musical elements are essential for obtaining content-based information from musical signals.

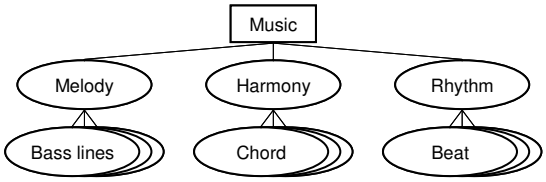

Figure 1. Musical elements

A key principle in analyzing musical signals is that musical elements are related to each other. Because composers exploit the interrelationship among musical elements, this interrelationship should be considered when analyzing the elements as well. Most studies in the literature have dealt with these elements independently.

This paper exploits the interrelationship between chord sequences and bass lines to improve the performance of chord recognition. The chord sequence is regarded as an element of the harmony, while bass lines are regarded as an element of the melody. The chord sequence consists of a chord symbol sequence and chord boundary sequence. As the chord sequence may represent the mood of music, it can be used to calculate the similarity in mood between musical pieces. This similarity is important in MIR and music recommendation. On the other hand, the bass line represents a melody in the bass register; thus, it leads the chord progression.

A recent approach adopted recently by many researchers for automated description of the chord sequence is the use of Hidden Markov Models (HMMs). Several methods have been suggested to explore the analogy between speech recognition and chord recognition and to consider the temporal connection of chords [1, 2, 3]. Sheh *et al.* [1] proposed a method that uses the extended Pitch Class Profile (PCP) [4] as a feature vector. They used an HMM that had one state per chord with a large set of classes (147 chord types). However, they were not able to obtain good enough results for recognition. Bello *et al.* [2] used chroma features and an HMM; they improved accuracy by incorporating musical

knowledge into the model. Lee *et al.* [3] built key-specific models for automatic chord transcription. They used a 6-dimensional feature vector, called Tonal Centroid that is based on Tonnetz [5]. Higher accuracies were obtained by limiting the number of chord types that could be recognized.

Yoshioka *et al.* pointed out that chord symbols affect chord boundary recognition and vice versa [6]. They developed a method that concurrently recognizes chord symbols and boundaries by using a hypothesis search that recognizes the chord sequence and key.

While previous studies have treated only the features of chords, we focus on the interrelationship among musical elements and integrate information about bass lines into chord recognition in a probabilistic framework. The framework enables us to deal with multiple musical elements uniformly and integrate information obtained from statistical analyses of real music.

This paper is organized as follows: Section 2 describes our motivation for developing an automatic chord recognition system, the issues in involved in doing so, and our solution. Section 3 explains our method in concrete terms. Section 4 reports the experimental results and describes the effectiveness of our method. Our conclusions are discussed in Section 5.

2 CHORD RECOGNITION USING BASS PITCH ESTIMATION

2.1 Motivation

A bass line is a series of tonal linear events in the bass register. We focus on it specifically because it is strongly related to the chord sequence. Bass sounds are the most predominant tones in the low frequency region, and bass lines have the following important properties:

- They are comprised of the bass register of the musical chords.
- They lead the chord sequence.
- They can be played with a single tone and have a pitch that is relatively easy to estimate.

We improve chord recognition by exploiting the above properties. Information about bass lines is obtained through bass pitch estimation.

To process multiple musical elements simultaneously, we use a probabilistic framework that enables us to deal with them uniformly because each evaluation value is scaled from 0 to 1. In addition, with statistical training based on probability theory, it is possible to apply information obtained from the analysis of real music to the recognition. Thus, the framework has both scalability and flexibility.

2.2 Issues

We use the hypothesis-search-based (HSB) method proposed by Yoshioka *et al.* to recognize chord symbols and chord boundaries simultaneously. We chose this method over the HMM-based one because it expressly solves the mutual dependency problem between chord symbols and chord boundaries. Furthermore, the HSB method makes it easier to probabilistically integrate various musical elements. That is, we are able to integrate bass pitch estimation into the chord recognition. In this paper, Yoshioka's HSB method is called the baseline method. However, two issues remain in using it to calculate the evaluation value of the hypothesis.

2.2.1 Usage of bass pitch estimation

Although information about bass sounds is used in the baseline method, it is not used in a probabilistic framework. When the predominant single tone estimated is different from the harmonic tones of the chord, penalties are imposed on the certainty based on bass sounds. Errors in estimating single tones also tend to produce errors in chord recognition.

2.2.2 Non-probabilistic certainties

The certainties based on musical elements for the evaluation function are not probabilistic in the baseline method. When the observation distribution of chroma vectors [7] is approximated with a single Gaussian, the Mahalanobis generalized distance between acoustic features is used as the certainty. Another certainty based on chord progression patterns is defined as a penalty. The criterion for applying this penalty is related to progression patterns that appear several times. However, as the scales of the values was inconsistent, it becomes difficult to integrate multiple elements. Additionally, optimizing the weighting factors of each value takes a lot of time and effort.

2.3 Our Solution

To resolve the above issues, we use bass pitch probability (BPP) and define hypothesis reliability by using a probabilistic function.

2.3.1 Bass Pitch Probability

We utilize BPP as information about bass lines to reduce the effect of bass pitch estimation errors on chord recognition. BPP can be estimated using a method called PreFEst [8]. Because BPP is uniform in non-bass-sound frames, it does not significantly affect chord recognition in these frames.

2.3.2 Probabilistic design of hypothesis reliability

It is necessary to reformulate hypothesis reliability in order to use BPP in the reliability calculation. Like the baseline

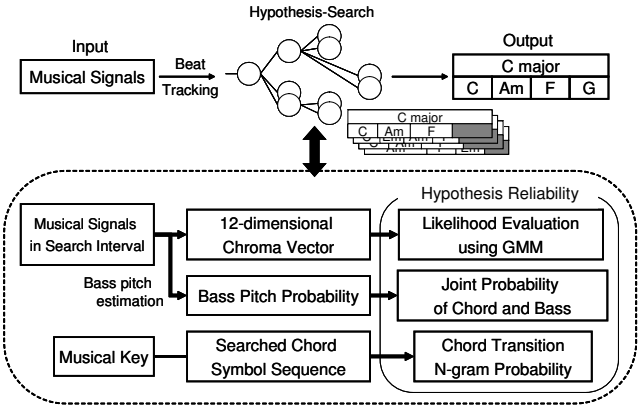

Figure 2. System Overview

method, we use acoustic features and chord progression patterns. However, we define the reliabilities based on these features by using a probabilistic framework. The values of the acoustic features are based on the likelihood of Gaussian Mixture Models (GMMs) for 12-dimensional chroma vectors. The values for the progression patterns are based on the transition probability obtained from statistical N-gram models. This reformulation enables three reliabilities to be integrated: those based on acoustic features, on BPP, and on transition probability.

3 SYSTEM

By applying information about BPP when calculating hypothesis reliability, we integrate the baseline method and bass pitch estimation in our chord recognition system. The three reliabilities based on the three elements are formulated probabilistically so that the system deals with them uniformly.

Figure 2 shows an overview of our automatic chord recognition system. First, the beat tracking method from [9] is used to detect the eighth note level beat times of an input musical piece. Second, the hypotheses are expanded over this beat time, and hypothesis reliability is calculated based on these three reliabilities. Third, a beam-search method using hypothesis reliability as the cost function is used to prune the expanded hypotheses. These operations are repeated until the end of the input signal. Finally, we obtain a tuple comprising the chord symbol sequence, chord boundary sequence, and key from the hypothesis that has the highest reliability.

3.1 Specification of Chord Recognition System

We define automatic chord recognition as a process of automatically obtaining a chord symbol sequence, a chord boundary sequence, and a key. These elements are defined as follows:

- **Chord symbol sequence**
  $\mathbf{c} = [c_1 \cdots c_M], c_i \in \mathbb{C} \equiv \mathbb{R} \times \mathbb{T}$
  The system uses 48 classes (major, minor, diminished, and sus4 for each of the 12 roots). Triads, sevenths, and so on are included as subclasses of these larger classes. we focused on discriminating the larger classes. For MIR applications we believe the larger classes would be sufficient to capture the characteristics or moods of accompaniments of musical pieces.
- **Chord boundary sequence**
  $\mathbf{t} = [t_0 \cdots t_M], t_i \in \mathbb{N}$
  where M is the number of chord symbols,
  t_i denotes the boundary time of c_i and c_{i+1},
  t_0 denotes the beginning time of the input signal,
  and t_M denotes the end time of the input signal.
- **Key**
  $k, k \in \mathbb{K} \equiv \mathbb{R} \times \mathbb{M}$

$\mathbb{R}$, $\mathbb{M}$, $\mathbb{T}$ are defined as follows:

$$\mathbb{R} \equiv \{\mathrm{C}, \mathrm{C\#}, \cdots, \mathrm{B}\}, \quad \mathbb{M} \equiv \{\mathrm{Major}, \mathrm{Minor}\},$$
$$\mathbb{T} \equiv \{\mathrm{Major}, \mathrm{Minor}, \mathrm{Diminished}, \mathrm{Sus4}\}$$

We also assume the tempo stays constant, the beat is a common measure (four-four time), and the key does not modulate.

3.2 Formulating of Hypothesis Reliability

Denoting the observed feature sequence over frames $\tau = (\tau_s, \tau_e)$ as X_τ, we can probabilistically define hypothesis reliability Rel_τ as follows:

$$Rel_\tau = p(c_\tau | X_\tau) \quad (X_\tau = [x_{\tau_s} \cdots x_{\tau_e}]) \tag{1}$$

The BPP, β_f^τ, during a duration τ is defined from the frame-by-frame BPP, $w^{(t)}(f)$, as follows:

$$\beta_f^\tau = \sum_{i=\tau_s}^{\tau_e} w^{(i)}(f)/\alpha = \{w^{(\tau_s)}(f) + \cdots + w^{(\tau_e)}(f)\}/\alpha \tag{2}$$

where f denotes the frequency of the bass pitch and α is a normalization constant. Hypothesis reliability integrating BPP on the duration τ is defined as follows:

$$Rel_\tau = p(c_\tau | X_\tau) = \sum_f p(c_\tau, \beta_f^\tau | X_\tau) \tag{3}$$

This hypothesis reliability is converted with Bayes' theorem into another form as follows:

$$\sum_f p(c_\tau, \beta_f^\tau | X_\tau) \propto \sum_f p(X_\tau | c_\tau, \beta_f^\tau) p(c_\tau, \beta_f^\tau) \tag{4}$$

$$= \sum_f p(X_\tau | c_\tau, \beta_f^\tau) p(c_\tau | \beta_f^\tau) p(\beta_f^\tau) \tag{5}$$

We use 12-dimensional chroma vectors as the observation features. Since the vectors only depend on the chord symbol, we set the following expression.

$$p(X_\tau | c_\tau, \beta_f^\tau) = p(X_\tau | c_\tau) \tag{6}$$

Thus, hypothesis reliability over τ becomes as follows:

$$Rel_\tau = p(X_\tau|c_\tau) \sum_f p(c_\tau|\beta_f^\tau)p(\beta_f^\tau) \quad (7)$$

where $p(X_\tau|c_\tau)$ denotes the reliability based on acoustic features, and $\sum_f p(c_\tau|\beta_f^\tau)p(\beta_f^\tau)$ denotes the reliability based on BPP.

The key k is independent of chord boundaries. With the conditional probability of a chord symbol sequence given a key, overall hypothesis reliability Rel_{all} is defined as follows:

$$Rel_{\text{all}} = p(\mathbf{c}|k) \prod_\tau Rel_\tau \quad (8)$$

$$= p(\mathbf{c}|k) \prod_\tau p(X_\tau|c_\tau) \sum_f p(c_\tau|\beta_f^\tau)p(\beta_f^\tau) \quad (9)$$

where $p(\mathbf{c}|k)$ denotes the reliability based on transition probability of chords.

3.3 Reliability based on Acoustic Features

We use 12-dimensional chroma vectors as acoustic features; these vectors approximately represent the intensities of the 12-semitone pitch classes. As the chord symbols are identified by the variety of tones, it is essential for chord recognition.

Because we focus on four chord types, major, minor, diminished and sus4, we use four M-mixture GMMs (Maj-GMM, Min-GMM, Dim-GMM, and Sus4-GMM). The parameters of each GMM, λ_t, are trained on chroma vectors calculated at the frame level. Note that chords of different roots are normalized by rotating chroma vectors. This normalization reduces the number of GMMs to four and effectively increases the number of training samples. The EM algorithm is used to determine the mean, covariance matrix, and weight for each Gaussian.

After chroma vectors from the input signals are rotated to adapt to the 12 chords having different root tones, we calculate the log likelihood between them and the 4 GMMs. The likelihood is equal to $p(X_\tau|c_\tau)$. Thus, the reliability divided by the number of frames in the hypothesis, g_c is defined as follows:

$$g_c = \log p(X_\tau|c_\tau) = \log p(X_r|\lambda_t), \quad (10)$$

where r denotes the number of rotating chroma vector indexes and t denotes the number of GMM types.

3.4 Reliability based on Bass Pitch Probability

BPP is obtained by bass pitch estimation, and it is used to represent the degree of each pitch of bass sounds. Since the bass lines determine the chord sequence, they should be simultaneously analyzed for recognizing the chord sequence.

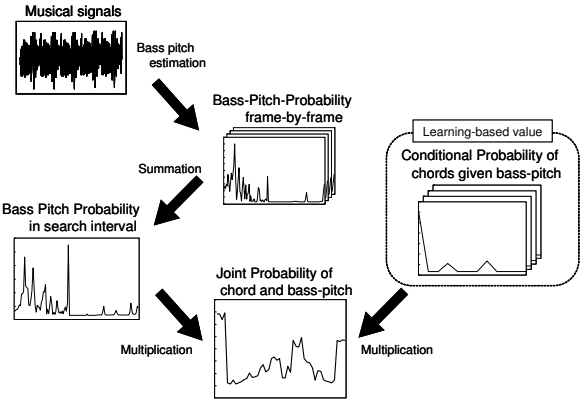

Figure 3. Bass-Pitch Processing

Figure 3 shows an overview of the process for calculating the reliability based on BPP. The prior probability of bass sounds $p(\beta_f^\tau)$ is defined by using BPP $w^{(t)}(f)$ as follows:

$$p(\beta_f^\tau) = w^{(\tau)}(f) = \sum_{j=\tau_s}^{\tau_e} w^{(j)}(f) \quad (11)$$

On the other hand, the conditional probability of chord c_i given bass pitch β_f^τ is obtained from real music by using correct chord labels and the results of PreFEst for the particular duration. We statistically calculate the frequency of appearance of each bass pitch for each chord.

As the reliability based on BPP is the log joint probability of the chord and bass pitch, the reliability g_b is defined in terms of the BPP $w^{(\tau)}(f)$ and the conditional probability $p(c_i|\beta_f^\tau)$.

$$g_b = \log(\sum_f p(c_i|\beta_f^\tau)w^{(\tau)}(f)) \quad (12)$$

3.5 Reliability based on Transition Probability

Music theory indicates that the genre and the artist usually determine the chord progression patterns for a given musical piece. The progression patterns are obtained from the key and scale degree. We probabilistically approximate the frequency of a chord symbol appearing, thus reducing the ambiguity of chord symbols.

We use two 2-gram models, one for major keys and one for minor keys. They are obtained from real music in advance. In the learning stage, we obtain the 2-gram probabilities from common chord symbol sequences consisting of the key and correct chord labels. We calculate the 2-gram probability $p(c_i|c_i^{(-1)})$ from the number of progression patterns and use smoothing to handle progression patterns not appearing in the training samples.

We estimate the transition 2-gram probability on a log scale by using the hypothesis's key. The reliability g_p is defined as follows:

$$g_p = \log p(c_i|k) = \log p(c_i|c_i^{(-1)}) \quad (13)$$

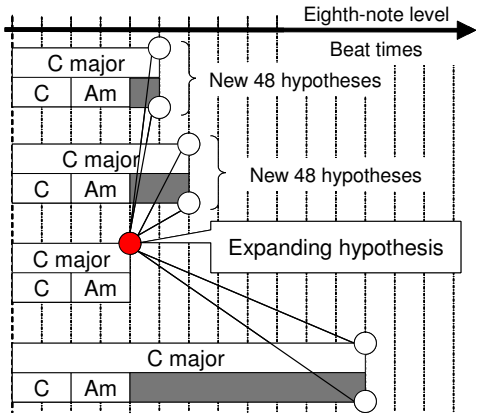

Figure 4. Hypothesis Expansion

3.6 Integrating three reliabilities

Because hypothesis reliability on a log scale is the weighted sum of the three log reliabilities described above, Rel is defined as follows:

$$Rel = w_c \times g_c + w_b \times g_b + w_p \times g_p \qquad (14)$$

where w_c, w_b, and w_p are weight constants.

3.7 Updating the Hypothesis

3.7.1 Hypothesis Expansion

Figure 4 shows the hypothesis expansion process. We consider the minimum size of boundary intervals to be eighth-note level beat times, which is the same as the baseline method. For each unit between one and eight beats, 48 hypotheses are generated for 48 chord symbols.

3.7.2 Hypothesis Pruning

We use a beam search to prune the expanded hypotheses; this prevents the number of hypotheses from expanding exponentially. In the beam search, hypotheses are pruned using a beam width BS. After the pruning, the BS hypotheses with higher reliability than the rest of the hypotheses are expanded. Furthermore, by delaying the expansion of hypotheses until it comes time to evaluate them, we can apply pruning to newly generated hypotheses. As a result, by decreasing wasteful expansions of hypotheses we can reduce both the amount of computation and memory required.

3.7.3 Update of Hypothesis Reliability

When a hypothesis is expanded, we need to update hypothesis reliability. To treat hypotheses with different numbers of intervals fairly, the reliability of a new hypothesis Rel_{new} is defined as follows:

$$Rel_{new} = \frac{Rel_{prev}}{N_h} + Rel_{next}, \qquad (15)$$

Table 1. Parameter values

$BS = 25$	$M = 16$	
$w_c = 1.0$	$w_b = 2.0$	$w_p = 0.3$

Table 2. Results of 5-fold cross validation
[1]:Baseline Method, [2]:Ac, [3]:Ac+Ba, [4]:Ac+Pt, **[5]:Our Method**

Groups	[1]	[2]	[3]	[4]	**[5]**
1st	65.1	62.9	71.7	68.3	**78.3**
2nd	62.7	62.6	70.6	65.7	**74.9**
3rd	57.7	60.4	66.8	61.2	**69.5**
4th	61.0	61.3	70.2	64.0	**72.7**
5th	61.6	60.9	69.2	64.8	**71.4**
Total	61.6	61.6	69.7	64.8	**73.4**

where Rel_{prev} denotes the reliability of the previously expanded hypothesis, Rel_{next} denotes the hypothesis reliability of the newly expanded interval, and N_h denotes the number of previously expanded intervals.

4 EXPERIMENTAL RESULTS

We tested our system on one-minute excerpts from 150 songs in 12 Beatles albums (a total of 180 songs), which have the properties described in Section 3.1. These songs were separated into 5 groups at random for 5-fold cross validation. For training the parameters of the GMMs, we used chroma vectors calculated from audio signals of 120 of these songs and audio signals of each chord played on a MIDI tone generator. These songs were also used to train the conditional probability of chords given the bass pitch (Section 3.4). We utilized 150 songs (137 in the major key, 13 in the minor key) as training data for the 2-gram models. As the correct chord labels, we used ground-truth annotations of these Beatles albums made by C. A. Harte [10]. The implementation for experiments used the parameters listed in Table 1.

To evaluate the effectiveness of our method, we compared the frame-rate accuracies of the following five methods of computing the hypothesis reliability:

1. Baseline method
2. Using only acoustic features
3. Using acoustic features and BBP
4. Using acoustic features and transition probability
5. Our method (three elements)

The results are listed in Table 2. With our system, the average accuracy for the 150 songs was 73.4%. Compared with using only acoustic features, the method using both acoustic features and bass pitch probability improved the recognition rate by 8.1 points. Furthermore, the method using acoustic features, BBP, and transition probability improved the recognition rate by 11.8 points. In addition, our system's accuracy was higher than that of the baseline method. This is because the probabilistic integration enabled us to

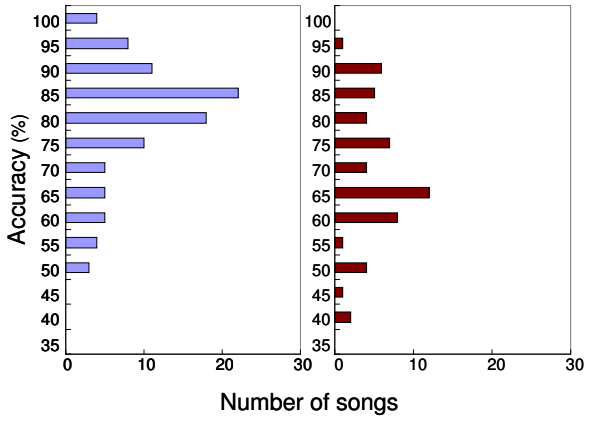

Figure 5. Accuracy Histograms.(Left) Histogram for the songs with correct key. (Right) Histogram for the songs with incorrect key.

utilize information about bass lines as a clue in chord recognition. Thus, the results prove both the importance of considering the interrelationship between chord sequence and bass lines and the effectiveness of the probabilistic integration of these elements.

We compared our results with those obtained from other systems proposed by Bello [2] and Lee [3]. They used two Beatles albums (*Please Please Me*, *Beatles for Sale*) as the test data set. Our system had an average accuracy of 77.5%. Although both Bello's and Lee's system used different training data from those that we used, their systems had 75.0% and 74.4% accuracy, respectively.

Upon investigating the accuracy distribution for all songs, we fount that the accuracy histogram for all songs is polarized with two peaks split at approximately 70%. Figure 5 shows two accuracy histograms, one for songs where the key was estimated correctly and another where it was incorrect. As these histograms describe the polarization, it is clearly important to correctly estimate the key for chord recognition. To improve key estimation, we plan to develop a method that searches the hypotheses by using not only a forward search but also backtracking.

5 CONCLUSION

We presented a chord recognition system that takes into account the interrelationship among musical elements. Specifically, we focus on bass lines and integrate hypothesis-search-based chord recognition and bass pitch estimation in a probabilistic framework. To evaluate hypotheses, our system calculates the hypothesis reliability, which is designed by the probabilistic integration of three reliabilities, based on acoustic features, bass pitch probability, and chord transition probability. The experimental results showed that our system had a 73.4% frame-rate accuracy of chord recognition in 150 songs. They also showed an increase in accuracy when the three reliabilities were integrated compared with the baseline method and a method using only acoustic features. This shows that to recognize musical elements (not only musical chords but also other elements), it is important to consider the interrelationship among musical elements and to integrate them probabilistically. To obtain more information about how to recognize chord sequences more effectively, we will design a way to integrate other musical elements such as rhythm.

6 ACKNOWLEDGEMENTS

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ON RHYTHMIC PATTERN EXTRACTION IN BOSSA NOVA MUSIC

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ABSTRACT

The analysis of expressive performance, an important research topic in Computer Music, is almost exclusively devoted to the study of Western Classical piano music. Instruments like the acoustic guitar and styles like Bossa Nova and Samba have been little studied, despite their harmonic and rhythmic richness. This paper describes some experimental results obtained with the extraction of rhythmic patterns from the guitar accompaniment of Bossa Nova songs. The songs, played by two different performers and recorded with the help of a MIDI guitar, were represented as strings and processed by FlExPat, a string matching algorithm. The results obtained were then compared to a previously acquired catalogue of "good" patterns.

1 INTRODUCTION

It is common sense that playing music in the exact way it is written in the score results in a mechanical and uninteresting succession of sounds. To make written music interesting, the musician is required to make variations on low level musical parameters, such as: Local tempo (*accelerandi*, *ritardandi*, *rubato*); dynamics; notes articulation (*staccati*, ligatures, etc.); micro-silences between the notes, etc. [12].

Several researchers stress the importance, as well as the the difficulties, of studying this phenomenon, also known as *expressive performance* [14]. These researches, in general, focus on building relationships between different musical elements (such as harmony and melody or even the low level parameters previously mentioned), and expressive performance itself. These relationships can be described in many ways, from different points of view or *levels of abstraction*, and including various musical parameters. Examples of such relationships are rules like *"lengthen a note if it is followed by a longer note and if it is in a metrically weak position"* or *"stress a note by playing it louder if it is preceded by an upward melodic leap larger that a perfect fourth"* [13].

With some exceptions [3, 6], the role of rhythm in expressive performance has not been thoroughly studied so far, despite its intuitive importance. Moreover, the research is almost exclusively devoted to the Western Classical Music composed for the piano. We are interested in studying the *Música Popular Brasileira* (Brazilian Popular Music)—MPB, represented by artists like João Gilberto, Tom Jobim, Caetano Veloso, Gilberto Gil, etc. We are particularly interested in the guitar accompaniment, that is, in how the guitar player accompanies the singer or solo instrument.

This paper presents an experiment that focus on the discovery of rhythmic patterns in Bossa Nova music. For this, two different performers played several songs on a MIDI guitar, which were processed in the form of strings. These strings were then processed using FlExPat, a pattern matching algorithm [8], and the results were then compared to a catalogue of patterns that reflects the rhythmic patterns used by João Gilberto, the "inventor" of the Bossa Nova style.

The remainder of the paper is organized as follows: In Section 2, we discuss what motivated us to try such an experiment. In Section 3, we describe how the data was acquired and the representation we used. In Section 4, we present the experiment itself and discuss the results we obtained. Finally, in Section 5, we present some conclusions and future directions for this work.

2 MOTIVATION

Apart from obvious differences (instrument, genre/style and player's role), there is a much more fundamental difference between research focusing on Western Classical Music and research that deals with MPB: While in the Western Classical Music case there is some sort of "official" notated version of musical pieces (the score, namely), in MPB there is no such thing. What may be available is the chord grid (chord sequence that should be played), and, in some rare cases, the score of the melody. Even when the chord grid is available, the musician is usually allowed to change the harmony and play something different from what was initially notated. Considering that, it becomes clear that the guitar player has a major role in the accompaniment's construction.

This importance becomes even clearer when the rhythm is taken into consideration, because, although the musician may have the chord sequence notated (*i.e.*, information about the harmony may be somehow specified), there is no indication whatsoever of the rhythm the musician should

play. It is entirely up to him/her to decide about the rhythm.

Some studies pointed out, however, that the guitar accompaniment in styles like *Bossa Nova* and *Samba* is built by the concatenation of certain rhythmical groups or *patterns* [4, 9]. There are, however, several aspects of the accompaniment construction that are only known by practitioners of these styles. Moreover, the knowledge about the accompaniment construction is mainly subjective. Due to this lack of formalized knowledge, there are many open questions such as:

- Are there rhythmic patterns that are preferred by a certain performer or required for a certain musical style? In which situations and in which frequency do they show up?
- Are there variations of these patterns? Is it possible to group these variations in meaningful way? Which variations (timing, dynamics, etc.) are acceptable within a pattern?
- Is it really the case that everything is a pattern, i.e., are there parts that are not recurrent?
- Is it possible to justify the choice of a pattern in terms of other musical features (melody, harmony, tempo, musical structure, style, etc.)?
- Is it possible to build a dictionary of patterns for a given player? Does this dictionary changes when the style changes (Bossa Nova and Samba, for instance)? Do different players have different dictionaries?
- Is it possible to build a grammar or a set of rules that is able to describe formally how the patterns are chained and/or the rhythmical transformations done by a given performer? If so, what are the relations between grammars from performers p_1 and p_2?

More general questions could also be posed: To which extent patterns used in Bossa Nova music are different from patterns used in Samba? How different is Samba today (in terms of patterns and pattern usage) from Samba in the 1920's or 1930's? Is Bossa Nova today still played as it was in the 1950's, when it was created? How can we describe those differences, if any?

3 DATA ACQUISITION AND REPRESENTATION

For the experiment, two different players, referred to from now on as $G1$ and $G2$, were invited to record the accompaniment of some Bossa Nova songs on a MIDI guitar [1]. $G1$ performed the following songs: *Bim Bom*, *O Barquinho*, *Insensatez* (How Insensitive), *Garota de Ipanema* (Girl from Ipanema), *Só Dançco Samba*, and *Wave*. From $G2$, we recorded *A Felicidade*, *Chega de Saudade*, *Corcovado*, *Desafinado*, *Eu Sei Que Vou Te Amar*, *Samba de uma Nota Só*, *Garota de Ipanema*, *Só Dançco Samba*, *Insensatez*, *Tarde em Itapoã*, and *Wave*. In the total, we collected 16 recordings (ca. 30 minutes of music). It was requested for the performers to play the songs according to a provided notation (the chord grid as notated by Chediak [1]).

The acquired data was, however, not ready for usage. Probably due to technological restrictions, the resulting MIDI files were noisy and it was necessary to clean the collected songs before using them. Noisy files contained notes that were not played by the guitarist and these notes could be grouped into two basic types: Very short notes (usually high pitched) and notes with very low volume (loudness). There was yet a second type of problem, namely events that were somehow misplaced by the recording equipment (usually a semitone up or down the actually played note). The first type of noise was removed automatically, but the second one required manual correction, which was done with the help of the recording's audio files [2]. After this step, the data was beat tracked at the eighth note level using *BeatRoot* [3], an interactive system that outputs the MIDI file beat tracked.

As we are interested in the discovery of rhythmic patterns, an essential information is the moment when a note is played or its *onset*. It would be very hard, however, to use this information only in a meaningful way. We should, then, associate some other information with onsets in order to better represent the songs. Pitches may come to the reader's mind as the most relevant information that could be used with onsets, but they are not intrinsically linked to the rhythm. If we, however, observe how sound is produced by the guitar player, we may link each onset to the finger that generated it. So, the finger used to pluck the string and produce sound may prove a much more interesting *abstraction* than, for instance, the pitches.

But, this abstraction was not readily available in the files we collected [3]. So, we had to develop an algorithm to automatically determine the right hand *fingering* [11]. Roughly speaking, we first introduced the concept of *hand position*, that is, the fingers' position regarding the string or strings they are about to pluck. Then, we created a *hand position set* that contains all relevant hand positions. Considering that the fingering is formed by transitions between consecutive hand positions, we assigned a cost to each transition [4].

[1] The equipments we used in the recordings were the following ones: An acoustical guitar with an RMC Poly-Drive II pick-up installed (`http://www.rmcpickup.com/polydriveii.html`) that was connected to a Roland GR-33 guitar synthesizer responsible for the pitch to MIDI conversion (`http://www.roland.com/products/en/GR-33/index.html`).

[2] We recorded, at the same time, MIDI and audio information.

[3] Note that the MIDI files we had at hand contained no information at all about the fingering. But, we collected them in a way that each string was recorded separately on its own MIDI channel.

[4] This cost represents, in fact, the amount of effort required to change from hand position HP_i to hand position HP_{i+1}.

Each possible hand position in the hand position set can now be represented as a node in a graph, whereas the costs as edges weights. The fingering problem can then be reduced to the discovery a path that minimizes the overall cost, and our algorithm simply tries to find an optimum path in this graph [5].

Our algorithm outputs the songs as depicted in Figure 1. Here, letters T, F, M and R represent, respectively, the thumb, fore, middle and ring fingers, crosses (+) represent the beats, and pipes (|) represent the measure bars. Each beat was equally divided by four, so each letter, cross or minus (–) represents the duration of a 32nd. Except for the last line, that represents exclusively the beats, each of the remaining lines represents one guitar string, ordered from higher to lower (i.e., first line represents the high E string, second line the B string, and so on until the low E string).

```
|-----------------|--------
|R---R-----R-----|R---R---
|M---M-----M-----|M---M---
|F---F-----F-----|F---F---  [...]
|-----------------|--------
|T-------T-------|T-------
|+---+---+---+---|+---+---
```

Figure 1. Right hand fingering for song *Insensatez*, played by $G2$

This representation, however, can be viewed as a polyphonic one (each guitar string being one "voice"). Polyphonic pattern matching, however, is a very difficult task [5] and we would like to avoid these difficulties, at least at our initial steps and experiments. So, we further reduced this initial representation to a one-dimensional string with minimum loss of information [6]. This simplified string is formed by the alphabet $\Sigma = \{b, B, p, P, l, a, A, s, S, -, +, |\}$. The meaning of each symbol is the following:

- Uppercase letters stand for events that occur on-beat, while lowercase letters for off-beat events;
- Letter b stands for "bass", i.e., events played with the thumb only, and letter p stands for "chord" (*sic*), i.e., events that are played with some combinations of two or more of fingers F, M and R [7];
- Letter l also stands for "chord", but a chord whose duration goes beyond the measure it was played (*i.e.*, we make a difference between a chord that is completely within a single measure and a chord that starts in one measure and ends in the next one);
- Letter a stands for "all", i.e., b and p played together, and letter s stands for "single note", i.e., events that are played with only one of fingers F, M and R; and
- Symbols $+$, $-$, and $|$ have the same meaning stated before.

It is important to note that this kind of reduction is also done by musicians themselves: They usually describe the rhythmic patterns as sequences of "baixos" or *basses* (events played with the thumb only) and "puxadas" or *chords* (events played with some combinations of two or more fingers). Figure 2 depicts part of the fingering for song Insensatez. Above the thick black line is the fingering as outputted by the fingering algorithm. Under it is the resulting simplified string.

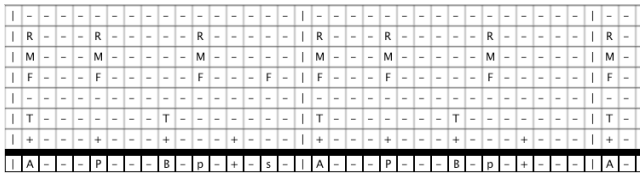

Figure 2. Fingering and one-dimensional string for song *Insensatez*, played by $G1$

4 EXPERIMENT

Considering that patterns are the building blocks used to create the accompaniment, finding rhythmic patterns in Bossa Nova songs is a basic step towards understanding how the accompaniment is built by musicians.

We have, however, one initial problem: In order to find patterns automatically, we need to use an existing algorithm or, case it fails to find patterns, we need to develop one that is able to find them. The big issue behind this problem is the following: How can we assess the results? How can we say that one algorithm performs better or worse than another one? How can we say that an algorithm is unsuitable for finding rhythmic patterns?

There is one particularity in the Bossa Nova domain: João Gilberto, the "inventor" of the style, is considered a model, and, as such, the musicians try to imitate him, playing the patterns he plays [8]. So, it is perfectly reasonable to assume that an algorithm that finds, in any Bossa Nova data set, the patterns used by João Gilberto has a minimum acceptable performance level.

[5] Note that our algorithm follows a similar approach used by Sayeg, as described in [10].

[6] Actually, with the alphabet we used we just can not recover the string where the note was played, what was not relevant for the experiments we made. We could easily avoid this information loss introducing new symbols in the alphabet.

[7] The terms "baixo" and "puxada" may explain more clearly the origin of letters b and p!

[8] A musicologist could ask, then: If someone does not play like the inventor is he/she playing Bossa Nova at all?

But, what are the patterns João Gilberto plays? In the literature we were able to find several transcriptions of patterns played by João Gilberto [4, 9]. So, we have built a catalogue containing 21 different patterns (labeled P1, P2, etc.), all manually transcribed by musicologists from João Gilberto's recordings. Examples of patterns in this catalogue are showed in Figure 3.

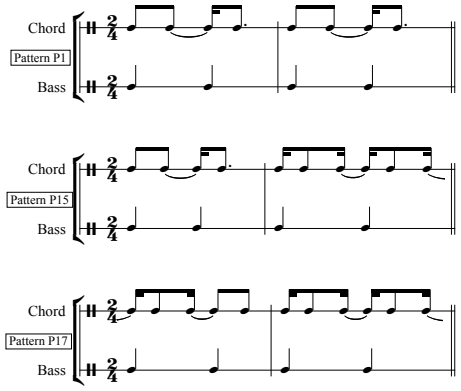

Figure 3. Examples of patterns in the catalogue

The experiment was, then, to explore the data set we had acquired, verifying if patterns from this catalogue could be found in the data set. For the experiment the algorithm FlExPat [8] was chosen. It is an inexact string matching algorithm that was inspired by algorithms from the Computational Biology field, but that also incorporates results from previous research on musical similarity [7]. Given an input (here the simplified string previously described) and using the edit distance as its similarity measure [9], the algorithm outputs a collection of patterns, organized in classes. Each class has a prototype, that is the most representative pattern of the class, and several occurrences, possibly inexact, of this prototype. It is also possible for the user to provide the algorithm with some constraints, such as the maximal and minimal length of patterns, the similarity threshold, the maximum difference between two candidates to be compared, etc.

In our first experiment FlExPat was used "as is", that is, we used the algorithm without any sort of modifications and/or adaptations. Although the algorithm found patterns from the catalogue, including patterns with some small modifications, the results, in this case, were not so good as we previously expected. Table 1 summarizes the results we obtained with FlExPat.

The algorithm's configuration we used was the following: $m_{min} = 17$ (minimal pattern length), $m_{max} = 34$ (maximal pattern length), and similarity threshold of 0.75 (normalized values). The lengths m_{min} and m_{max} correspond to patterns varying from one to two measures [10]. On

Pat	$G1$	$G2$	**Pat**	$G1$	$G2$
P1	×	×	P12		
P2	×		P13		
P3			P14		
P4	×		P15	×	
P5			P16		
P6			P17		
P7			P18		
P8			P19		
P9	×	×	P20		
P10	×	×	P21		
P11			/	/	/

Table 1. Patterns from the catalogue found in data set

average, each song has 48 classes of patterns (smallest number of classes was 40 and greatest number was 78).

In song *Garota de Ipanema*, played by $G2$, the algorithm was able to identify the following pattern: `|A---P---B-p-+---|P---P---Bp--+l`. The corresponding pattern in the catalogue is `|A---P---B-p-+---|A---P---B-p-+-l-`. Looking closer at the modifications done by the performer, it is possible to notice that he usually substitutes `A` by `P` in the second measure of the pattern, and that he anticipates both final chords in this same measure. Figure 4 depicts these variations.

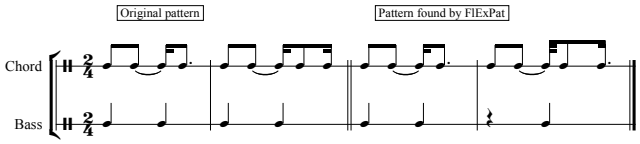

Figure 4. Pattern found by FlExPat

In song *Wave*, as played by $G1$, FlExPat identified as the prototype of class 18 the following pattern: `|A---P---B-p-+l--|B---P---Bp--+p-`. The corresponding pattern in the catalogue is `|A---P---B-p-+-l-|B---P---B-p-+-l-`. As in the previous case, the modifications done by the performer are mainly anticipations.

The main problem we found was that the extracted patterns, many times, start from the middle of a measure, such as in `--B---B---+---|P---B---B---+---|P` (FlExPat also found the expected pattern, `|P---B---B---+---|P---B---B---+--`). This brings some problems to the evaluation of the results: Due to the great number of patterns with this kind of structural malformation, it becomes difficult to validate the patterns.

[9] Dynamic programming is used to compute it efficiently.

[10] It does not mean, however, that the patterns necessarily start at the beginning of the measure. There is no way to specify such a constraint in FlExPat.

This problem is even bigger when most of a class is formed by such patterns (it happened many times, unfortunately).

From these results we may rush and conclude that FlExPat performed poorly, since it found just a few patterns from the catalogue in the data set and because these patterns were grouped into too many classes. However, as pointed out in the previous paragraph, there was a sort of structural malformation in many patterns found by FlExPat. So, what would happen if we could "help" the algorithm, somehow providing it with some structural information about the patterns?

Unfortunately, it is not possible to describe for the algorithm the desired aspect of the pattern: We can not say that it should only consider patterns that begin on the measure bar, for instance. We can only describe the minimum and maximum pattern lengths, as well as the minimum similarity threshold. If we, however, look more carefully at the patterns in the catalogue, it is possible to identify some common structure: They are all two measures long and they start either at the measure bar or at a "puxada" right before the measure bar [11] .

We then introduced a step just before the actual pattern matching where all songs were segmented according to this structure. It is important to note that with this modification, we changed the abstraction level at which the pattern extraction happens: Instead of comparing event by event (basses or chords), now we compare groups of events (i.e., the segments).

After we adjusted our implementation of the algorithm to accept the segments described before as input, we ran the experiment again. In this case, differently from the previous one, the algorithm found systematically in every song patterns from the catalogue. This time, more than half of the catalogue's patterns was found, which significantly improves the results we obtained before. On average, each song had 14 classes of patterns (smallest number was 6 and greatest number was 34). Table 2 shows the patterns found in this second part of the experiment.

After running the experiments and having in mind that we have used a small data set, it is now possible to discuss a little bit the results we obtained. First of all, let us examine the patterns used by the performers. As described by Dahia et al. [2], patterns in the catalogue belong to one of four groups: cyclical ($P2$ to $P8$), beginning ($P9$ to $P12$), special ($P13$ and $P14$) and fill-in patterns ($P15$ to $P21$). Pattern $P1$ is the main Bossa Nova pattern and forms its own group. From the results presented before (Tables 1 and 2), it is possible to see that the players used patterns belonging to all groups, but the special one. This group, however, contains patterns that are rarely used and it would not be a problem if they do not show up. Given that the catalogue contains Bossa Nova patterns, we can say, then, that the performers do follow the Bossa Nova style in the rhythmic accompaniment construction.

[11] It means that all segments look like `|<pattern>` or like `l*|<pattern>`, where the * symbol represents a sequence of zero or more minuses only.

Pat	$G1$	$G2$	**Pat**	$G1$	$G2$
P1	×	×	P12		
P2	×	×	P13		
P3	×	×	P14		
P4	×		P15		×
P5	×		P16	×	×
P6	×		P17		
P7			P18		×
P8			P19	×	
P9	×	×	P20		
P10	×	×	P21		
P11			/	/	/

Table 2. Patterns found by the modified version of FlExPat

There are occasions, however, that the pattern used does not belong to the catalogue. $G1$, for instance, uses the pattern `|P---B---B---+---|P---B---B---+---` in song *Eu Sei Que Vou Te Amar*. We can interpret this fact in many ways, such as: the player deliberately used a non Bossa Nova pattern, the catalogue may not be complete, or even that Bossa Nova style has changed over the time and players started to use or create their own patterns.

5 FINAL REMARKS AND FUTURE WORK

This paper described an experiment that dealt with the extraction of rhythmic patterns from Bossa Nova songs. Sixteen beat tracked MIDI files, representing the recording of several songs by two different players, were represented as a string and thereafter processed by a string matching algorithm called FlExPat. The objective was to identify in the data set patterns from a previously acquired catalogue of patterns.

First results showed that, although FlExPat could find some patterns, many patterns, including some that could be clearly heard, were not found. After examining the structure of the catalogue's patterns, we were able to pre-segment the songs and use the pre-segmented version of the song as input for the algorithm. This time the results were much better, and more than half of the patterns was found by the algorithm.

It is important to stress the relevance of the abstractions we have built. First of all, we could reduce the complexity of the pattern matching process by rewriting a polyphonic song as a monophonic line, with minimum information loss. This first abstraction was, however, not sufficient, as can be seen with the first experiment we made, and we had to build a second abstraction, going up one structural level: From single events to groups of events (*segments*). Only due to these

abstractions we could explore the data set we collected.

Of course, there are several points for improvement. One that is particularly important is the representation. We used a very simple representation of the events. Attributes like tempo, structural or harmonic information are not represented. The more attentive reader may have even noticed that the actual duration of the events is not represented at all. We just used the onset information, which turned out to work for this experiment. To further investigate the particularities of the patterns, however, we surely need to represent appropriately the duration of each event. Another point related to the representation is the following: what kind of results would we have if we had represented the downbeat explicitly? Would they be better than the ones we presented here?

The evaluation of the algorithm's results was done in an *ad hoc* manner: Results were compared, one by one, to the patterns in the catalogue. This procedure takes too much time, is error prone, and, therefore, must be improved. We plan to implement a tool to help us with this task.

The data acquisition is an important and non-trivial problem. It is important because if we want to draw relevant and significant conclusions about Bossa Nova (and any musical style, in fact), we must have a much more representative data set. And it is non-trivial because there are many factors involved varying from the availability and willingness of certain performer to record for us, to copyright issues of the collected material. One possible way to remedy this problem is to use audio recordings as start point, but, unfortunately, there are other problems (separating guitar and singing voice signals most notably).

FlExPat's problem (structural malformation) should be examined more carefully, since, instead of a problem, it can mean another thing. Several patterns of the catalogue have a common substructure (i.e., sub-parts of these patterns are equal). It may be the case that the second measure of patterns whose second measure are equal, are frequently concatenated with patterns whose first measure are similar. So, it may be the case that these concatenations are so typical that they are "promoted" to patterns by the algorithm.

Finally, we hope that the questions previously formulated were provocative enough. We do believe that answers to them will bring much more understanding of how the musicians bring their genres to life, as well as register how they, musicians and styles, evolved through the years.

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LEARNING MUSICAL INSTRUMENTS FROM MIXTURES OF AUDIO WITH WEAK LABELS

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ABSTRACT

We are interested in developing a system that learns to recognize individual sound sources in an auditory scene where multiple sources may be occurring simultaneously. We focus here on sound source recognition in music audio mixtures. Many researchers have made progress by using isolated training examples or very strongly labeled training data. We consider an alternative approach: the learner is presented with a variety of weaky-labeled mixtures. Positive examples include the target instrument at some point in a mixture of sounds, and negative examples are mixtures that do not contain the target. We show that it not only *possible* to learn from weakly-labeled mixtures of instruments, but that it works significantly *better* (78% correct labeling compared to 55%) than learning from isolated examples when the task is identification of an instrument in novel mixtures.

1 INTRODUCTION

We are interested in developing a system that can learn to recognize individual sound objects in an auditory scene where multiple sound sources may be occurring simultaneously. A system able to identify what particular sound sources are present in a mixture would enable automatic tagging of audio recordings with meta-data. A system capable of modeling sources in audio mixtures, without having to first learn the sounds in isolation, would be useful to researchers working on separating audio mixtures into their component sources.

In a typical supervised learning paradigm, isolated instances of a particular class of data are presented to a learner. Learning in this case is mostly limited to a stage of the design process. With a more ecologically realistic set of requirements, where learning can occur in an environemnt after deployment, this isolation of training instances can not be guaranteed. While similar problems have already been studied in the field of computer vision [1], auditory data presents its own unique problems because mixtures of sound can result in a composite of multiple sources at a given time.

In this paper, we focus on identification of musical instruments in a musical mixture. Almost all systems that learn to identify musical instruments require isolated examples of the target instrument at some stage in the training process (see [9] for a review). This could mean, isolated notes [12], or more recently, solo phrases from an instrument [11]. A number of these systems have been evaluated using polyphonic audio [5, 11, 15], i.e. audio with multiple simultaneous notes, but this work is still limited by the requirement that instruments be *learned* in isolation.

We are aware of two systems that *learn* solely from polyphonic audio. However, they differ significantly from the task we consider here. In [6] the system is trained on polyphonic audio, but each unique combination of instruments must be learned individually. The authors admit this approach is only realistic when learning is done before deployment. Given n instruments, this results in 2^n possible combinations that must be individually learned, significantly increasing the number of required training examples. This approach also leaves open the question of how to recognize individual instruments when presented in novel audio contexts. In addition, the training data requires labels indicating all instruments currently playing for every two second segment of audio, a fairly intensive labeling task. This is true even if a score is available (since it must be correctly aligned with a particular audio recording).

Kitahara et al [10] learn from polyphonic data, avoiding the combinatorial problems of [6]. However, they require the user to input a musical score for the recording that exactly labels every note with pitch, onset time and offset time, i.e. perfect segmentation of each instrument into individual notes is assumed.

What we would really like is a system that can learn from weakly labeled mixtures. We call a label weak if only the presence or absence of the target sound object is indicated for some N second length of audio. A weakly labeled positive example will contain the target object at some point in the example, but a significant portion of the time may not contain audio from the target. A weakly labeled negative example does not contain the target sound object at any point in the example. In this scenario a useful positive training example for the "saxophone" class would be

an acoustic recording of a saxophone and bass duo playing at a gallery opening. A negative example would be some other audio clip that contains no saxophone in the recording. Training could be accomplished on-line by recording a few audio clips that contain the desired sound. Our proposed input and output have important differences from that of previous systems that learn from polyphonic audio [6, 10] (weakly vs. strongly labeled input and multi-class vs. a presence/absence label as output).

As a first step towards our objective, in this paper we evaluate a system that provides presence/absence labels for a single instrument over short time intervals (~2 seconds). We do not require labeling all instruments in the mixtures at short intervals, nor do we require detailed scores. Instead, the system learns from weakly labeled examples that include distractor sounds.

Our instrument classes are learned independently of the background sounds: the classifier can be trained in one background environment and used in another. Although all the tools we use to accomplish this task already exist (as descibed in Section 2), we are not aware of anyone who has yet put these pieces together the way we do. In our experiments (in Section 4), we show that it is not only *possible* to learn from a mixture of instruments (constructed as in Section 3), but that it works significantly *better* than learning from isolated examples when identifying instruments in novel mixtures.

2 CLASSIFICATION METHOD

We use the segment-and-combine [7] framework for our classification system, defined below.

1. **Segment** each example X_i by selecting n potentially overlapping segments $\{\mathbf{x}_{i1}, \mathbf{x}_{i2}, \ldots, \mathbf{x}_{in}\} \in X_i$ at random (as per [7]), using the piece extraction operator described below.

2. **Learn** a classifier function $f(\mathbf{x}) \in [0, 1]$ which returns higher values for segments more likely to be part of the target object.

3. **Classify** segments with the learned function $f(\mathbf{x})$.

4. **Combine** each of the classifications of $f(\mathbf{x})$, $F(X_i) = C[f(\mathbf{x}_{i1}), f(\mathbf{x}_{i2}), \cdots, f(\mathbf{x}_{in})] \in \{0, 1\}$. The combinator $C[\cdot]$ returns a positive result if the average of $f(\mathbf{x})$ for all segments is greater than 0.5 (as per [7]).

We chose this approach because it has shown promise, even compared to more sophisticated approaches, in similar problems, such as learning to recognize visual objects from cluttered data [7]. We describe the extraction operator in Section 2.1 and the segment classifiers $f(x)$ in Section 2.2.

Feature	Description/Details
MFCCs	Mel frequency cepatral coefficients
Brightnes	Ratio of energy above 1500Hz to energy below.
Spectral Stats	First four statistical moments of spectrum
Flatness	Ratio of geometric to arithmetic mean of spectrum
Pitch	Calculated using first peak of autocorrelation function.
Amplitude	Root mean squared average.
Flux	Difference of spectrum between subsequent frames.
Inharmoncity	The strength of the pitch: amount of energy far from $f_0 \cdot n$

Table 1. The features used to represent each audio segment.

2.1 Extraction Operator

Our extraction operator to select possibly overlapping audio segments in Step 1 selects a sub-portion of the audio, based on two parameters: time(t) and length(l). The range of possible values for our start time (t) is from 0 to the length of the file in frames. The length of the extracted segment (l) ranges from 0.1 seconds to 0.5 seconds. Given an audio example X_i, we select 5 segments per second of audio in the example. Thus, for a 10 second example there would be 50 segments. We randomly select these segments, picking values for t and l using a uniform probability distribution over time.

The *j*th segment drawn from example *i* is represented as a vector of features $\mathbf{x}_{ij} \in \mathbb{R}^d$.In this case $d = 22$ (13 MFCCs, 4 spectral statistics, plus the other 6 features). We choose to use a set of features found to be commonly useful in past work on musical instrument identification. All features used are listed in Table 1.

2.2 Segment Classifiers

We compare three classifiers: Extra Trees [8], which, as we will explain, should be fairly appropriate for this task, to a very simple classifier, the K-nearest neighbors algorithm [4], and a fairly sophisticated classifier, the Support Vector Machine [2] with a radial basis kernel. This comparison is detailed in Section 4.

It is important to recognize the noisy nature of the labeling data: it is possible that a segment from a positive example, which would be labeled positively, contains no energey from the target instrument. Since labels are weak, a positive example for "saxophone" could be a jazz recording containing many portions where the saxophone is silent.

The results in [3] suggest that methods such as bagging or randomization work well under such noisy labeling con-

ditions. The Extra Tree algorithm [8] is an ensemble method that uses randomized decision trees. It differes from other randomized decision trees, in that, with the lowest parameter setting, the algorithm would completely randomize both the split and the attribute selected at each decision node, yielding a tree completely independent of training data (Extra Tree is short for extremley randomized tree). The construction of a single Extra Tree is summarized below.

1. Let the subset of training examples be S, the feature-split pool size be K and the minimum subset be n_m.
2. If $|S| < n_m$ or if all features values are constant, or all examples in S share the same label.
   (a) Define this tree's output $f(\mathbf{x})$ as the value of the most frequent label in S.
3. Otherwise:
   (a) Select K randomly uniform split s_i and feature f_i pairs.
   (b) Select the best split-feature pair s_{max}, f_{max} by entropy measures.
   (c) Let the output $f(\mathbf{x})$ be the output of the right tree if $f_{max} < s_{max}$ for $\mathbf{x}$ and the output of the left tree if $f_{max} \geq s_{max}$.
   (d) Create the right tree, with S equal to the subset of examples with $f_{max} < s_{max}$.
   (e) Create the left tree, with S equal to the subset of examples with $f_{max} \geq s_{max}$.

Results in [8] suggest that with appropriate parameter settings, as detailed in Section 4.1.1, the Extra Tree ensemble outperforms other random tree ensemble methods.

3 DATA

For our initial tests we wanted to tightly control the difficulty of our examples, allowing us to observe how each of our classifiers degrades as the data becomes more challenging. So, rather than use commercial recordings, we trained and tested on synthesized mixtures of randomly selected notes from four instruments. Each positive example contains a mixture of instruments, one of which corresponds to the label the system should learn. Thus, a positive example for "Cello" might contain a Cello, Violin and Oboe. The four instrument classes used for our source sounds were Cello, Violin, Oboe, and Bb Clarinet. Instrument recordings were taken from a set of instrument samples made available by the University of Iowa [1]. The ranges of pitches used in our experiments for the instruments were as follows: for Cello we used notes from C2 to C7, for Violin from G3 to Db7, for Clarinet from D3 to B6, and for Oboe from Bb3 to C6. These were played at three different dynamic levels, *pp, mf* and *ff*. All samples were encoded as linear PCM 16bit wav files at a sampling rate of 44100Hz. The Iowa recordings were segmented into individual notes. Training and testing examples were constructed as follows.

1. Select a set of instruments
2. For each instrument, create a melody by randomly picking a series of note recordings from the database
3. For each instrument melody, intersperse the notes with silences of random length from 0 to some maximum value.
4. Adjust instrument volumes to obfuscate the target instrument to the desired level.
5. Sum the melodies to create a mixture.

Examples were a minimum of 10 seconds long. The average length of an example was 11.5 seconds and the longest example was 17.3 seconds. To vary the a level of target obfuscation we varied the maximum spacing between notes in the target instrument melody (s_t) and notes in the distractor instrument melodies(s_d), as well as the relative amplitude of the distractors notes (a_d) across three target obfuscation conditions. These values were selected so that we would obtain a range of target to mixture ratios, as defined in Section 4.2. For **easy obfuscation**, $s_t = 0.5$, $s_d = 1$ and $a_d = 1$; for **medium obfuscation**, $s_t = 1.5$, $s_d = 1$ and $a_d = 1$; and for **hard obfuscation** $s_t = 1.5$, $s_d = 0.5$, and $a_d = 1.5$. Thus, for easy obfuscation, target instrument notes were more frequent than distractors and of equal amplitude. For hard obfuscation, distractors were significantly more frequent and louder than targets.

Conditions varied along three dimensions: target instrument (one of the four instruments), ensemble of distractor instruments (one to two of the other instruments in all possible combinations), and target obfuscation (from easy to hard): this resulted in a total of $4 \times 6 \times 3 = 72$ data conditions. We created 10 positive examples and 10 negative examples for each of the conditions, yielding a total of 1440 example audio files.

Some examples of typical correctly and incorrectly classified examples for the mixed condition (from Section 4.1) can be found at the following URL:
`http://www.cs.northwestern.edu/~dfl909/ismir2008sounds`

4 EVALUATION

In our evaluation we sought to answer to the following questions.

[1] `http://theremin.music.uiowa.edu/index.html`

1. What effect does training on mixtures, rather than isolated notes have on accuracy when identifying instruments in novel mixtures of sound?

2. How does accuracy degrade across our three segment classifiers as the target class is increasingly obfuscated by the distractors in the mixture?

3. How does performance using weak labels compare to performance using stronger labels?

We begin our evaluation by addressing questions 1 and 2 in Section 4.1. Questions 3 is addressed in Section 4.2.

4.1 Experiment 1

In this experiment, we wanted to compare performance across isolated and mixture training conditions, across our three different segment classifiers (ExtraTrees, K-nearest neighbor and SVM).

We constructed the various ensembles of instruments as described in Section 3. We then ran an experiment under two conditions, *mixed* and *isolated* training , using the segment-and-combine algorithm descibed in Section 2 with $f(\mathbf{x})$ learned using one of the three segment classifiers (Section 2.2).

4.1.1 Parameter Settings

To select parameters for our K-nearest-neighbor (KNN) and the radial basis kernel SVM classifiers we evaluated a range of parameters using the data for our most challenging target obfuscation condition (the hard obfuscation condition as described in Section 3). We evaluated the KNN classifier with K = 1, 2, 5, 10, 20, or 50, of which 5 was the best K we found. For the parameters of the SVM, γ (variance of the Gaussian in the radial basis kernel) and C (the error penality), we considered γ= 0.001, 0.01, 0.1, 1, 10, 100, or 1000, and C=0.001,0.01,1,10,100, or 1000 (yielding $6 \times 6 = 36$ possible parameters settings for the SVM), we found $\gamma = 0.1$ and $C = 100$ was the best setting.

For our ensemble of ExtraTree's we used the recomended parameters, as described in [8]: thus $K = \sqrt{N}$ where N is the number of features (22), and $n_{min} = 5$. Generally the more trees in the ensemble the better this algorithm, so we chose a large number that did not unduely compromise speed (100).

We extracted segments from the audio files at a rate of 5 segments per second. This value was chosen based on a run of the Extra Tree algorithm using the hard obfuscation condition.

4.1.2 Procedure

In the first (*mixed*) training condition, we used the following procedure for each target instrument class T and each target obfuscation condition.

1. Train the learner on the example from 5 of the 6 possible ensembles of distractors using target T, totaling 50 positive and 50 negative examples.

2. Test on the remaining 20 examples from the final ensemble condition.

3. Repeat steps 1-2, excluding a different condition each time.

In the second (*isolated*) training condition, we used this procedure for each target instrument class T and each target obfuscation condition:

1. Train the learner on all isolated notes of an instrument. For negative examples, we use an equal number of randomly selected notes from the remaining instruments.

2. Test on the 10 negative and 10 positive mixture examples with target T, for each of the 6 ensemble conditions, for a total of 120 examples tested.

This condition included from 138-170 unique isolated notes (depending on the instrument) during training: this is as many or more notes than used in the *polyphonic* training condition.

Thus for each classifier we had a total of $24 \times 3 = 72$ trials of our system (24 mixture conditions $\times$ 3 target obfuscation conditions).

4.1.3 Results

We calculated system accuracy in each trial based on how well the system could identify approximately every two seconds of audio: each example being tested was divided into sixths: for every sixth (or about two seconds) of an example given a correct label we noted that as a success. Successes were then tallied across all examples in a trial. Note that for all obfuscation conditions there was no more than 1.5 second silence between target notes.

Since this is a balanced binary labeling task, random chance performance for a given condition is 50%. Overall performance for the systems across the various target obfuscation conditions is shown in Figure 1.

We performed a $2 \times 3 \times 3$ repeated measures ANOVA within the 24 mixture conditions, across the training condition, segment classification method, and target obfuscation condition. This analysis showed main effects for training condition ($F = 124.90$, $p < 0.001$), classification method ($F = 48.51$, $p < 0.001$), and target obfuscation condition ($F = 57.48$, $p < 0.001$), as well as interactions between training condition with classification method ($F = 12.27$, $p < 0.001$) and training condition with target obfuscation ($F = 7.72$, $p = 0.006$). Since results were poor for the

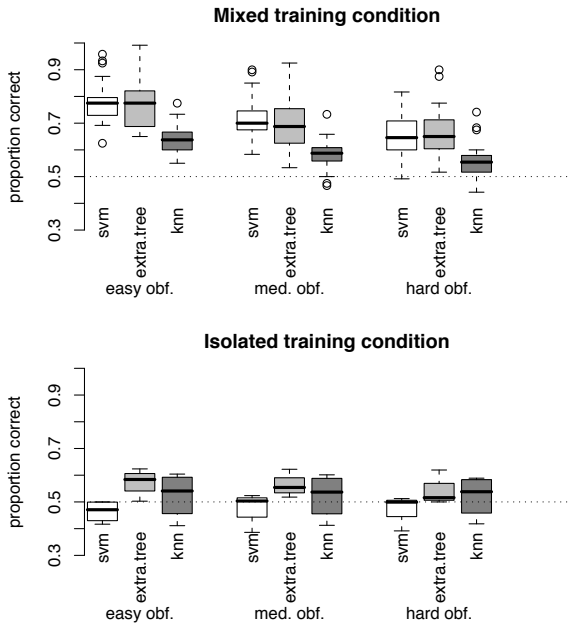

Figure 1. Performance across the three segment classification methods for both the isolated and mixed training conditions. Chance performance is indicated by the dotted line. Here, "obf." stansds for "obfuscation."

isolated conditions, it is not surprising that the other main effects should interact with the training condition.

Post-hoc permutation tests [13] showed that there was no statistically significant difference in performance between the ExtraTree and SVM segment classifiers ($p = 0.845$) on the mixture case, but performance with the KNN classifier was worse from both SVMs and Extra Trees ($p < 0.001$). The the KNN method is known to be sensitive to irrelevant features, since we have yet to consider feature selection, this is one possible reason the KNN method failed to work. All three obfuscation conditions differed from each other ($p < 0.02$). We also confirmed the that the isolated condition was worse than the mixed training condition ($p < 0.001$). We also examined the pattern of performance across different sorts of mixtures: not surprisingly the more similar a target is to its distractors the worse performance is: due to space considerations we have left the breakdown of performance by instrument out.

Only in the mixed training condition was performance better than chance (0.5), and then only for the SVM and Extra Tree classifier ($p < 0.001$). Taken togethor, these results suggest that the SVM and the Extra Tree algorithm can be used to classify instruments in novel mixtures using the segment-and-combine algorithm when the training data are mixtures of sound rather than isolated notes. As obfuscation of the target became greater, accuracy dropped similarly across the three segment classifiers.

4.2 Experiment 2

We performed a second experiment to determine how a more exact labeling of segments would affect our results. To do this we determined the ratio of the target instrument to the total mixture (R) in each segment selected for classification. This was found using the following equation:

$$R = \frac{\|proj_{\mathbf{t}}\mathbf{m}\|^2}{\|\mathbf{m}\|^2} \quad (1)$$

The linear projection of the target onto the mixture, $proj_{\mathbf{t}}\mathbf{m}$, is used as an estimate of the target that remains in the mixture as per [14]. An R of 0.5 means half the energy in a segment is from the target. Given an R for each segment our system is provided a positive label only if it is above some threshold t. This new labeling will be more accurate because segments from positive examples that contain no energy from the target instrument will not be included as a labeled segment. Using this procedure, we trained the Extra-Tree classifier as per the mixture condition in Experiment 1 with the data from the hardest target obfuscation condition. Thus each test at a given threshold consists of 24 trials: 4 instruments with 6 possible arrangements of distractors. Results across various values for the threshold t can be seen in Figure 2.

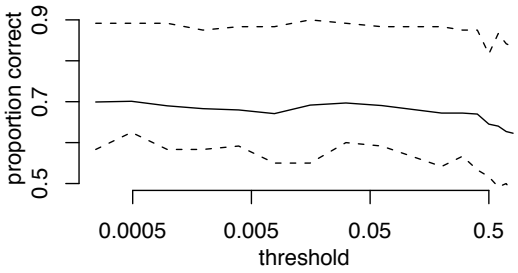

Figure 2. Classification accuracy as a function of the threshold R used to label segments. Dotted lines indicate the maximum and minimum; the solid line indicates the mean.

We can conclude from this that although performance can be improved if we select a fairly inclusive t (i.e. one that includes any segment that has even a small fraction of the target), this will will not yield performance gains of much more than about 4-6%. We would note that the poor performance for high values of t may be attributable to the small sample of segments that actually have a high enough R. For example with $t = 0.5$ only about 19% of the segments from positive examples were included.

5 CONCLUSIONS AND FUTURE WORK

Many researchers have tried to address the problem of identifying instruments in mixtures of sound by trying to mitigate the effect the other instruments have on the target in-

strument to be identified, through, for instance, source separation methods [15], missing feature approaches [5] or weighting of features [10]. Here we presented an alternative approach: rather than being smart about how we use isolated training data, we have simply used a variety of mixtures as our training data.

This simple method makes classificiation possible in mixtures of random notes using only weakly labeled data, when it is not possible to do so when trained with isolated training data, without one of the above mentioned methods. We believe this is because the training data, in the mixed case, is more representative of the testing data, even when the training mixtures do not use the same set of instruments as the testing mixtures. It's possible that the use of mixtures during training could be used in conjuction with these other methods to further improve performance.

We found that when using more strongly labeled data, our results did not improve much (Section 4.2). We believe the most likely explanation of this result is that we have hit a performance ceiling for the chosen features set. This may also explain the similar performance of Extra Trees and the SVM. In future work, now that we have identified a potentially useful training method and learning algorithm for this task, we will begin to explore a more effective set of features.

We will also need to confirm these results on more musical arrangements, and more challenging recordings (e.g. varied articualation, echoes and non-instrumental background, etc...). Since our algorithm does not a priori depend on these simplifications, we suspect the conclusions drawn here will be meaningful in more challenging environments.

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CHORD RECOGNITION USING INSTRUMENT VOICING CONSTRAINTS

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ABSTRACT

This paper presents a technique of disambiguation for chord recognition based on *a-priori* knowledge of probabilities of chord voicings in the specific musical medium. The main motivating example is guitar chord recognition, where the physical layout and structure of the instrument, along with human physical and temporal constraints, make certain chord voicings and chord sequences more likely than others. Pitch classes are first extracted using the Pitch Class Profile (PCP) technique, and chords are then recognized using Artificial Neural Networks. The chord information is then analyzed using an array of *voicing vectors* (VV) indicating likelihood for chord voicings based on constraints of the instrument. Chord sequence analysis is used to reinforce accuracy of individual chord estimations. The specific notes of the chord are then inferred by combining the chord information and the best estimated voicing of the chord.

1 INTRODUCTION

Automatic chord recognition has been receiving increasing attention in the musical information retrieval community, and many systems have been proposed to address this problem, the majority of which combine signal processing at the low level and machine learning methods at the high level. The goal of a chord recognition system may also be low-level (identify the chord structure at a specific point in the music) or high level (given the chord progression, predict the next chord in a sequence).

1.1 Background

Sheh and Ellis [6] claim that by making a direct analogy between the sequences of discrete, non-overlapping chord symbols used to describe a piece of music and word sequence used to describe speech, much of the speech recognition framework in which hidden Markov Models are popular can be used with minimal modification. To represent the features of a chord, they use Pitch Class Profile (PCP) vectors (discussed in Section 1.2) to emphasize the tonal content of the signal, and they show that PCP vectors outperformed cepstral coefficients which are widely used in speech recognition. To recognize the sequence, hidden Markov Models (HMMs) directly analogous to sub-word models in a speech recognizer are used, and trained by the Expectation Maximization algorithm. Bello and Pickens [1] propose a method for semantically describing harmonic content directly from music signals. Their system yields the Major and Minor triads of a song as a function of beats. They also use PCP as the feature vectors and HMMs as the classifier. They incorporate musical knowledge in initializing the HMM parameters before training, and in the training process. Lee and Slaney [5] build a separate hidden Markov model for each key of the 24 Major/Minor keys. When the feature vectors of a musical piece are presented to the 24 models, the model that has the highest possibility represents the key to that musical piece. The Viterbi algorithm is then used to calculate the sequence of the hidden states, i.e. the chord sequence. They adopt a 6-dimensional feature vector called the Tonal Centroid [4] to detect harmonic changes in musical audio. Gagnon *et al* [2] propose an Automatic Neural Network based pre-classification approach to allow a focused search in the chord recognition stage. The specific case of the 6-string standard guitar is considered. The feature vectors they use are calculated from the Barkhausen Critical Bands Frequency Distribution. They report an overall performance of 94.96% accuracy, however, a fatal drawback of their method is that both the training and test samples are synthetic chords consisting of 12 harmonic sinusoids for each note, lacking the noise and the variation caused by the vibration of the strings where partials might not be in the exact multiple of their fundamental frequencies. Yoshioka *et al* [7] point out that there exists mutual dependency between chord boundary detection and chord symbol identification: it's difficult to detect the chord boundaries correctly prior to knowing the chord; and it's also difficult to identify the chord name before the chord boundary is determined. To solve this mutual dependency problem, they propose a method that recognizes chord boundaries and chord symbols concurrently. PCP vector is used to represent the feature. When a new beat time is examined (Goto's[3] method is used to obtain the beat times), the hypotheses (possible chord sequence cadidate) are updated.

1.2 Pitch Class Profile (PCP) Vector

A musical note can be characterized as having a global pitch, identified with a note name and an octave (*e.g. C*4) or a pitch color or pitch class, identified only by the note name independent of octave. The pitch class profile (PCP) technique detects the color of a chord based on the relative content of pitch classes. PCP begins from a frequency representation, for example the Fourier transform, then maps the frequency components into the 12 pitch classes. After the frequency components have been calculated, we get the corresponding notes of each frequency component and find its corresponding pitch class.

2 CHORD ANALYSIS

Our approach deviates from the approaches presented in Section 1 in several key areas. The use of voicing constraints (described below) is the primary difference, but our low-level analysis is also somewhat different from current work. First, current techniques will often combine PCP with Hidden Markov Models. Our aproach analyzes the PCP vector using Neural Networks, using a viterbi algorithm to model chord sequences in time. Second, current techniques normally use window sizes on the order of 1024 samples (23.22 ms). Our technique uses comparatively large window sizes (22050 samples, 500ms). Unlike Gagnon, Larouche and Lefebvre [2], who use synthetic chords to train the network, we use real recordings of chords played on a guitar. Although the constraints and system development are based on guitar music, similar constraints (with different values) may be determined for other ensemble music.

2.1 Large Window Segmentation

Guitar music varies widely, but common popular guitar music maintains a tempo of 80–120 beats per minute. Because chord changes typically happen on the beat or on beat fractions, the time between chord onsets is typically 600–750 ms. Segmenting guitar chords is not a difficult problem, since the onset energy is large compared to the release energy of the previous chord, but experimentation has shown that 500ms frames provide sufficient accuracy when applied to guitar chords for a number of reasons. First, if a chord change happens near a frame boundary, the chord will be correctly detected because the majority of the frame is a single pitch class profile. If the chord change happens in the middle of the frame, the chord will be incorrectly identified because contributions from the previous chord will contaminate the reading. However, if sufficient overlap between frames is employed (*e.g.* 75%), then only one in four chord readings will be inaccurate, and the chord sequence rectifier (see Section 2.3) will take care of the erroneous measure: based on the confidence level of chord recognition and changes in analyzed feature vectors from one frame to the next, the rectifier will select the second-most-likely chord if it fits better with the sequence.

The advantages of the large window size are the accuracy of the pitch class profile analysis, and, combined with the chord sequence rectifier, outweigh the drawbacks of incorrect analysis when a chord boundary is in the middle of a frame. The disadvantage of such a large window is that it makes real-time processing impossible. At best, the system will be able to provide a result half a second after a note is played. Offline processing speed will not be affected, however, and will be comparable to other frame sizes. In our experience, real-time guitar chord detection is not a problem for which there are many real-world applications.

2.2 PCP with Neural Networks

We have employed an Artificial Neural Network to analyze and characterize the pitch class profile vector and detect the corresponding chord. A network was first constructed to recognize seven common chords for music in the keys of C and G, for which the target chord classes are [*C, Dm, Em, F, G, Am, D*]). These chords were chosen as common chords for "easy" guitar songs. The network architecture was set up in the following manner: 1 12-cell input layer, 2 10-cell hidden layers, and 1 7-cell output layer. With the encouraging results from this initial problem (described in Section 4), the vocabulary of the system was expanded to Major (I, III, V), Minor (I, iii, V) and Seventh (I, III, V, vii) chords in the seven natural-root(♮) keys (*C, D, E, F, G, A, B*), totaling 21 chords. Full results are presented in Section 4.

A full set of 36 chords (Major, Minor and Seventh for all 12 keys) was not implemented, and we did not include further chord patterns (Sixths, Ninths etc.). Although the expansion from seven chords to 21 chords gives us confidence that our system scales well, additional chords and chord patterns will require further scrutiny. With the multitude of complex and colorful chords available, it is unclear whether it is possible to have a "complete" chord recognition system which uses specific chords as recognition targets, however a limit of 4-note chords would provide a reasonably complete and functional system.

2.3 Chord Sequence Rectification

Isolated chord recognition does not take into account the correlation between subsequent chords in a sequence. Given a recognized chord, the likelihood of a subsequent frame having the same chord is increased. Based on such information, we can create a sequence rectifier which corrects some of the isolated recognition errors in a sequence of chords. For each frame, the neural network gives a rank list of the possible chord candidates. From there, we estimate the chord transition possibilities for each scale pair of Major and rel-

ative Minor through a large musical database. The Neural Network classification result is provided in the S matrix of size $N \times T$, where N is the size of the chord dictionary and T is the number of frames. Each column gives the chord candidates with ranking values for each frame. The first row of the matrix contains the highest-ranking individual candidates, which, in our experience, are mostly correct identifications by the neural network. Based on the chords thus recognized, we calculate the most likely key for the piece. For the estimated key we develop the chord transition probability matrix A of size $N \times N$. Finally, we calculate the best sequence fom S and A using the Viterbi Algorithm, which may result in a small number of chord estimations being revised to the second or third row result of S.

2.4 Voicing Constraints

Many chord recognition systems assume a generic chord structure with any note combination as a potential match, or assume a chord "chromaticity," assuming all chords of a specific root and color are the same chord, as described above. For example, a system allowing any chord combination would identify [*C*-*E*-*G*] as a *C* Major triad, but would identify a unique *C* Major triad depending on whether the first note was middle *C* (*C*4)or *C* above middle *C* (*C*5). On the other hand, a system using chromaticity would identify [*C*4-*E*4-*G*4] as identical to [*E*4-*G*4-*C*5], the first voicing [1] of a *C* Major triad. Allowing any combination of notes provides too many similar categories which are difficult to disambiguate, and allowing a single category for all versions of a chord does not provide complete information. What is necessary, then, is a compromise which takes into account statistical, musical, and physical constraints for chords.

The goal of our system is to constrain the available chords to the common voicings available to a specific instrument or set of instruments. The experiments that follow concentrate on guitar chords, but the technique would be equally applicable to any instrument or ensemble where there are specific constraints on each note-production component. As an example, consider a SATB choir, with standard typical note ranges, *e.g.* Soprano from *C*4 to *C*6. Key, musical context, voice constraints and compositional practice means that certain voicings may be more common. It is common compositional practice, for example, to have the Bass singing the root (I), Tenor singing the fifth (V), Alto singing the major third (III) and Soprano doubling the root (I).

This *a-priori* knowledge can be combined with statistical likelihood based on measurement to create a bayesian-type analysis resulting in greater classification accuracy using fewer classification categories. A similar analysis can be performed on any well-constrained ensemble, for example a string quartet, and on any single instrument with multiple variable sound sources, for example a guitar. At first, the Piano does not seem to benefit from this method, since any combination of notes is possible, and likelihoods are initially equal. However, if one considers musical expectation or human physiology (hand-span, for example), then similar voicing constraints may be applied.

One can argue that knowledge of the ensemble may not be reasonable *a priori* information—will we really know if the music is being played by a wind ensemble or a choir? The assumption of a specific ensemble is a limiting factor, but is not unreasonable: timbre analysis methods can be applied to detect whether or not the music is being played by an ensemble known to the system, and if not, PCP combined with Neural Networks can provide a reasonable chord approximation without voicing or specific note information.

For a chord played by a standard 6-string guitar, we are interested in two features: what chord is it and what voicing of that chord is it [2]. The PCP vector describes the chromaticity of a chord, hence it does not give any information on specific pitches present in the chord. Given knowledge of the relationships between the guitar strings, however, the voicings can be inferred based the *voicing vectors* (VV) in a certain category. VVs are produced by studying and analyzing the physical, musical and statistical constraints on an ensemble. The process was performed manually for the guitar chord recognition system but could be automated based on large annotated musical databases.

Thus the problem can be divided into two steps: determine the category of the chord, then determine the voicing. Chord category is determined using the PCP vector combined with Artificial Neural Networks, as described above. Chord voicings are determined by matching harmonic partials in the original waveform to a set of context-sensitive templates.

3 GUITAR CHORD RECOGNITION SYSTEM

The general chord recognition ideas presented above have been implemented here for guitar chords. Figure 1 provides a flowchart for the system. The feature extractor provides two feature vectors: a PCP vector which is fed to the input layer of the neural net, and an voicing vector which is fed to the voicing detector.

Table 1 gives an example of the set of chord voicing arrays and the way they are used for analysis. The fundamental frequency (f_0) of the root note is presented along with the f_0 for higher strings as multiples of the root f_0.

The Guitar has a note range from *E*2 (82.41Hz, open low string) to *C*6 (1046.5Hz, 20th fret on the highest string). Guitar chords that is above the 10th fret (*D*) are rare, thus we can restrict the chord position to be lower than the 10th

[1] A voicing is a chord form where the root is somewhere other than the lowest note of the chord

[2] Although different voicings are available on guitar, a reasonable assumption is that they are augmented with a root bass on the lowest string

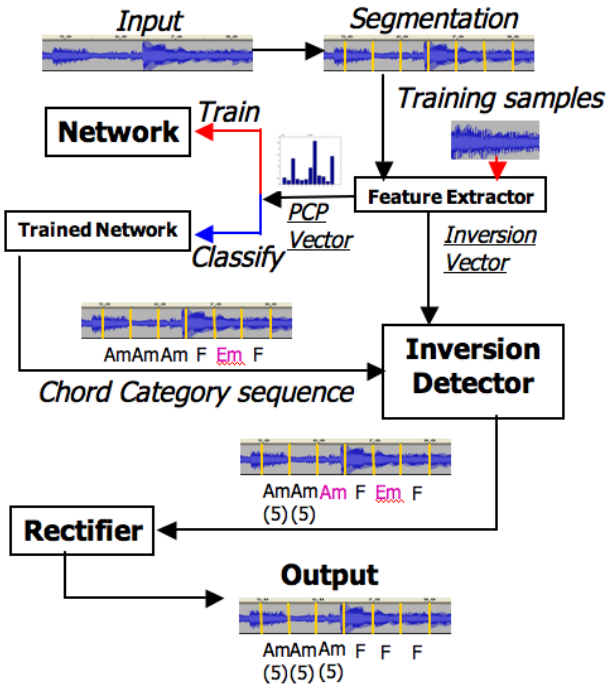

Figure 1. Flowchart for the chord recognition system.

fret, that is, the highest note would be 10th fret on the top string, i.e. *D*5, with a frequency of 587.3Hz. Thus if we only consider the frequency components lower than 600Hz, the effect of the high harmonic partials would be eliminated.

Each chord entry in Table 1 provides both the frequencies and first harmonics of each note. "Standard" chords such as Major, Minor and Seventh, contain notes for which f_0 is equal to the frequency of harmonic partials of lower notes, providing consonance and a sense of harmonic relationship. This is often be seen as a liability, since complete harmonic series are obscured by overlap from harmonically related notes, but our system takes advantage of this by observing that a specific pattern of harmonic partials equates directly to a specific chord voicing. Table 1 shows this by detailing the pattern of string frequencies and first harmonic partials. First harmonic partials above *G6* are ignored since they will not interact with higher notes. Harmonic partials above 600Hz are ignored, since there is no possibility to overlap the fundamental frequency of higher notes (as described above). These are indicated by the symbol "ø". In this way, we construct a pattern of components that are expected to be present in a specific chord as played on the guitar. A string that is not played in the chord is indicated by "–". Boxes and underlines are detailed below. Table 2 shows the same information for voicings of a single chord in three different positions, showing how these chords can be disambiguated.

Chord	S1	S2	S3	S4	S5	S6
f_0 (Hz)	H1	H2	H3	H4	H5	H6
F	1	1.5	2	2.52	3	4
87.31	2	3	4	ø	ø	ø
Fm	1	1.5	2	2.38	3	4
87.31	2	3	4	ø	ø	ø
F7	1	1.5	1.78	2.52	3	4
87.31	2	3	3.56	ø	ø	ø
G	1	1.26	1.5	2	2.52	4
98	2	2.52	3	4	ø	ø
Gm	1	1.19	1.5	2	3	4
98	2	2.38	3	4	ø	ø
G7	1	1.26	1.5	2	2.52	3.56
98	2	2.52	ø	ø	ø	ø
A	–	1	1.5	2	2.52	3
110	–	2	3	ø	ø	ø
Am	–	1	1.5	2	2.38	3
110	–	2	3	ø	ø	ø
A7	–	1	1.5	1.78	2.52	3
110	–	2	3	ø	ø	ø
C	–	1	1.26	1.5	2	2.52
130.8	–	2	2.52	ø	ø	ø
Cm	–	1	1.19	1.5	2	–
130.8	–	2	ø	ø	ø	–
C7	–	1	1.26	1.78	2	2.52
130.8	–	2	2.52	ø	ø	ø
D	–	–	1	1.5	2	2.52
146.8	–	–	2	ø	ø	ø
Dm	–	–	1	1.5	2	2.38
146.8	–	–	2	ø	ø	ø
D7	–	–	1	1.5	1.78	2.52
146.8	–	–	2	ø	ø	ø

Table 1. Chord pattern array, including three forms of five of the natural-root chords in their first positions. S1–S6 are the relative f_0 of the notes from the lowest to highest string, and H1–H6 are the first harmonic partial of those notes. See text for further explanation of boxes and symbols.

Chord	S1	S2	S3	S4	S5	S6
f_0 (Hz)	H1	H2	H3	H4	H5	H6
G	1	1.26	1.5	2	2.52	4
98	2	2.52	3	4	ø	ø
G(3)	1	1.5	2	2.52	3	4
98	2	3	4	ø	ø	ø
G(10)	–	1	1.5	2	2.52	3
196.0	–	2	3	ø	ø	ø

Table 2. Voicing array for the *Gmaj* chords played on different positions on the guitar.

3.1 Harmonic Coefficients and Exceptions

It can be seen from Table 1 that there are three main categories of chords on the guitar, based on the frequency of the second note in the chord. The patterns for the three categories are: (1.5), where the second note is (V): *F, Fm, F7, E, Em, E7, A, Am, A7, B, Bm, D, Dm, D7*; (1.26), where the second note is (III): *B7, C7, G, G7*; and (1.19), where the second note is (iii): *Cm, Gm*.

Thus, from the first coefficient (the ratio of the first harmonic peak to the second) we can identify which group a certain chord belongs to. After identifying the group, we can use other coefficients to distinguish the particular chord. In some situations (*e.g.*, *F* and *E*; *A* and *B*), the coefficients are identical for all notes in the chord, thus they cannot be distinguished in this manner. Here, the chord result will be disambiguated based on the result of the Neural Network and the f_0 analysis of the root note. Usually, all first harmonic partials line up with f_0 of higher notes in the chord. When the first harmonic falls between f_0 of higher notes in the chord, they are indicated by $\boxed{\text{boxed}}$ coefficients.

Underlined coefficients correspond to values which may be used in the unique identification of chords. In these cases, there are common notes within a generic chord pattern, for example the root (1) and the fifth (1.5). String frequencies corresponding to the Minor Third (1.19, 2.38) and Minor Seventh (2.78) are the single unique identifiers between chord categories in many cases.

4 RESULTS

Chord detection errors do not all have the same level of "severity". A *C* chord may be recognized as an *Am* chord (the relative Minor), since many of the harmonic partials are the same and they share two notes. In many musical situations, although the *Am* chord is incorrect, it will not produce dissonance if played with a *C* chord. Mistaking an *C* chord for a *Cm* chord, however, is a significant problem. Although the chords again differ only by one note, the note in question is more harmonically relevant and differs in more harmonic partials. Further, it establishes the mode of the scale being used, and, if played at the same time as the opposing mode, will produce dissonance.

4.1 Chord Pickout

Chord Pickout [3] is a popular off-the-shelf chord recognition system. Although the algorithm used in the *Chord Pickout* system is not described in detail by the authors, it is reasonable to compare with our system since Chord Pickout is a commercial system with good reviews. We applied the same recordings to both systems and identified the accuracy of each system. We were more forgiving with the analysis for Chord Pickout in order to better detail the types of errors that were made. If Chord Pickout was able to identify the root of the chord, ignoring Major, Minor or Seventh, it is described as "correct root." If the chord and the chord type are both correct, it is described as "correct chord." Errors between correct root and correct chord included Major→Minor, Major→Seventh, and Minor→Major. For our system, all chord errors regardless of severity, are considered incorrect. The complete results for 6 trials are presented in Table 3.

[3] `http://www.chordpickout.com`

4.2 Independent accuracy trials

To detect the overall accuracy of our system, independent of a comparison with another system, we presented a set of 40 chords of each type to the system and evaluated its recognition accuracy. Two systems were trained for specific subsets of chord detection.

The first system was trained to detect chords in the a single key only assuming key recognition has already taken place. Seven chord varieties are available as classification targets, and the system performed well, producing 96.8% accuracy over all trials. Misclassifications were normally toward adjacent chords in the scale.

The second system was trained to recognized Major, Minor and Seventh chords of all seven natural-root keys, resulting in 21 chord classification targets. This system produced good results: 89% for Major versus Minor, and 75% accuracy for Major versus Seventh chords. Table 4 provides a confusion matrix between single-instance classification of Major and Seventh chords, which had the lower recognition rate. There are two reasons for this: in some cases the first three notes (and correspondingly the first three harmonic partials detected) are the same between a chord and its corresponding Seventh; and in some cases the first harmonic of the root note does not line up with an octave and thus contributes to the confusion of the algorithm. Recognition accuracy is highest when only the first two notes are the same (as in *C* and *G* chords). Recognition accuracy is low in the case of *F7*,when the root is not doubled, and the pattern can be confused with both the corresponding Major chord and the adjacent Seventh chord. Recognition accuracy is also low in the case of *G7*, where the difference between the Major and the Seventh is in the third octave, at 3.56 times the fundamental of the chord. In this case, the Seventh chord is frequently mistaken for the Major chord, which can be considered a less "severe" error since the Seventh chord is not musically dissonant with the Major chord.

A more severe case is with *D7*, which contains only 4 sounded strings, one of which produces a harmonic that does not correspond to a higher played string. From Table 1, we can see that the string frequency pattern for *D7* is [1, 1.5, 1.78, 2.25], and the first harmonic partial of the root note inserts a 2 into the sequence, producing [1, 1.5, 1.78,

		Voicing Constraints		Chord Pickout			
Trial	Frames	Correct	Rate	Correct Root	Rate	Correct Chord	Rate
1	24	23	95.8%	20	83.3%	1	5.0%
2	46	44	95.6%	30	65.2%	12	26.1%
3	59	54	91.5%	38	64.4%	7	11.9%
4	50	49	98.0%	31	62.0%	30	60.0%
5	65	51	78.4%	51	78.4%	21	32.3%

Table 3. Comparison of our system to *Chord Pickout*, an off-the-shelf chord recognition system.

Chord	Rate	C	C7	D	D7	E	E7	F	F7	G	G7	A	A7
C	40/40	40											
C7	35/40		35		2	2						1	
D	40/40			40									
D7	13/40			1	13		1	3	20				2
E	40/40					40							
E7	37/40						37				3		
F	38/40					1	1	38					
F7	5/40					3	16	16	5				
G	40/40									40			
G7	17/40				2					21	17		
A	30/40			1	8							30	1
A7	25/40	5	2								8		25

Table 4. Confusion Matrix for Major and Seventh chords of natural-root keys. Overall accuracy is 75%.

2, 2.25]. This is very similar to the sequence for *F7*, which is why the patterns are confused. It would be beneficial, in this case, to increase the weight ascribed to the fundamental frequency when the number of strings played is small. Unfortunately, detecting the number of sounded strings in a chord is a difficult task. Instead, f_0 disambiguation can be applied when a chord with fewer strings is one of the top candidates from the table, since that information is known.

5 CONCLUSIONS

A chord detection system is presented which makes use of voicing constraints to increase accuracy of chord and chord sequence identification. Although the system is developed for guitar chords specifically, similar analysis could be performed to apply these techniques to other constrained ensembles such as choirs or string, wind, or brass ensembles, where specific chords are more likely to appear in a particular voicing given the constraints of the group.

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APPLICATION OF THE FUNCTIONAL REQUIREMENTS FOR BIBLIOGRAPHIC RECORDS (FRBR) TO MUSIC

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ABSTRACT

This paper describes work applying the Functional Requirements for Bibliographic Records (FRBR) model to music, as the basis for implementing a fully FRBR-compliant music digital library system. A detailed analysis of the FRBR and Functional Requirements for Authority Data (FRAD) entities and attributes is presented. The paper closes with a discussion of the ways in which FRBR is gaining adoption outside of the library environment in which it was born. This work benefits the MIR community by demonstrating a model that can be used in MIR systems for the storage of descriptive information in support of metadata-based searching, and by positioning the Variations system to be a source of robust descriptive information for use by third-party MIR systems.

1. THE FUNCTIONAL REQUIREMENTS FOR BIBLIOGRAPHIC RECORDS (FRBR)

Functional Requirements for Bibliographic Records (FRBR) is a conceptual model released in 1997 by the International Federation of Library Associations and Institutions (IFLA). FRBR provides a "framework that would provide a clear, precisely stated, and commonly shared understanding of what it is that the bibliographic record aims to provide information about, and what it is that we expect the record to achieve in terms of answering user needs" [9] (p. 2). In short, FRBR provides a model for the "bibliographic universe" by describing the entities and attributes that make up library metadata.

FRBR entities are divided into three groups. Group 1 entities are the "products of intellectual or artistic endeavour," and include Work ("a distinct intellectual or artistic creation"), Expression ("the intellectual or artistic realization of a Work"), Manifestation ("the physical embodiment of an Expression of a Work") and Item ("a single exemplar of a manifestation"), as seen in Figure 1 [9] (p. 13). The FRBR Group 2 entities are those that create or act upon Group 1 entities: Person and Corporate Body. FRBR Group 3 entities are the subjects of Works: Concept, Object, Event, and Place.

The application of FRBR to music has been described (though mostly in passing) at past ISMIR conferences [4][8][18][19][22][23]. More in-depth studies of FRBR for musical materials appear in the library and information science literature. LeBoeuf performs a semiotic analysis of FRBR for musical works, raising crucial questions regarding the boundaries of Works and proposing revised definitions for the FRBR Group 1 entities [12]. Miller and LeBoeuf [14] analyze the application of the FRBR Group 1 entities to live events of performing arts, focusing on mixed-media works including drama or choreography. Vellucci, in the most systematic published analysis to date, considers what the Work and Expression entities and their various attributes mean for music [25].

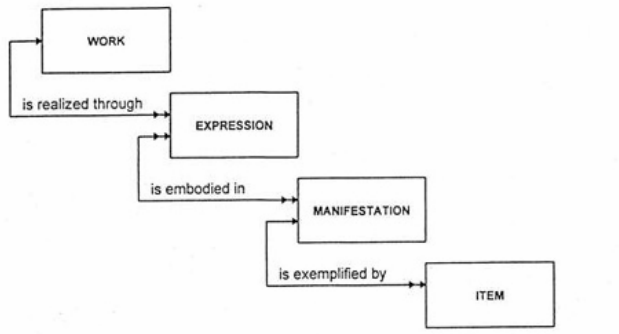

Figure 1. The FRBR group 1 entities [9] (p. 14)

FRBR has also served as the underlying conceptual model for a small number of music digital libraries, most notably MusicAustralia [1] and the evolving Probado system at the Bavarian State Library [4]. While research reports from some of these initiatives are available, specific details needed as a guide to local FRBR adoption efforts are generally more difficult to obtain.

Despite the work that has been done to study and test implementation of FRBR in music digital libraries, it is still not clear how the model should be applied to a newly-developed system. Even taking into account the move of some FRBR implementations for music towards production-level systems, the music library and MIR communities' understanding of FRBR exists largely at a theoretical rather than a practical level. The work presented in this paper represents a significant step forward in demonstrating how this theory can be turned into practice.

2. THE VARIATIONS SYSTEM AND FRBR

2.1. Variations music metadata

Like FRBR, the Variations digital music library system at Indiana University has been discussed at previous ISMIR conferences [5][6][8][19][22], demonstrating the value of contributions from the academic music library and digital library communities to MIR research. Variations2 was a research and development project investigating such wide areas as system architecture, metadata standards, component-based application architecture, and network services. A system that provides access to streaming audio and scanned scores developed as part of the Variations2 project has been put into production at the Cook Music Library in the Indiana University Jacobs School of Music. A follow-on project known as Variations3 is currently underway, which aims to provide a version of the software that can be deployed at other institutions, and continues some of the music metadata research started in the earlier project. The source code for the Variations Audio Timeliner[1] and the Variations metadata model, described below, both show promise for use within MIR systems.

The Variations system uses a work-based metadata model which has been described in more detail at previous ISMIR conferences [8][22] and elsewhere [16]. While the Variations metadata model is similar in many ways to FRBR, it was not envisioned as a FRBR implementation at the time of its creation, relying heavily instead on the music modeling work of Richard Smiraglia [24]. Table 1 shows a partial summary of the differences between the Variations model and FRBR.

Variations2/3 Entity	FRBR Group 1 Entity
Work (more concrete than FRBR Work)	Work
Instantiation (can only appear on one Container)	Expression
Container (includes some copy-specific data)	Manifestation
Media Object (defined as a digital file)	Item

Table 1. Comparison of Variations and FRBR models

The FRBR model is increasingly taking hold in the library environment, and shows promise for improving interoperability of data within libraries and beyond, including with the MIR community. The increasing uptake of FRBR led the Variations team to embark upon an in-depth study of how the FRBR model applies to musical materials. In order to make our work as useful as possible to others outside of our home institution, we structured our investigations as direct analysis of FRBR, rather than presenting our analysis as incremental changes to our existing data model.

We also limited our work to applying the FRBR model to scores and recordings, rather than to other types of material held by music libraries or managed by MIR systems. With this approach we did not model musical works as separate from texts included in them, in contrast to the method that some have proposed as ideal [1]. This decision is largely a practical one, limiting the scope of study to materials in the Variations system.

To function as a production service, Variations also requires a variety of structural and technical metadata. While some attributes described in the FRBR report could be used to support these functions, FRBR is primarily focused on descriptive metadata. Our team took the approach of using FRBR for the descriptive metadata in Variations only, with plans to model structural and technical metadata separately.

2.2. FRBR Group 1 Entities

The Variations team's initial efforts to study the FRBR model focused on the FRBR Group 1 entities: Work, Expression, Manifestation, and Item. We created operational definitions of each of the Group 1 entities as they applied to music, evaluated each attribute and relationship for its utility for the description of musical materials, and noted areas of description needed for music that were not covered in the FRBR report. Detailed results of our analysis are documented in a white paper freely available on the project web site [20].

2.2.1. Defining Work and Expression

One area of particular interest for music is the boundary between FRBR Works and Expressions. Phrases in the FRBR report suggest a Work for music should be interpreted broadly rather than strictly, including passages such as "addition of parts or an accompaniment to a musical composition" and "musical transcriptions and arrangements" as referring to new Expressions rather than new Works [9] (p. 17). Based on this approach, our operational definition for a Work is liberal and abstract, following the FRBR characterization of "adaptation" as a relationship between two Works, and "arrangement" as a relationship between two Expressions of the same Work.

This basic definition functions well for music in the canon of Western art music, where the composition takes precedence over any given performance of it. For other types of music, the notion of an abstract work is less natural. The Variations team proceeded nevertheless to create definitions of "Work" for music from these other traditions, believing that compromises for the modeling of

[1] http://sourceforge.net/projects/variations/

specific cases are worth the benefits of using a single model throughout the system. For jazz, our operational definition of a FRBR Work for music is the "tune." Performances of that tune, even widely diverging in nature, would be Expressions of that same Work. A fundamental transformation of the work would be considered a new Work. For pop music (itself a difficult-to-define category), we defined the "song" as a Work. Covers and different performances by the same artist or group are therefore considered separate Expressions of that same Work. An album, when it represents a cohesive artistic whole, can also be considered a Work, with a whole-part relationship to the individual songs on the album. World and non-Western music, including systems that operate in an oral rather than a written tradition, pose a particular challenge to a work-based model. The Variations team's definition of a Work for music from these traditions is necessarily vague, and specifies that the cultural or ethnic group responsible for the music and the location and context in which the music is generally performed should be major factors in the decision as to whether two performances reflect two Works or different Expressions of the same Work.

A specific definition of Expression for music requires even more interpretation of the FRBR report than Works. It is clear from the FRBR report that Expressions have a defined form; that is, music represented in visual notation is a different Expression than music represented as an audio recording. Yet since different arrangements are considered different Expressions of the same Work, for audio recording Manifestations, each embodied Expression represents *both* an arrangement and a specific interpretation of that arrangement as a musical performance. The combining of the abstract notion of arrangement together with the slightly more concrete idea of realizing a Work in a particular notation or performance seems to reflect a weakness of the FRBR model for the performing arts. Vellucci in response to this issue proposes that FRBR entities have "subentities." In this approach, notation and performance are separate Expressions, each with "sub-Expressions" for particular notations or performances [25] (p. 142-4). The Variations team did not take this approach, as it is possible to connect performances of the same arrangement Expression with mechanisms described in the FRBR report through the use of Expression-Expression relationships. While our solution might not give the notion of arrangement the intellectual primacy it holds in typical discussions of Western art music, it appears to be sufficient to support the requirements of Variations.

2.2.2. Balancing rigor with practicality

In our work applying FRBR to music, the Variations team faced difficult compromises between rigorous data modeling and practical considerations. In some cases our team determined additional levels of complexity were not required, and in others we believed there was benefit to following the FRBR model to its fullest. We took the former approach in two notable cases. First, due to our early decision that our model would only create entities representing musical content, we decided to track language of liner notes and other material appearing on the Manifestation as an attribute of that Manifestation, although a different FRBR interpretation might model the liner notes as a separate Work with an Expression that appears together on the recording Manifestation. Second, although multiple volumes in a set can be modeled as separate Manifestations with whole-part relationships to a set-level Manifestation according to the FRBR report, the Variations team determined that this would not be necessary in most cases for our purposes.

We took the approach of following more complex features of FRBR more often, allowing for more robust services to be built on our data. The current Variations system allows us to track the date of first performance of a musical Work (i.e., its first performance Expression), as this data can be used to support a wider range of research questions than most current library catalogs. In order to support this type of discovery in a FRBR-based system, an Expression should be modeled for the original performance, even if the Variations system did not include a recording of that performance. We also determined that the recording held by the library should be modeled as a separate Manifestation than the WAV file generated from it, and that the MP3 file for streaming to end-users was a third Manifestation.

FRBR Items presented a challenge in our analysis. The FRBR report describes Items as physical entities, and does not explicitly cover the case where a Manifestation exists only as a digital file rather than in physical form. While there is some disagreement in the community as to how digital files should be handled [7], the Variations team concluded that the intent of the FRBR authors was that digital files should be FRBR Items. The Item-Item "reproduction" relationship will be used to record which copy of a physical recording was digitized to create a particular WAV file. For digitized scores, each individual page might be modeled as a separate Item[2], although our team is reserving a final decision on this issue until we complete our work on specifying technical metadata needed for effective system operation. These decisions represent a fairly complex model for Items, beyond what is implied by the text of the FRBR report. We expect, however, that most Item attributes and relationships will be handled by Variations at the system level and not require manual intervention by a cataloger.

[2] The Probado project's implementation of FRBR introduces a "file" entity to avoid this situation. [4]

2.2.3. Additional attributes needed

The Variations team excluded a number of attributes and relationships defined in the FRBR report from local interpretation of the model, as they were not appropriate to the description of musical material. Our analysis also uncovered a few attributes that were not present in the FRBR model, but nevertheless would be necessary for the Variations system. The "missing" attributes generally fell into three categories: those necessary due to simplifications to FRBR we instituted for practical reasons, those needed for efficient system operation, and those that appear to have simply been overlooked in the development of the FRBR report.

The first case, attributes necessary due to our decision to preference musical content over other types, can be seen in the need to record a language for a Work, since text settings are an integral part of musical works and in our interpretation are not considered to be separate Works with their own Expressions. The second case, attributes to assist with system operation, represent both new attributes and refinements of FRBR-defined attributes. An identifier attribute for Work and location, call number, and copy number for Item are examples of this second case.

The third case, attributes missing entirely from FRBR, represent our most significant departure from the FRBR model. Based on initial reactions to our findings from members of the FRBR Review Group[3], we are optimistic that our findings will influence the growth of the FRBR model into the future. The attributes we noted as necessary for our local implementation but not obviously present in the FRBR report are place of composition and genre/form/style for Works, and place of performance, key, and genre/form/style for Expressions. We also noted that data essential for the understanding of non-Western music, such as geography, culture, event, and function could be recorded within FRBR-defined Work attributes, but that FRBR could be adjusted to model this type of information in a more prominent way.

2.3. Next Steps and Implementation Plans

The Variations team's initial study of the FRBR model extended only to the FRBR Group 1 Entities, assuming that if they matched the needs of the Variations system, then the rest of the FRBR model would as well. Our analysis demonstrated that moving the Variations system to a fully FRBRized model would be both possible and desirable. This decision was only the first step in what will be a much longer process.

Our next area of study was to extend our detailed analysis of the applicability of FRBR to musical material to the Group 2 (Person and Corporate Body) and Group 3 (Concept, Object, Event, and Place) entities and their attributes. We also considered in this second phase of our analysis the model presented by the FRBR companion report *Functional Requirements for Authority Data* (FRAD), released in draft in April 2007 [10]. This second phase of our FRBR analysis proved more straightforward than the first, with relatively little interpretation of FRBR/FRAD entities and attributes needed for musical materials. A report describing this phase two analysis is available from the Variations3 project web site [21].

The Variations team concluded that modeling Names/Identifiers and Controlled Access Points as separate entities, as outlined in FRAD, was not necessary for a single-language system of the scale of Variations. We adjusted our phase one interpretations in a few cases, modeling features such as form/genre and instrumentation as FRBR/FRAD Group 3 Concept entities rather than as attributes on Work or Expression, but stopped short of large-scale conversion of attributes to relationships to Group 3 entities, in keeping with the spirit of the FRBR report limiting Group 3 entities to "subjects" of Works.

We expect to embark upon a third planning stage in the late summer and fall of 2008, where we will adopt a FRBR encoding syntax, or develop our own if necessary. FRBR as a conceptual model does not define a formal data structure for recording FRBR data, and one has not emerged from the library community from a trusted maintenance agency that would represent an obvious choice for adoption by Variations. A few FRBR encodings have emerged from efforts outside of the core library community, however, and the Variations team plans to study each of these to determine if they meet the functional requirements for our system, and to analyze their likely sustainability and supportability over time.

3. ADOPTION OF FRBR AND IMPLICATIONS FOR MUSIC INFORMATION RETRIEVAL

While FRBR originated within libraries as an effort to better understand the bibliographic universe, it has also attracted attention from outside the library community. An expression of FRBR in RDF, allowing FRBR data to interact in Semantic Web environments, has arisen from a small but diverse group of information professionals [2]. The museum community recognized similarities between FRBR and its CIDOC Conceptual Reference Model[4], and organized an effort to create an object-oriented version of FRBR that aims to harmonize the two models [11]. The enthusiast- and researcher-driven Music Ontology, expressed in the OWL Web Ontology Language[5], uses FRBR together with other existing frameworks such as Friend of a Friend (FOAF)[6] to provide a "formal framework for dealing with music-related information on

[3] http://www.ifla.org/VII/s13/wgfrbr/

[4] http://cidoc.ics.forth.gr/index.html

[5] http://www.w3.org/2004/OWL/

[6] http://www.foaf-project.org/

the Semantic Web, including editorial, cultural and acoustic information" [18] (p. 417). The Music Ontology especially falls within the area of interest for the MIR community, and these developments can be interpreted as a sign that the FRBR approach has proven useful beyond libraries, including in MIR research.

Within the library environment, the FRBR model is gaining significant traction as well. While a number of systems have implemented FRBR or FRBR-like models in test environments, more production-scale implementations are starting to emerge, such as OCLC's WorldCat.org.[7] A recent report made to the Library of Congress concerning the future of library cataloging recognized the potential of the FRBR model for enhancing library services, but called for more testing of it in production situations [13]. A new content standard for library cataloging records called Resource Description and Access (RDA)[8] is in development, which will be based on FRBR principles. These and other changes are driving investigations into the use of record structures other than the current MARC standards in ubiquitous use in libraries [3][15]. It seems clear that FRBR has taken hold as a fundamental guiding model for ongoing metadata development in libraries. We expect that Variations, as a FRBR-based system, will serve as a model for these developments.

The library community has recognized that this time of change in our system architectures and standards presents us with an opportunity to participate more fully in the global information environment. The activity in this area most relevant to the ISMIR community is the work of the Dublin Core Metadata Initiative (DCMI)/RDA Task Group. This group has three primary goals: the definition of the "elements" presented by RDA as Resource Description Framework (RDF)[9] properties and classes, the exposure of in-line vocabularies defined in RDA for use by Semantic Web tools, possibly in RDF Schema (RDFS)[10] or Simple Knowledge Organization System (SKOS)[11], and the development of a DCMI application profile for FRBR and FRAD.[12] When library-generated metadata is openly exposed in formats such as these, communities such as MIR can benefit.

Indiana University's plans to move Variations to a FRBR-based model, if a pending funding application is awarded to support the implementation phase of our work, would be of particular advantage to the MIR community in two ways. First, structured music metadata in standards-based and commonly-understood forms would be available to MIR researchers long before the vast amounts of library metadata as a whole could undergo a similar transformation. Such wide exposure of library-generated music metadata would allow it to be used in new environments, such as the emerging linked data movement [17]. This metadata could be used as a source of ground truth, as information for display to users in systems demonstrating MIR research, in support of production music discovery and use systems, or even as primary data worthy itself of analysis and study. Second, the interpretation of the FRBR model for musical materials that the Variations team has performed could serve as a model for MIR systems that wish to store descriptive metadata as the basis of metadata-based searching. The potential for advancing the state of MIR research as a result of the Variations/FRBR harmonization and other related initiatives in the library sector is indeed vast.

[7] http://www.worldcat.org. Note that this system is not a full FRBR implementation; it lacks a robust Expression entity.
[8] http://www.collectionscanada.ca/jsc/rda.html
[9] http://www.w3.org/RDF/
[10] http://www.w3.org/TR/rdf-schema/
[11] http://www.w3.org/2004/02/skos/
[12] http://dublincore.org/dcmirdataskgroup/

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ONLINE ACTIVITIES FOR MUSIC INFORMATION AND ACOUSTICS EDUCATION AND PSYCHOACOUSTIC DATA COLLECTION

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ABSTRACT

Online collaborative activities provide a powerful platform for the collection of psychoacoustic data on the perception of audio and music from a very large numbers of subjects. Furthermore, these activities can be designed to simultaneously educate users about aspects of music information and acoustics, particularly for younger students in grades K-12. We have created prototype interactive activities illustrating aspects of two different sound and acoustics concepts: musical instrument timbre and the cocktail party problem (sound source isolation within mixtures) that also provide a method of collecting perceptual data related to these problems with a range of parameter variation that is difficult to achieve for large subject populations using traditional psychoacoustic evaluation. We present preliminary data from a pilot study where middle school students were engaged with the two activities to demonstrate the potential benefits as an education and data collection platform.

1 INTRODUCTION

Recently, a number of web-based games have been developed for the purpose of large-scale data labeling [1]. Similar activities can be used to collect psychoacoustic data from a large number of users, which is difficult to obtain using traditional evaluations. We present two such activities that explore the perception of audio (instrument timbre and the cocktail party problem), with the additional aim of educating users, particularly K-12 students, about aspects of music and acoustics. These web-based interfaces are designed as game activities with minimal complexity so they can be easily used by students without previous training. To maintain widespread accessibility, the activities require only internet access through a web browser and run independently of external applications. We believe this platform will enable us to collect a very large number of samples exploring myriad parameter variations to better define perceptual boundaries and quantitative models of perceived features.

2 BACKGROUND

Relatively little research has been conducted on human performance in the identification of musical instruments after timbral modifications. Saldanha and Corso demonstrated that the highest performance is achieved when the test tone consists of the initial transient and a short steady-state segment and the lowest performance occurs using test tones consisting of only the steady-state component and the ending transient [2]. Iverson examined the dynamic attributes of timbre by evaluating the effects of the transient in sound similarity tests. While results of instrument identification experiments depended on the presence of the initial transient in the sound, the results of similarity tests using tones with and without the initial transient suggest that the initial transient is not required to imply similarity. This research suggests that similarity judgments may be attributed to acoustic properties of an instrument's sound other than the transient component [3]. Martin demonstrated that humans can identify an instrument's family with greater accuracy than the instrument itself [4]. Other studies on instrument identification suggest that musically inclined test subjects perform better than non-musically inclined subjects and in particular, subjects with orchestra experience perform better than those without [5].

The performance of automatic speaker and speech recognition algorithms often use human performance as a benchmark, but it is difficult to obtain a large human subject population for comparisons. Atal [6] provides a summary of human speaker recognition evaluations before 1976 in which no more than 20 human listeners are employed for each evaluation. Stifelman [7] performed a speech recognition evaluation that simulated the environment of the cocktail party problem, testing the listening comprehension and target monitoring ability of 3 pilot and 12 test subjects. Lippmann [8] provides a summary of several human vs. machine speech recognition comparisons, all of which distinctly show humans outperforming machines with a limited number of test subjects. In the area of speaker recognition, Schmidt-Nielsen [9] conducted an evaluation in which 65 human listeners were tested with speech data from the 1998 NIST automatic speaker recognition evaluations. The experiment was administered in the same manner as the automated sys-

tems to create a direct comparison, and the results show that humans perform at the level of the best automated systems and exceed the performance of typical algorithms.

3 DEVELOPED ACTIVITIES

3.1 Timbre Game

The Timbre Game is an online activity designed to illustrate the effects of particular acoustic modifications on musical instrument timbre and to collect evaluation data on the perception of timbre. The user interface for the Timbre Game was developed in Java and is accessed by downloading applets within a client's web browser, requiring no additional software. The Timbre Game has two user interfaces, labeled "Tone Bender" and "Tone Listener". The objective of Tone Bender is primarily educational in that a player is allowed to modify time and frequency characteristics of musical sounds and immediately hear how those sounds are affected. Upon modification, the player submits the resulting sound they have created to a server database. In Tone Listener, other players will listen to the sounds whose timbre has been modified by a player in the prior component. Players will then attempt to determine the original instrument source from the modified timbre. Points are awarded to both players (modifier and listener) if the listening player enters the sound source's identity correctly. A more detailed description of each component of the activity follows.

3.1.1 Tone Bender

The Tone Bender game involves time and frequency analysis of a single instrument sound and provides a visual interface which allows a player to modify the sound's timbre. The sounds available for analysis are 44.1 kHz recordings of various musical instruments, each producing a single note with a duration of less than five seconds.

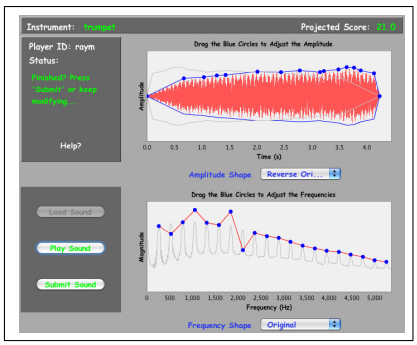

Figure 1. The Tone Bender interface

A player starts a session in Tone Bender by requesting a sound file from the server database. The sound is then analyzed in both the time and frequency domains to generate control points suitable for modifying the sound's timbre. In the time domain analysis, a heuristic method is employed to approximate the sound's amplitude envelope by picking amplitude peaks within small time intervals of the sound wave. These peaks are used as the initial amplitude envelope control points.

In the frequency domain, the sound is analyzed via an efficient Short-Time Fourier Transform (STFT) implementation optimized for Java, using 45 msec Hanning windows with 50 percent overlap. The STFTs are used to generate a time-averaged spectrum of the sound. Linear prediction is employed to establish a threshold, which is used to extract twenty of the most prominent spectral peaks from the time-averaged spectrum. These spectral peaks are used as the initial control points for the harmonic weights that the player will modify.

The visual representation of the sound's timbre is displayed to the player in two separate 'XY' plots in the Java applet as shown in Figure 1. In the amplitude plot, the sound wave is shown with the extracted amplitude control points. The player is allowed to manipulate the shape of the amplitude envelope as they wish by clicking and dragging the control points within the plot window. In the frequency plot, the time-averaged spectrum of the sound wave is shown with the spectral control points. The player is allowed to move the control points vertically so that they are only adjusting the harmonic weights of the sound without affecting pitch. After modifying the spectral envelope, the sound wave is resynthesized using additive sinusoidal synthesis and redrawn in the time plot.

After each modification in Tone Bender, the player is presented with a potential score based on the difference between the original sound and the altered sound, calculated using the signal-to-noise ratio (SNR):

$$SNR = 10 \log_{10} \left| \sum_{n=0}^{N} \frac{p[n]^2}{(p[n] - \widehat{p}[n])^2} \right| \qquad (1)$$

This score is intended to reflect the potential difficulty in correctly identifying the original instrument from the modified sound where $p[n]$ and $\widehat{p}[n]$ are the original and modified sounds, respectively. The resulting difficulty score ranges from 1-25, where 1 corresponds to a high SNR (little change) and 25 represents a low SNR (significant change). The player has the incentive to modify the timbre of the sound as greatly as possible while still maintaining the identity of the original instrument. They will be awarded points based on the difficulty score of their modifications only if a listener can correctly guess the original instrument. This encourages the player to be creative with their timbre adjustments yet still produce recognizable musical sounds.

When a player is finished altering the timbre of a specific instrument, they submit their information, including user ID and modified time and spectral envelopes, to the server

database, which collects all the players' modified sounds. The player can then load another sound from the database to continue with timbre modification.

3.1.2 Tone Listener

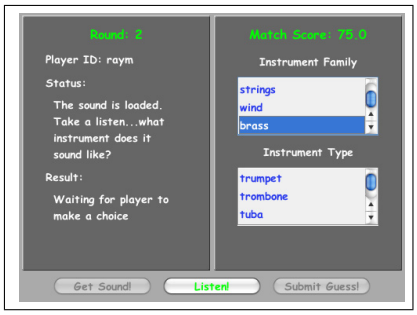

Figure 2. The Tone Listener interface

In the Tone Listener interface, a player is presented with a Java applet that allows them to listen to sounds created by other players in the Tone Bender component. A player's objective is to correctly guess the family and identity of the instrument from the modified sound with which they are presented. The modified sounds are randomly selected from the database and the sounds are re-synthesized in the applet using the amplitude and spectral envelope data. The player is allowed to listen as many times as needed before submitting their guess.

The listening player is allowed to classify the sound among three instrument families: strings, wind, and brass. The player's choice populates another list consisting of individual instruments. After the player has listened to the sound and made a selection for the family and instrument, they submit their guess. The player will receive points only if they correctly guess either the instrument family or the specific instrument. If the player correctly guesses an instrument, they receive a score proportional to the difficulty rating. If the player correctly guesses within the instrument family, they receive half of the potential point value. After each guess, the results, including the user ID, original sound information and the player response are uploaded to a server database containing all players' listening results.

3.1.3 Educational objectives

The Timbre Game is designed to educate players about sound timbre, particularly the importance of time- and frequency-domain envelopes. The interface does not require any previous background in engineering or music, and the ability to listen to timbral changes in real-time encourages the user to learn by experimentation. Additionally, the simple scoring method is designed to provide players with an indication of the degree of change imparted upon the sound, while concealing technical details, such as SNR.

3.2 Cocktail Party Game

The Cocktail Party Game is a web-based activity designed to collect data from listeners on how source locations and acoustic spaces affect identification of a speaker and the intelligibility of speech. It also provides a method of examining the perception of complex timbres, such as the dynamically varying sounds of voices. This game consists of two components: room creation and listening room simulators. The room creation component introduces the concepts of the cocktail party problem and illustrates the effects of reverberation and interfering sounds. The listening room component evaluates the ability of a listener to detect a known person's voice within a room mixture. The two components are described in further detail in the sections that follow.

3.2.1 Room Creation

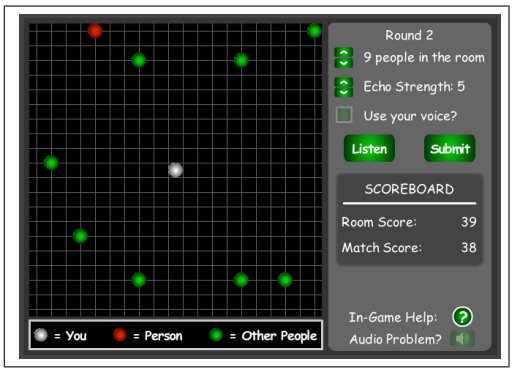

Figure 3. The Room Creation interface

In this component of the game, the player simulates a "cocktail party" situation by positioning multiple talkers, including the target person of interest, in a reverberant room, thus making it more difficult to hear the voice of the target speaker. The goal is to create a situation where the speaker of interest is obscured, but still identifiable, and more points potentially will be awarded to the player based on the "degree of difficulty" of the designed room. Initially, the game displays a room (20' x 20' x 10') containing two people: the listener and the person of interest, represented as white and red circles, respectively. The player has the option to add or remove people from the room, change the strength of the room reverberation, and alter the position of the people in the room. Audio for each of the speakers in the room is randomly drawn from the well-known TIMIT speech database [10]. The browser-based user interface communicates with the server to download the relevant speech files.

A room's potential score is based on the resulting SINR, treating the target voice as the signal and the others as interferers. These points are only added to the player's score if another player correctly determines if the speaker is in the room.

3.2.2 Room Listening

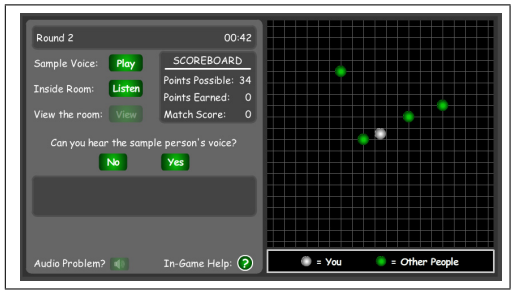

Figure 4. The Listening Room interface

In the listening component of the game, the player's goal is simply to determine if the target person's voice is in the mixture of voices in the room. The game communicates with the server to obtain parameters and sound sources for a previously created room. The game randomly determines whether or not the target voice will be present in the room. The player then listens to the mixed room audio with the option to graphically view the configuration of the people in the room. The room audio is generated in exactly the same manner as in the room creation component. Then the player decides whether or not the sample person's voice is in the room.

After the player decides if the person of interest is in the room, the response is sent to the server and the player is informed of the correct answer. The points are based on the difficulty of the room as calculated by the SINR and only awarded when the person chooses the correct answer. After the information is submitted, the player continues on to the next round until all rounds are completed for the current match.

3.2.3 Educational Objectives

The Cocktail Party Game is designed to educate and inform students about the cocktail party effect and basic room acoustics. Like the Timbre Game, the controls are designed to be simple and minimal so that players can easily experiment with sound source locations and listen to the results. The graphical layout of the game represents actual acoustic space so that players can visually correlate speaker positions with the resulting sound. The activity is intended for players without any specific training in engineering or music, which broadens its appeal.

3.3 Technical implementation details

Both the Timbre Game and Cocktail Party game employ a client-server architecture that allows distributed online game play, so that players of both games may be widely separated. The data for both activities is stored and served from a web server using a MySQL database. PHP web server scripts on the server respond to client queries with XML formatted responses from the database, such as sound and room parameters and audio file URLs.

The Timbre Game is a single Java applet, containing the interface, sound processing, and server communication code. The overall client-server architecture for the Timbre Game shown in Figure 5.

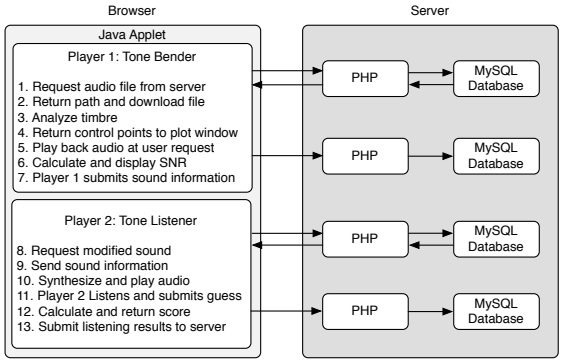

Figure 5. Diagram of Timbre Game

The user interface for the components of the Cocktail Party Game is implemented in Adobe Flash, which has rather limited audio processing and computational capabilities. Therefore, it was necessary to also use a Java helper applet for audio playback flexibility and its concurrent computation capabilities using threading, which is generally sufficient to handle the computation required for room acoustics simulation. Calculation and playback of the room audio in both components of the game is initiated via a call in Flash that sends the room configuration and speaker audio file paths to the Java applet via a JavaScript bridge. The Flash application also communicates with the server to obtain game parameters and audio file paths using Asynchronous JavaScript and XML (AJAX) calls.

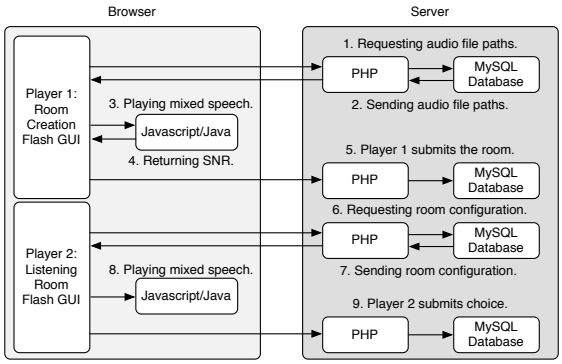

Figure 6. Diagram of Cocktail Party Game

The generation of the room audio for listening is a multi-

step process, requiring multiple components and conversions. First, the room impulse response for each source speaker location is determined based on the person's position in the room using a Java implementation of the well-known room image model [11]. Next, the resulting room impulse response is convolved with the respective speech audio using fast block convolution via the FFT in blocks of approximately 8192 samples ($\sim$ 0.5 seconds at 16 kHz sampling). An efficient Java FFT library was used to optimize the calculation speed by employing concurrent threading. In the final step, the convolved, reverberant speech from each source is combined to obtain the overall room audio for the current configuration. This audio is then played through the Java applet in the client's browser. The overall architecture of the Cocktail Party Game is given in Figure 6.

4 ACTIVITY EVALUATIONS

Evaluations of both activities were performed on a population of 56 eighth grade students attending a magnet school specializing in music performance. Activity sessions focused on the Room Simulation Game and the Timbre Game on separate days. The students were divided into groups of approximately 10 students for each of six sessions lasting 40 minutes per day. The students were switched from the creative component of the game to the listening component midway through the session to have an opportunity to both create and objectively listen to the sounds. The students played the games alone using headphones to avoid confusion in the sound creation and listening processes.

Prior to playing each component, the students were given a 2-3 minute demonstration covering the game objectives, instructions, and user controls. The students were given the opportunity to ask questions throughout the sessions.

4.1 Quantitative results

4.1.1 Timbre Game

Figure 7 provides the results of 800 listening trials from the Tone Listener game: percentage of correct identification of the instrument and instrument family vs. varying SNR levels. The plots demonstrate a very slight upward trend in percentage of correct detection with increasing SNR, with the percentage of correct family detection being greater than correct instrument detection across all SNR values. This result is expected since, in general, it is easier for a listener to identify an instrument's family. The wide variation in performance, however, is likely due to the difference between SNR as a measure of sound quality and actual perceived sound quality. It should be noted that the majority of sounds listened to were created with an SNR value under 20 dB due to players seeking to earn a high modification score by creating more difficult instrument sounds.

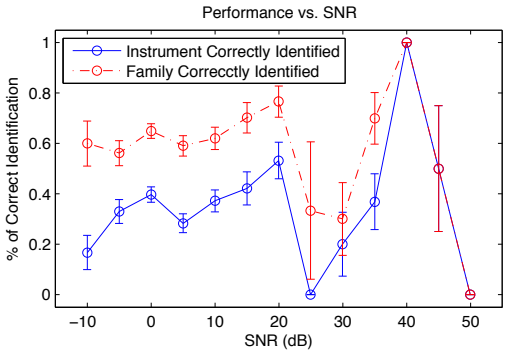

Figure 7. Timbre Game Results

4.1.2 Cocktail Party Game

The results of 817 unique listening tests were analyzed and the results are presented in Figure 8. As the figure shows, the percentage of correct detection generally increased with the SINR. This result is somewhat expected since rooms with higher SINR indicate situations where the greater energy of the target listener should make them easier to identify. This upward trend, however, was not strictly monotonic, indicating that factors other than SINR affect overall performance. The confusion matrix indicates that false negatives were far more likely than false positives, an interesting result that warrants further study.

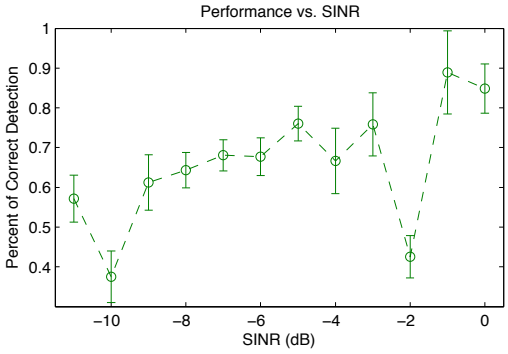

Figure 8. Cocktail Party Game Results

		Correct Answer	
		In Room	Not In Room
Player Guess	In Room	229	84
	Not In Room	215	289

Table 1. Cocktail Party Evaluation confusion matrix

4.2 Qualitative Observations

The Tone Listener interface had the broadest appeal among the students and provided the most instant gratification. This was likely due to the simple objective of the activity, which only required them to listen to sounds and guess the original instrument producing them. Additionally, the instant feedback and scoring added a distinct competitive aspect that encouraged the students to keep playing and comparing scores with each other. In the case of Tone Bender, student reactions to the game objective were mixed. Some students appeared intimidated by or uninterested in the visual representations of instrument timbre. This behavior was evident in students repeatedly asking the demonstrators (graduate students) to explain the game or simply not participating. The students who were more engaged with the activity, however, attempted to modify many different sounds without assistance.

Similarly, the room creation component of the Cocktail Party Game raised more questions and requests for clarification from the students than the listening component. This was expected since the game requires students to be creative and to achieve some understanding of the task in order to successfully design a challenging room. The activity could be improved by altering the audio processing chain so that the room audio responds in real-time to game parameter changes (position and number of sources, etc.). Then the player would receive instant audio feedback, reducing the time to iterate a particular room design. The lack of information provided to room creators regarding the performance of their rooms when listened to by other players was also frustrating and reduced one of the motivating competitive aspects of the game. Overall, the room creation component may need to be simplified in order for middle school students to better understand the objectives of the activity.

5 FUTURE WORK

The websites for both activities will eventually be made publicly available through the web. Another improvement we believe will enhance the activities is to allow players to record their own sounds for immediate use in the games. For examples, an instrumentalist could record their own instrument to be used in the Timbre Game, and players could record their own voices for use in the Cocktail Party game. This feature would enable continuous expansion of the sound databases, keeping the game fresh for frequent users.

A relatively straightforward extension of the Cocktail Party Game would be to extend the sound sources to musical instruments, providing a method of examining the perception of individual instruments within mixtures. We are also investigating the utility of time limits for different phases of the game, in order to keep the activity moving and to increase competition. We particularly wish to pursue a detailed analysis of acquired performance data for cases that deviate from anticipated "difficulty" in terms of SNR. We plan to investigate other acoustic factors affecting listening performance as well as other metrics that may be better correlated to perceptual task performance than SNR.

6 ACKNOWLEDGEMENTS

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ARMONIQUE: EXPERIMENTS IN CONTENT-BASED SIMILARITY RETRIEVAL USING POWER-LAW MELODIC AND TIMBRE METRICS

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ABSTRACT

This paper presents results from an on-going MIR study utilizing hundreds of melodic and timbre features based on power laws for content-based similarity retrieval. These metrics are incorporated into a music search engine prototype, called Armonique. This prototype is used with a corpus of 9153 songs encoded in both MIDI and MP3 to identify pieces similar to and dissimilar from selected songs. The MIDI format is used to extract various power-law features measuring proportions of music-theoretic and other attributes, such as pitch, duration, melodic intervals, and chords. The MP3 format is used to extract power-law features measuring proportions within FFT power spectra related to timbre. Several assessment experiments have been conducted to evaluate the effectiveness of the similarity model. The results suggest that power-law metrics are very promising for content-based music querying and retrieval, as they seem to correlate with aspects of human emotion and aesthetics.

1. INTRODUCTION

We present results from an on-going project in music information retrieval, psychology of music, and computer science. Our research explores power-law metrics for music information retrieval.

Power laws are statistical models of proportions exhibited by various natural and artificial phenomena [13]. They are related to measures of self-similarity and fractal dimension, and as such they are increasingly being used for data mining applications involving real data sets, such as web traffic, economic data, and images [5]. Power laws have been connected with emotion and aesthetics through various studies and experiments [8, 9, 12, 14, 16, 18].

We discuss a music search engine prototype, called *Armonique*, which utilizes power-law metrics to capture both melodic and timbre features of music. In terms of input, the user selects a music piece. The engine searches for pieces similar to the input by comparing power-law proportions through the database of songs. We provide an on-line demo of the system involving a corpus of 9153 pieces for various genres, including baroque, classical, romantic, impressionist, modern, jazz, country, and rock among others. This corpus was originally encoded in MIDI, which facilitated extraction of melodic features. It was then converted to MP3 for the purpose of extracting timbre features.

In terms of assessment, we conducted an experiment by measuring human emotional and physiological responses to the music chosen by the search engine. Analyses of data indicate that people do indeed respond differently to pieces identified by the search engine as similar to the participant-chosen piece, than to pieces identified by the engine as different. For similar pieces, the participants' emotion while listening to the music is more pleasant, their mood after listening is more pleasant; they report liking these pieces more, and they report them to be more similar to their own chosen piece. These results support the potential of using power-law metrics for music information retrieval.

Section 2 presents relevant background research. Sections 3 and 4 describe our power-law metrics for melodic and timbre features, respectively. Sections 5 and 6 discuss the music search engine prototype, and its evaluation with human subjects. Finally, section 7 presents closing remarks and directions for future research.

2. BACKGROUND

Tzanetakis et al. [15] performed genre classifications using audio signal features. They performed FFT analysis on the signal and calculated various dimensions based on the frequency magnitudes. Also, they extracted rhythm features through wavelet transforms. They reported classification success rates of 62% using six genres (classical, country, disco, hiphop, jazz and rock) and 76% using four classical genres.

Aucouturier and Packet [1] report the most typical audio similarity technique is timbre via spectral analysis using Mel-frequency cepstral coefficients (MFCCs). Their goal was to improve on the overall performance of timbre similarity by varying parameters associated with these techniques (e.g. sample rate, frame size, number of MFCCs used, etc.) They report that there is a "glass ceiling" for timbre similarity that prevents any major improvements in performance via this technique. Subsequent research on this seems to either be a verification of this ceiling or an attempt to pass it using additional similarity dimensions (e.g., [10]).

Lidy et al. [6] discuss a genre classification experiment utilizing both timbre and melodic features. They utilize the typical timbre measures obtained through spectral analysis. They also employ a music transcription system they developed to generate MIDI representations of audio music files. From these files, they extract 37 different features, involving attributes of note pitches, durations and non-diatonic notes. The combined timbre and melodic features are used to conduct genre classification experiments. They report classification accuracies with combined feature sets ranging from 76.8% to 90.4% using standard benchmarks (e.g., ISMIR 2004 audio data).

Cano et al. [4] report on a music recommendation system called *MusicSurfer*. The primary dimensions of similarity used are timbre, tempo and rhythm patterns. Using a corpus of 273,751 songs from 11,257 artists their system achieves an artist identification rate of 24%. On the ISMIR 2004 author identification set they report a success rate of 60%, twice as high as that of the next best systems. MusicSurfer has a comprehensive user interface that allows users to find songs based on artist, genre, and similarity among other characteristics, and allows selection of different types of similarity. However, it does not include melodic features.

2.1. Power Laws and Music Analysis

Our music recommender system prototype employs both melodic and timbre features based on power-laws. Although power laws, fractal dimension, and other self-similarity features have been used extensively in information retrieval (e.g., [5]), to the best of our knowledge, there are no studies of content-based music recommendation systems utilizing power law similarity metrics.

A power law denotes a relationship between two variables where one is proportional to a power of the other. One of the most well-known power laws is *Zipf's law*:

$$P(f) \sim 1 / f^{n} \quad (1)$$

where $P(f)$ denotes the probability of an event of rank f, and n is close to 1. Zipf's law is named after George Kingsley Zipf, the Harvard linguist who documented and studied natural and social phenomena exhibiting such relationships [18]. The generalized form is:

$$P(f) \sim a / f^{b} \quad (2)$$

where a and b are real constants. This generalized form is known as the Zipf-Mandelbrot law, after Benoit Mandelbrot.

Numerous empirical studies report that music exhibits power laws across timbre and melodic attributes (e.g., [8, 16, 18]). In particular, Voss and Clarke [16] demonstrate that music audio properties (e.g. loudness and pitch fluctuation) exhibit power law relationships. Using 12 hours worth of radio recordings, they show that power fluctuations in music follow a *1/f* distribution.

Manaris et al. [8] report various classification experiments with melodic features based on power laws. These studies include composer identification with 93.6% to 95% accuracy. They also report an experiment using emotional responses from humans. Using a corpus of 210 music excerpts in a 12-fold cross-validation study, artificial neural networks (ANNs) achieved an average success rate of 97.22% in predicting (within one standard deviation) human emotional responses to those pieces.

3. MELODIC FEATURE EXTRACTION

As mentioned earlier, we employ hundreds of power-law metrics that calculate statistical proportions of music-theoretic and other attributes of pieces.

3.1. Melodic Metrics

We have defined 14 power-law metrics related to proportion of pitch, chromatic tone, duration, distance between repeated notes, distance between repeated durations, melodic and harmonic intervals, melodic and harmonic consonance, melodic and harmonic bigrams, chords, and rests [9]. Each metric calculates the *rank-frequency* distribution of the attribute in question, and returns two values:

- the slope of the trendline, b (see equation 2), of the rank-frequency distribution; and
- the strength of the linear relation, r^2.

We also calculate *higher-order* power law metrics. For each regular metric we construct an arbitrary number of higher-order metrics (e.g., the difference of two pitches, the difference of two differences, and so on), an approach similar to the notion of derivative in mathematics.

Finally, we also capture the difference of an attribute value (e.g., note duration) from the local average. Local variability, *locVar[i],* for the i^{th} value is

$$locVar[i] = abs(vals[i] - avg(vals, i)) / avg(vals, i) \quad (3)$$

where *vals* is the list of values, *abs* is the absolute value, and *avg(vals, i)* is the local average of the last, say, 5 values. We compute a local variability metric for each of the above metrics (i.e., 14 regular metrics x the number of higher-order metrics we decide to include).

Collectively, these metrics measure hierarchical aspects of music. Pieces without hierarchical structure (e.g., aleatory music) have significantly different measurements than pieces with hierarchical structure and long-term dependencies (e.g., fugues).

3.2. Evaluation

Evaluating content-based music features through classification tasks based on objective descriptors, such as artist or genre, is recognized as a simple alternative to listening tests for approximating the value of such features for similarity prediction [7, 11].

We have conducted several genre classification experiments using our set of power-law melodic metrics to extract features from music pieces. Our corpus

consisted of 1236 MIDI-encoded music pieces from the Classical Music Archives (www.classicalarchives.com). These pieces were subdivided into 9 different musical genres (listed here by timeline): Medieval (57 pieces), Renaissance (150 pieces), Baroque (160 pieces), Classical (153 pieces), Romantic (91 pieces), Modern (127 pieces), Jazz (118 pieces), Country (109 pieces), and Rock (271 pieces).

First, we carried out a 10-fold cross validation experiment training an ANN to classify the corpus into the 9 different genres. For this, we used the Multilayer Perceptron implementation of the Weka machine learning environment. Each piece was represented by a vector of 156 features computed through our melodic metrics. The ANN achieved a success rate of 71.52%.

Figure 1 shows the resulting confusion matrix. It is clear that most classification errors occurred between genres adjacent in timeline. For example, most Renaissance pieces misclassified (32/43) were either falsely assigned to the Medieval (7) or Baroque (25) period. Most misclassified Baroque pieces (40/64) were incorrectly classified as Renaissance (23) or Classical (17), and so on. This is not surprising, since there is considerable overlap in style between adjacent genres.

To verify this interpretation, we ran several binary classification experiments. In each experiment, we divided the corpus into two classes: *class 1* consisting of a particular genre (e.g., Baroque), and *class 2* consisting of all other genres non-adjacent in timeline (e.g., all other genres minus Renaissance and Classical). For Jazz, as well as Rock, the other class included all other genres.

Table 1 shows the classification accuracies for all of these experiments. Since the two output classes in each experiment were unbalanced, the ANN accuracy rates should be compared to a majority-class classifier.

4. TIMBRE FEATURE EXTRACTION

We calculate a *base audio metric* employing spectral analysis through FFT. This metric is then expanded through the use of higher-order calculations and variations of window size and sampling rate. Since we are interested in power-law distributions within the human hearing range, assuming CD-quality sampling rate (44,1KHz), we use window sizes up to 1-sec. Interestingly, given our technique, the upper frequencies in this range do not appear to be as important for calculating timbre similarity; the most important frequencies appear to be from 1kHz to 11kHz.

For each of these windows we compute the power spectrum per window and then average the results, across frequencies. We then extract various power-law proportions from this average power spectrum. Again, each power-law proportion is captured as a pair of slope and r^2 values.

a	*b*	*c*	*d*	*e*	*f*	*g*	*h*	*i*	*<-- classified as*
92	17	1	2	4	4	23	8	9	| *a = baroque*
20	107	0	1	2	5	5	4	9	| *b = classical*
1	1	96	0	0	0	0	9	2	| *c = country*
0	3	1	96	0	2	1	13	2	| *d = jazz*
8	3	0	0	28	0	15	3	0	| *e = medieval*
9	9	1	0	0	78	2	10	18	| *f = modern*
25	8	1	0	7	0	107	1	1	| *g = renaissance*
1	2	7	11	0	4	1	242	3	| *h = rock*
9	12	3	3	1	19	1	5	38	| *i = romantic*

Figure 1. Confusion matrix from 9 genre multi-classification ANN experiment.

Class 1	Class 2	ANN Accuracy	Majority Classifier
Baroque	Non-Adjacent	91.29 %	82.85 %
Classical	Non-Adjacent	92.08 %	84.46 %
Rock	Non-Rock	92.88 %	78.07 %
Jazz	Non-Rock	96.66 %	90.45 %

Table 1. ANN success rates for binary classification experiments.

We have explored various ways to calculate *higher-order* quantities involving both signal amplitudes and power spectrum (i.e., frequency) magnitudes. Again, the idea of a higher-order is similar to the use of derivatives in mathematics where one measures the rate of change of a function at a given point. Through this approach, we have created various derivative metrics, involving raw signal amplitude change, frequency change within a window, and frequency change across windows.

To further explore the proportions present in the audio signal, we vary the window size and the sampling rate. This allows us to get measurements from multiple "views" or different levels of granularity of the signal. For each new combination of window size and sampling rate, we recompute the above metrics, thus getting another pair of slope and r^2 values. Overall, we have defined a total of 234 audio features.

4.1. Evaluation

To evaluate these timbre metrics, we conducted a classification experiment involving a corpus of 1128 MP3 files containing an equal number of classical and non-classical pieces.

We carried out a 10-fold cross-validation, binary ANN classification experiment using a total of 234 audio features. For comparison, each classification was repeated using randomly assigned classes. The ANN achieved a success rate of 95.92%. The control success rate was 47.61%. We are in the process of running additional classification experiments to further evaluate our timbre metrics (e.g., ISMIR 2004 audio data).

5. A MUSIC SEARCH ENGINE PROTOTYPE

Musicologists consider melody and timbre to be independent/complementary aesthetic dimensions (among others, such as rhythm, tempo, and mode). We have developed a music search-engine prototype, called Armonique, which combines melodic and timbre metrics to calculate sets of similar songs to a song selected by the user. For comparison, we also generate a set of dissimilar songs. A demo of this prototype is available at http://www.armonique.org.[1]

The corpus used for this demo consists of 9153 songs from the Classical Music Archives (CMA), extended with pieces from other genres such as jazz, country, and rock. These pieces originated as MIDI and were converted for the purpose of timbre feature extraction (and playback) to the MP3 format. As far as the search engine is concerned, each music piece is represented as a vector of hundreds of power-law slope and r^2 values derived from our metrics. As input, the engine is presented with a single music piece. The engine searches the corpus for pieces similar to the input, by computing the mean squared error (MSE) of the vectors (relative to the input). The pieces with the lowest MSE are returned as best matches.

We have experimented with various selection algorithms. The selection algorithm used in this demo, first identifies 200 similar songs based on melody, using an MSE calculation across all melodic metrics. Then, from this set, it identifies the 10 most similar songs based on timbre, again, using an MSE calculation across all timbre metrics. Each song is associated with two gauges providing a rough estimate of the similarity across the two dimensions, namely melody and timbre. It is our intention to explore more similarity dimensions, such as rhythm, tempo, and mode (major, minor, etc.).

6. EVALUATION EXPERIMENT

We conducted an experiment with human subjects to evaluate the effectiveness of the proposed similarity model. This experiment evaluated only the melodic metrics of Armonique, since the timbre metrics had not been fully incorporated at the time.[2]

6.1. Participants

Twenty-one undergraduate students from Bethel College participated in this study. The participants consisted of 11 males and 10 females, with a mean age of 19.52 years and a range of 5 years. They had participated in high school or college music ensembles for a mean of 3.52 years with a standard deviation of 2.21, and had received private music lessons for a mean of 6.07 years with a standard deviation of 4.37.

[1] Due to copyright restrictions, some functionality is password-protected.
[2] We are planning a similar assessment experiment involving both melodic and timbre metrics.

6.2. Design

A single repeated-measures variable was investigated. This variable consisted of the seven different excerpts, one of them participant-chosen (hereafter referred to as the original piece), three of them computer-selected to be similar to the participant-chosen piece, and three selected to be different (see next section). Of primary interest was the comparison of responses to the original with those to the three similar pieces, of the original to the three different pieces, and of the three similar to the three different pieces.

6.3. Data Set

At the time they were recruited, participants were asked to list in order of preference three classical music compositions that they enjoyed. The most preferred of these three compositions that was available in the CMA corpus was chosen for each participant. The search engine was then employed to select three similar and three different pieces. Similarities were based upon the first two minutes of each composition. This corpus is available at http://www.armonique.org/melodic-search.html .

6.4. Procedure

The resulting seven two-minute excerpts (original plus six computer-selected pieces), different for each participant, were employed in a listening session in which participant ratings of the music were obtained during and after the music, and psychophysiological responses were monitored (i.e., skin conductance, heart rate, corrugator supercilii electromygraphic recording, respiration rate, and 32 channels of electroencephalographic recording). The physiological data are not reported here. In some instances the entire piece was shorter than two minutes. A different random order of excerpts was employed for each participant.

Ratings of mood were obtained immediately prior to the session using the Self-Assessment Manikin [3] represented as two sliders with a 1-9 scale, one for pleasantness and one for activation, on the front panel of a LabVIEW (National Instruments, Austin, TX) virtual instrument. Physiological recording baseline periods of one minute were included prior to each excerpt. By means of a separate practice excerpt, participants were carefully instructed to report their feelings during the music by using a mouse to move a cursor on a two-dimensional emotion space [2]. This space was displayed on the front panel of the LabVIEW instrument, which also played the music file and recorded x-y coordinates of the cursor position once per second. After each excerpt participants again rated their mood with the Self-Assessment Manikin, and then rated their liking of the previous piece using another slider on the front panel with a 1-9 scale ranging from "Dislike very much" to "Like very much."

At the conclusion of the listening session, participants rated the similarity of each of the six computer-selected

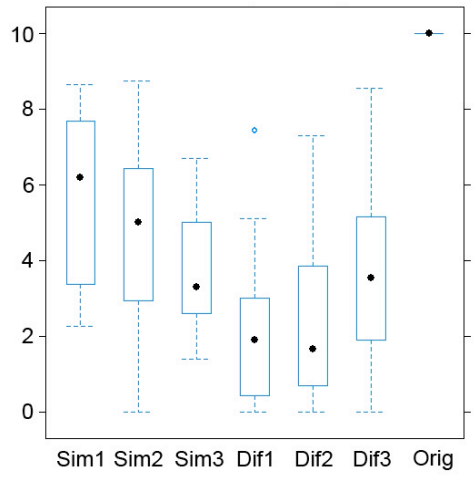

Figure 2. Boxplots of similarity ratings across all subjects for the three similar songs recommended by the search engine and the three different songs.

excerpts to the original; these ratings were accomplished on a LabVIEW virtual instrument with six sliders and 0-10 scales ranging from "Not at all" to "Extremely."

6.5. Data Analysis

Data from the ratings during the music were averaged over time, yielding an average pleasantness and activation measure for each excerpt. A procedural error resulted in loss of data for two participants on these measures. Data from each behavioral measure were subjected to planned contrasts based upon a repeated-measures analysis of variance (SYSTAT software, SYSTAT, Inc., San Jose, CA). These contrasts compared the original song with each of the other two categories of music, and each of those categories with each other.

6.6. Results

The results for similarity ratings are shown in Figure 2. A contrast between the three similar and three different pieces indicated that the similar pieces were indeed judged to be more similar than were the different ones ($F(1, 20) = 20.98$, $p < 0.001$). Interestingly, the similar songs recommended by the search engine were ordered by humans the same way as by the search engine.[1]

In terms of the average ratings for pleasantness recorded during the music, the contrast between the similar pieces and the original was significant ($F(1, 18) = 5.85$, $p = 0.026$), while that between the different pieces and the original showed an even more marked difference ($F(1, 18) = 11.17$, $p = 0.004$). The contrast between similar and different pieces approached significance ($F(1, 18) = 3.04$, $p = 0.098$). No significant differences were found for these contrasts on the average activation measure.

In terms of the ratings for pleasantness recorded after the music, the contrast between the similar pieces and the original was not significant ($F(1, 20) = 1.21$, $p = 0.285$), while that between the different pieces and the original was significant ($F(1, 20) = 7.64$, $p = 0.012$). The contrast between similar and different pieces again approached significance ($F(1, 20) = 4.16$, $p = 0.055$). No significant differences were found for these contrasts on the activation measure recorded after the music.

Finally, in terms of the ratings for liking recorded after the music, the contrast between the similar pieces and the original showed a clear difference ($F(1, 20) = 20.31$, $p < 0.001$), as did that between the different pieces and the original ($F(1, 20) = 42.09$, $p < 0.001$). The contrast between similar and different pieces was not significant ($p > 0.2$).

7. CONCLUSION AND FUTURE WORK

These assessment results involving human subjects suggest that the model under development captures significant aesthetic similarities in music, evidenced through measurements of human emotional and, perhaps, physiological responses to retrieved music pieces. At the same time they indicate that there remains an important difference in affective responses – greater liking of the piece chosen by the person. Thus, these data provide new insights into the relationship of positive emotion and liking.

It is well documented experimentally that liking increases with exposure to music up to some moderate number of exposures and then decreases as the number of exposures becomes very large [17]. It may be that the pieces chosen by our participants are near that optimum number of exposures whereas the aesthetically similar pieces are insufficiently (or perhaps excessively) familiar to them. Thus, different degrees of familiarity may account for some of the liking differences that were found. It may be desirable in future studies to obtain some independent measure of prior exposure to the different excerpts in order to assess the contribution of this factor.

We plan to conduct additional evaluation experiments involving humans utilizing both melodic and timbre metrics. We also plan to explore different selection algorithms, and give the user more control via the user interface, in terms of selection criteria.

Finally, it is difficult to obtain high-quality MIDI encodings of songs (such as the CMA corpus). However, as Lidy et al. demonstrate, even if the MIDI transcriptions are not perfect, the combined (MIDI + audio metrics) approach may still have more to offer than timbre-based-only (or melodic-only) approaches [6].

This paper presented results from an on-going MIR study utilizing hundreds of melodic and timbre metrics

[1] It should be noted that the different songs used in this corpus were not the most dissimilar songs. These selections were used to avoid a cluster of most dissimilar songs, which was the same for most user inputs. However, they also somewhat reduced the distance between similar and dissimilar songs.

based on power laws. Experimental results suggest that power-law metrics are very promising for content-based music querying and retrieval, as they seem to correlate with aspects of human emotion and aesthetics. Power-law feature extraction and classification may lead to innovative technological applications for information retrieval, knowledge discovery, and navigation in digital libraries and the Internet.

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COLLECTIVE ANNOTATION OF MUSIC FROM MULTIPLE SEMANTIC CATEGORIES

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ABSTRACT

Music semantic annotation aims to automatically annotate a music signal with a set of semantic labels (words or tags). Existing methods on music semantic annotation usually take it as a multi-label binary classification problem, and model each semantic label individually while ignoring their relationships. However, there are usually strong correlations between some labels. Intuitively, investigating this correlation can be helpful to improve the overall annotation performance. In this paper, we report our attempts to collective music semantic annotation, which not only builds a model for each semantic label, but also builds models for the pairs of labels that have significant correlations. Two methods are exploited in this paper, one based on a generative model (Gaussian Mixture Model), and another based on a discriminative model (Conditional Random Field). Experiments show slight but consistent improvement in terms of precision and recall, compared with the individual-label modeling methods.

1 INTRODUCTION

Semantic annotation of music signals have become an important direction in music information retrieval. With music annotation approaches, a music signal is associated with a set of semantic labels (text, words), which is a more compact and efficient representation than the raw audio or low level features. It can also potentially facilitate a number of music applications, such as music retrieval and recommendation, since it is more natural for a user to describe a song by semantic words, and it is more flexible to measure music similarities with vectors of semantic labels.

Several methods have been proposed for automatic music semantic annotation, which basically can be classified into two categories: *non-parametric* and *parametric*. Non-parametric methods model the text-audio relations implicitly. For example, Slaney [10] created separate hierarchical models in the acoustic and text spaces and then linked the two spaces for annotation and retrieval. Cano and Koppenberger [2] proposed an approach to predict the semantic words based on nearest neighbor classification. On the other hand, parametric methods explicitly model the text-audio relations. For instance, Whitman *et al.* [14, 13] trained a one-versus-all discriminative model (a regularized least-square classifier or a support vector machine) for each word, based on which the audio frames were classified. Turnbull *et al.* [11] built a generative model for each semantic word, and calculated a multinomial distribution over the word vocabulary for each song. Eck *et al.* [3] used AdaBoost to predict the strength of the occurrence of each social tag (a word) on a large audio data set.

The methods abovementioned achieve good results by modeling the text-audio relations, however, they share two drawbacks. First, there lacks of a taxonomy to organize the semantic labels. Music has a number of important aspects affecting music perception and music similarity, such as genre, instrumentation, emotion, and tempo, etc. The classification or detection of these aspects, were also investigated in many previous methods [12, 4, 8, 9]. Semantic labels used for music annotation can be also naturally divided into groups corresponding to these aspects. However, although most of the annotation methods use a rather large semantic vocabulary that covers almost all the aspects, the words are not structurally organized. One consequence is that they cannot make sure that a song is annotated from all the aspects. For example, Turnbull *et al.* [11] calculated the posterior probability for each word in the vocabulary given a song, and selected the A (a constant) words with the largest posterior probability to annotate the song. Thus, suppose for some songs, the posterior probabilities of some words describing genre are larger than those of all the words describing instrumentation, this may cause the words describing instrumentation being absent in the annotation.

Second, in the previous methods, semantic labels are modeled individually, that is, the methods only build text-audio

This wok was performed when the first author was a visiting student in Microsoft Research Asia.

relations, but ignore the text-text relations between two labels. However, some labels do have strong correlations, and this information can be investigated to improve annotation schemes. For example, "hard rock" and "electronic guitar" tend to co-occur in the annotations of a song, while "happy" and "minor key", "fast tempo" and "gloomy" tend rarely to co-occur. Using the text-text relation information, strong evidence for the occurrence of "hard rock" may help to predict the presence of "electronic guitar". On the other hand, conflicts in the annotated labels such as the co-occurrence of "fast tempo" and "gloomy", which may happen using the individually annotation methods, could be mostly avoided by employing the text-text correlation.

To address the two issues above, this paper divides the semantic vocabulary into a number of categories, and proposes two collective annotation methods by exploiting the correlations within label pairs. Specifically, 50 web-parsed semantic labels are used to form the vocabulary, and are divided into 10 categories, each of which describes an aspect of music attributes, including genre, instrumentation, texture, vocal, arousal, affectivity, tempo, rhythm, tonality and production process. We also pose the restriction that the obtained annotation should contain labels from all the categories. In order to estimate the text-text relations, the normalized mutual information (NormMI) between the labels in the vocabulary is calculated. The label pairs whose NormMI values are larger than a threshold are selected to be modeled. Two methods are then exploited to integrate correlation modeling: one is a generative method, in which each selected label pair is modeled by a Gaussian Mixture Model (GMM); the other is a discriminative method, which is based on Conditional Random Field (CRF).

The rest of the paper is organized as follows: Section 2 describes the semantic vocabulary and the selection process of important word pairs. Section 3 describes audio feature extraction. The two proposed annotation methods are presented in Section 4. Section 5 presents the experimental results, and Section 6 concludes this paper.

2 SEMANTIC VOCABULARY

A vocabulary lists all the labels that can be used for semantic annotation. Currently there is not a standard vocabulary, and most researchers build their own vocabularies. Cano and Koppenberger [2] used the taxonomy provided by WordNet [1] . Whitman and Rifkin [14] extracted about 700 words from web documents associated with artists. Turnbull *et al.* [11] extracted 135 musically relevant words spanning six semantic categories from song reviews. These vocabularies are usually large enough to cover all the aspects of music. However, due to the large vocabulary, it is usually hard to avoid preference over words when acquiring the ground truth annotations.

[1] http://wordnet.princeton.edu/

Category	Words	Num
Genre	Blues, Country, Electronica, Folk, Funk, Gospel, HardRock, Jazz, Pop, Punk, Rap, R&b, Rock-roll, SoftRock	1-2
Instrument	Acoustic guitar, Acoustic piano, Bass, Drum, Electric guitar, Electric piano, Harmonica, Horn, Organ, Percussion, Sax, String	1-5
Texture	Acoustic, Electric, Synthetic	1-2
Vocal	Group, Male, Female, No	1-2
Affective	Positive, Neutral, Negative	1
Arousal	Strong, Middle, Weak	1
Rhythm	Strong, Middle, Weak	1
Tempo	Fast, Moderato, Slow	1
Tonality	Major, Mixed, Minor	1
Production	Studio, Live	1

Table 1. The vocabulary contains 50 quantized labels spanning 10 semantic categories. Each song is annotated using labels from all the categories with a number limitation.

In this paper, we build a simplified (but still general) vocabulary from a list of web-parsed musically relevant words. 50 commonly used labels are manually selected and quantized, covering 10 semantic categories (aspects) to describe characteristics of music signals. Table 1 lists the vocabulary. Using this vocabulary, each song will be annotated by labels from each category with label number limitations. This solves the problem that the annotations of a music signal are missing in some music aspects when the vocabulary is not organized as categories. It is noticed that multiple labels can be selected from the categories of genre, instrument, texture and vocal, while the labels within the other categories are exclusive with each other.

The same as existing methods, each label can be viewed as a binary variable in modeling and annotation. As mentioned previously, some labels have strong relations: *positive* or *negative* correlations. For example, the labels within some categories (e.g. Rhythm and Tempo) have negative correlations and are exclusive with each other. Moreover, some labels from different categories may have positive or negative correlations. For example, "Genre.HardRock" and "Arousal.Strong" tend to co-occur, while "Tonality.Major" and "Affective.Negative" tend rarely to co-occur.

The normalized mutual information (NormMI) is used to measure the correlations of each label pair (X, Y) as

$$\text{NormMI}(X,Y) = \frac{I(X,Y)}{\min\{H(X), H(Y)\}} \qquad (1)$$

Word pair	NormMI
(Production.Live, Production.Studio)	1.00
(Vocal.Female, Vocal.Male)	0.79
(Tonality.Major, Tonality.Minor)	0.69
(Tempo.Fast, Tempo.Moderato)	0.62
(Rhythm.Middle, Rhythm.Strong)	0.56
(Genre.Electronica, Texture.Synthetic)	0.25
(Arousal.Weak, Rhythm.Weak)	0.24
(Instrument.AcousticGuitar, Texture.Acoustic)	0.23
(Instrument.Drum, Rhythm.Weak)	0.23
(Genre.HardRock, Instrument.ElectricGuitar)	0.19

Table 2. Selected word pairs and their normalized mutual information. The label pairs in the first five rows are from the same semantic category, and those in the last five rows are from different categories.

where $I(X,Y)$ is the mutual information between X and Y

$$I(X,Y) = \sum_{x\in\{+1,-1\}} \sum_{y\in\{+1,-1\}} P(x,y) \log \frac{P(x,y)}{P_X(x)P_Y(y)} \quad (2)$$

and $H(x)$ is the entropy of label X defined by

$$H(X) = -\sum_{x\in\{+1,-1\}} P_X(x) \log P_X(x) \quad (3)$$

Here $+1$ and -1 represents the presence and absence of a label, respectively. The probability $P_X(x)$ and $P_Y(y)$, and $P(X,Y)$ can be estimated from a training set.

NormMI(X,Y) has the following properties:

1. $0 \leq$ NormMI$(X,Y) \leq 1$;
2. NormMI$(X,Y) = 0$ when X and Y is statistically independent;
3. NormMI$(X,X) = 1$.

NormMI considers label correlations only, and is irrelevant to the distributions of individual labels. The larger NormMI is, the stronger the correlation is. In our approach, only the label pairs whose NormMI values are larger than a threshold are selected to be modeled (see in Section 4). Table 2 lists some of the selected pairs and their NormMI values.

3 AUDIO FEATURE EXTRACTION

Each song is divided into frames with 20ms length and 10ms overlap. Tempo and beats are detected, then the song is divided into beat segments. Each segment contains a number of successive frames. A bag of beat-level feature vectors are used to represent a song. Each vector contains two sets of features: timbre features and rhythm features. Beat-level timbre features are the mean and standard deviation of the timbre features extracted in each frame. Rhythm features are extracted from the beat segments.

The reason to use beat-level features is that they are much more compact than the frame-level features to represent a song, and hence result in much lower computational complexity. Besides, the beat-level features cover a long period information, and may represent some high-level music characteristics so that it may be helpful for the annotation task.

3.1 Timbre Features

For each audio frame, three classes of spectral features are calculated. They are 8-order Mel-frequency cepstral coefficients (MFCCs), spectral shape features and spectral contrast features. The spectral shape features, including brightness, bandwidth, rolloff, and spectral flux, are commonly used in genre classification [12]. The spectral contrast features which was originally proposed in [6], are designed to be a complement of MFCCs on the sub-band information, and are shown successful in mood classification [8]. These three classes of features constitute a 47-dimensional vector. Finally, the mean and standard deviation of the frame-level timbre features in each beat segment compose the beat-level timbre feature vector, which is a 94-dimensional vector.

3.2 Rhythm Features

Rhythm is an important aspect of music. In our approach, a 20-second window (with current beat-segment in the middle) is used for rhythm feature extraction. Following Lu *et al.* [8], eight rhythm features are extracted, including average tempo, average onset frequency, rhythm regularity, rhythm contrast, rhythm strength, average drum frequency, drum amplitude and drum confidence.

In the end, a 102-dimensional (timbre plus rhythm) beat-level feature vector for each beat segment is extracted. Then the vectors are normalized to zero mean and unit variance along each dimension. Principle Component Analysis (PCA) is further employed to reduce the dimensionality of the feature vectors to 65, reserving 95% energy.

4 SEMANTIC ANNOTATION

Given a vocabulary $\mathcal{V}$ consisting of $|\mathcal{V}|$ labels (or words) $w_i \in \mathcal{V}$, and a song s represented by a bag of T real-valued feature vectors $\mathcal{X} = \{\mathbf{x}_1, \cdots, \mathbf{x}_T\}$, the goal of semantic annotation is to find a set $\mathcal{W} = \{w_1, \cdots, w_A\}$ of A words describing the song. It is convenient to represent the set $\mathcal{W}$ as an annotation vector $\mathbf{y} = (y_1, ..., y_{|\mathcal{V}|})$. Here y_i is a binary variable, valued 1 or -1 to represent "presence" or "absence" of label w_i. Therefore, a data set $\mathcal{D}$ is a collection of song-annotation pairs $\mathcal{D} = \{(\mathcal{X}_1, \mathbf{y}_1), \cdots, (\mathcal{X}_{|\mathcal{D}|}, \mathbf{y}_{|\mathcal{D}|})\}$.

In general, this annotation problem can be addressed by Maximum A Posterior (MAP), that is, to choose an annotation vector with maximum posterior: $\hat{\mathbf{y}} = \arg\max P(\mathbf{y}|\mathcal{X})$ [11]. In existing methods, the labels are treated independent, so that the posterior probability of the annotation vector can be decomposed into the multiplication of the posterior probability of each label as

$$P(\mathbf{y}|\mathcal{X}) = \prod_{i}^{|\mathcal{V}|} P(y_i|\mathcal{X}) \propto \prod_{i}^{|\mathcal{V}|} p(\mathcal{X}|y_i)P(y_i) \quad (4)$$

If further assume that feature vectors $\mathbf{x}_1, \cdots, \mathbf{x}_T$ in the bag $\mathcal{X}$ are independent, then

$$p(\mathcal{X}|y_i) = \prod_{t}^{T} p(\mathbf{x}_t|y_i) \quad (5)$$

where T is the number of feature vectors in the bag $\mathcal{X}$. The likelihood $p(\mathbf{x}_t|y_i)$ can be estimated using a parametric model such as a GMM from the training data. The prior probability $P(y_i)$ can also be estimated or as usual set to a uniform distribution.

However, as mentioned above, the labels are not independent, and we need to consider their correlations. In the following subsections, two approaches are exploited for correlation modeling: a GMM-based method, and a Conditional Random Field (CRF)-based method.

4.1 The GMM-based method

When the labels are not independent, the joint posterior probability $P(\mathbf{y}|\mathcal{X})$ cannot be decomposed into single label posteriors. Instead, we approximate it using the multiplication of single label posteriors and label-pair posteriors.

$$P(\mathbf{y}|\mathcal{X}) \sim \prod_{i}^{|\mathcal{V}|} P(y_i|\mathcal{X}) \left(\prod_{j}^{|\mathcal{E}|} P(y_{e_j^1}, y_{e_j^2}|\mathcal{X}) \right)^{\alpha} \quad (6)$$

$$\propto \prod_{i}^{|\mathcal{V}|} p(\mathcal{X}|y_i)P(y_i) \left(\prod_{j}^{|\mathcal{E}|} p(\mathcal{X}|y_{e_j^1}, y_{e_j^2})P(y_{e_j^1}, y_{e_j^2}) \right)^{\alpha} \quad (7)$$

where $\mathcal{E}$ is the set of the selected label pairs that have large normalized mutual information; e_j^1 and e_j^2 are the two labels in the pair; α is a the trade off between label posteriors and label pair posteriors. In our experiments it is set to 1 typically.

The likelihood $p(\mathcal{X}|y_i)$ and $p(\mathcal{X}|y_{e_j^1}, y_{e_j^2})$ can be computed based on Eq.(5), assuming the feature vectors within a bag are independent. The likelihood of each feature vector is estimated using a GMM model, where 8 kernels are arbitrarily selected in this paper.

Although the right part of Eq.(6) is not the exact decomposition of $P(\mathbf{y}|\mathcal{X})$, it represents the intuitive idea that the annotation should not only maximize the posterior probabilities of individual words, but should also consider the posteriors of the correlated label pairs.

4.2 The CRF-based method

Conditional Random Field (CRF) was firstly proposed by Lafferty *et al.* [7] to segment and label sequence data such as natural language. It is an undirected graphical model, where the nodes represent the label variables and the edges represent the relations between labels. Compared with Hidden Markov Model (HMM), one advantage of the chain CRF model is that it relaxes the strong independence assumptions. In fact, a general CRF can naturally model arbitrary dependencies between features and labels. Further, Ghamrawi and McCallum [5] proposed two multi-label classification models based on CRF, which directly parameterized correlations among labels and features.

Generally, given a sample $\mathbf{x}$ and its output label vector $\mathbf{y}$, using the multi-label classification CRF models, the posterior probability $p(\mathbf{y}|\mathbf{x})$ can be written as

$$p(\mathbf{y}|\mathbf{x}) = \frac{1}{Z(\mathbf{x})} \exp\left(\sum_k \lambda_k f_k(\mathbf{x}, \mathbf{y}) + \sum_l \mu_l g_l(\mathbf{x}, \mathbf{y}) \right) \quad (8)$$

where $Z(\mathbf{x})$ is the normalizing factor. $f_k(\mathbf{x}, \mathbf{y})$ and $g_l(\mathbf{x}, \mathbf{y})$, are two predefined real-valued functions, corresponding to a node and an edge respectively. They are usually referred to as *features* of the CRF. In principle, any real-valued function of sample $\mathbf{x}$ and label $\mathbf{y}$ can be treated as a feature. For example, it can be the frequency of a phrase in a text document, or one dimension of the beat-level feature in a music signal. λ_k and μ_l are the parameters to be estimated to maximize Eq. (8) using training data, where k and l enumerate the following indexes of features,

$$k \in < r_i, y_j >: 1 \le i \le |R|, 1 \le j \le |\mathcal{Y}| \quad (9)$$

$$l \in < r_i, y_j, y_{j'} >: 1 \le i \le |R|, 1 \le j, j' \le |\mathcal{Y}| \quad (10)$$

where R is a set of music characteristics (we do not call them features in order to avoid confusion with CRF features like f_k and g_l above), and $r_i \in R$; $|\mathcal{Y}|$ is the length of the label vector, and y_j is a label variable. It can be seen that each feature f_k corresponds to a pair consisting of a label and a characteristic, and each feature g_l corresponds to a triplet consisting of a label pair and a characteristic.

Note that k in Eq.(9) enumerates all the nodes, and l in Eq.(10) enumerates all the edges. Therefore, Eq. (8) corresponds to a full connected graph, where $\sum_k \lambda_k f_k(\mathbf{x}, \mathbf{y})$ represents the overall potential of nodes, and $\sum_l \mu_l g_l(\mathbf{x}, \mathbf{y})$ represents the overall potential of edges. However, in practice, not all the label pairs have close relations, and we only

%	Individual GMM	Collective GMM	Individual CRF	Collective CRF
Overall	60.7 / **61.0** / 60.8	61.2 / **61.0** / 61.1	68.0 / 60.5 / 64.0	**68.4** / **61.0** / **64.5**
Genre	43.4 / **50.9** / 46.9	44.5 / 50.2 / 47.2	54.2 / 40.9 / 46.6	**55.4** / 41.8 / **47.7**
Instrument	54.3 / 53.5 / 53.9	54.9 / **53.8** / 54.3	72.8 / 48.4 / 58.1	**72.9** / 48.6 / **58.3**
Texture	73.2 / **71.8** / 72.5	74.0 / 71.7 / 72.8	75.1 / 71.0 / 73.0	**75.2** / 71.0 / **73.1**
Vocal	76.5 / 71.3 / 73.8	76.7 / 71.2 / 73.8	**80.6** / 84.3 / 82.4	**80.6** / **85.4** / **82.9**
Affective	46.8	**47.5**	42.1	43.1
Arousal	56.5	56.6	57.0	**58.5**
Rhythm	63.0	62.3	64.0	**64.5**
Tempo	59.2	59.4	**63.3**	63.0
Tonality	56.9	57.4	**60.4**	60.1
Production	93.0	93.0	94.5	**94.6**

Table 3. Average per-category performance in the whole vocabulary and each semantic category. For each item, the three numbers are arranged in the format "Precision / Recall / F-measure". Note that for the lower 6 semantic categories, precision, recall and F-measure are the same, and hence written as one number.

consider those with strong correlations. In this case, the edges of the graph are sparse. Suppose the number of edges is E, and all the label variables have C possible values (in our case, C is 2, representing the presence and absence of each label.), then the number of parameters to be estimated in total is $|R||Y|C + |R|EC^2$.

In Eq.(10), the potential of edges are feature-dependent. It can also be degenerated to feature-independent as

$$l \in < y_j, y_{j'} >: 1 \leq j, j' \leq |\mathcal{Y}| \qquad (11)$$

In this case, the number of parameters to be estimated in total is $|R||Y|C + EC^2$.

In order to reduce the computational complexity, we adopt the degenerated CRF model, where the edges are feature-independent, as in Eq.(11). Besides, each song is treated as a sample, and a 115-dimensional song-level feature vector is calculated and set as the characteristic set R. It consists of two parts: a 65-dimensional vector, which is the mean of the beat-level features, and a 50-dimensional vector, with each dimension representing the likelihood of the song given an semantic label in vocabulary, calculated using Eq.(5). The 50-dimensional vector can also be seen as a song representation in an anchor space [1], where each anchor represents one of the 50 semantic labels in the vocabulary.

5 EXPERIMENTS

In this section, the two proposed collective annotation methods are evaluated and compared with two individual annotation methods: the individual GMM-based method with Eq.(4), and the individual CRF-based method which does not consider the edges (or label pairs) in the graph. The experimental data set consists of 4,951 Western popular songs, each of which was manually annotated with semantic labels from the vocabulary in Table 1, as described in Section 2. 25% of the songs are randomly selected as the training set and the left as the test set. For the collective methods, 49 label pairs in total, whose NormMI values in the training set are larger than 0.1, are selected to be modeled.

The annotation performance is measured from two aspects: the *category* aspect and the *song* aspect. From the category aspect, the annotation performances in different categories are measured. From the song aspect, the average annotation performance of each song is evaluated. For both aspects, the average *precision*, *recall* and *F-measure* are used as evaluation metrics, where F-measure is defined as

$$\text{F-measure} = \frac{2 \times \text{precision} \times \text{recall}}{(\text{precision} + \text{recall})} \qquad (12)$$

Table 3 lists the average per-category performances of the four methods. It can be seen that the CRF-based methods outperform the GMM-based methods generally, which accords with previous experiences that discriminative methods generally outperform generative methods in classification problems. Besides, the collective annotation methods slightly but consistently improve the performance, compared with their individual counterpart, both for GMM-based methods and CRF-based methods. This indicates that the label pair modeling helps annotatoin in some cases.

Table 4 presents the performances of song annotations, comparing with four methods. It can be seen that, while the recalls are similar for all the methods, the precision is improved significantly from the generative models to discriminative models. Besides, the collective methods slightly outperform their individual counterparts, which are consistent with the observations made above.

However, the performance improvements from individual modeling to collective modeling is not so much. Although the level of improvement accords with the experiments in [5], we still need to further discover reasons and

%	Precision	Recall	F-measure
Individual GMM	60.9	61.4	60.9
Collective GMM	61.4	61.4	61.2
Individual CRF	68.1	60.9	64.0
Collective CRF	**68.5**	**61.3**	**64.4**

Table 4. Performance of song annotation, comparing with four methods.

exploit solutions. One possible reason may be that, in individual modeling methods, although each label is modeled individually, the labels which are "correlated" share many songs in their training set (since each song has multiple labels). This makes the trained models of "correlated" labels are also "correlated", or in other words, the correlation is implicitly modeled.

6 CONCLUSION

In this paper, we presents our attempts to collective annotation of music signals, which not only models individual semantic labels , but also their correlations. In our approach, 50 musically relevant labels are manually selected for music annotation, covering 10 semantic aspects of music perception. Then, normalized mutual information is employed to measure the correlation between two semantic labels, and those label pairs with strong correlation are selected and modeled in two methods, one based on GMM, and the other based on CRF. Experimental results show slight but consistent improvement compared with individual label modeling methods.

There is still considerable room to improve the proposed approach. First, we need further exploit better methods to model label correlation in order to get higher performance improvement. Second, we need also exploit better features. In current approach, only a song-level feature vector is used for CRF-based methods. How to choose effective song-level features or to adapt the bag of features to CRF-based methods is still a challenging task. Finally, we will also try to apply the obtained annotations in various applications, such as music similarity measure, music search or music recommendation. We are also like to check the impact of annotation accuracy in these applications.

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TONAL PITCH STEP DISTANCE: A SIMILARITY MEASURE FOR CHORD PROGRESSIONS

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ABSTRACT

The computational analysis of musical harmony has received a lot of attention the last decades. Although it is widely recognized that extracting symbolic chord labels from music yields useful abstractions, and the number of chord labeling algorithms for symbolic and audio data is steadily growing, surprisingly little effort has been put into comparing sequences of chord labels.

This study presents and tests a new distance function that measures the difference between chord progressions. The presented distance function is based on Lerdahl's Tonal Pitch Space [10]. It compares the harmonic changes of two sequences of chord labels over time. This distance, named the Tonal Pitch Step Distance (TPSD), is shown to be effective for retrieving similar jazz standards found in the Real Book [3]. The TPSD matches the human intuitions about harmonic similarity which is demonstrated on a set of blues variations.

1 INTRODUCTION

Among musicians and music researchers, harmony is considered a fundamental aspect of western tonal music. For centuries, analysis of harmony has aided composers and performers in understanding the tonal structure of music. The chord structure of a piece alone reveals modulations, tonal ambiguities, tension and release patterns, and song structure [11]. Not surprisingly, the modeling of tonality and computational harmonic analysis have become important areas of interest in music research. Such models can play an important role in content based music information retrieval (MIR). There are obvious benefits in retrieval methods based on harmonic similarity. Melodies are often accompanied by a similar or identical chord progression, or songs may belong to a class with a specific harmonic structure, e.g. blues or rhythm changes, but also cover songs or variations over standard basses in baroque instrumental music could be identified by their harmony.

In this article we present a method for matching two sequences of symbolic chord labels that is based on Lerdahl's Tonal Pitch Space (TPS). Lerdahl [10] developed a formal music-theoretic model that correlates well with data from psychological experiments and unifies the treatment of pitches, chords and keys within a single model. Our proposed method uses this model to create step functions that represent the change of harmonic distance in TPS over time. Two step functions can be efficiently compared using an algorithm designed by Aloupis et al. [2]. Therefore the proposed measure is named the Tonal Pitch Step Distance (TPSD).

Contribution: We introduce a new distance function in the domain of polyphonic music that measures the difference between chord progressions. It is key invariant, independent of the sequences' lengths, allows for partial matching, can be computed efficiently, is based on a cognitive model of tonality, and matches human intuitions about harmonic similarity. We illustrate the soundness of the distance measure by applying it on a set of blues variations. The efficacy for retrieval purposes is demonstrated in an experiment on 388 Real Book [3] songs.

2 RELATED WORK

Theoretical models of tonality have a long tradition in music theory and music research. The first geometric representations of tonality date back at least two centuries. Some authors have investigated the formal mathematical properties of harmonic structures [14], but of particular interest for the current research are the models grounded in data from psychological experiments (see for reviews, [7] [8]). A notable model is Chew's [5] spiral array. The Spiral Array is founded on music-theoretical principles (the Riemannian *Tonnetz*) and, as the name suggests, places pitches, chords and keys on a spiral. Chords are represented as three-dimensional shapes within the spiral. Despite the fact that distances between pitches are incompatible with empirical findings [9] [10], Chew's model has yielded some useful and theoretically interesting algorithms. Another important model is Lerdahl's TPS [10], which correlates reasonably well with Krumhansl's empirical data [9] and matches music-theoretical intuitions. TPS serves as a basis of the distance function here presented and will be explained in the next section.

A related problem that has gained a lot of attention as well is the problem of automatic chord labeling. Chord labeling is the task of finding the right segmentation and la-

(a) octave (root) level:	0											(0)
(b) fifths level:	0					7						(0)
(c) triadic (chord) level:	0			4		7						(0)
(d) diatonic level:	0	2		4	5	7		9			11	(0)
(e) chromatic level:	0	1	2	3	4	5	6	7	8	9	10	11 (0)

Table 1. The basic space of the tonic chord in the key of C Major (C = 0, C# = 1, . . . , B = 11), from Lerdahl [10].

bels for a musical piece. A chord label consists of a chord root, triadic quality, inversion, and extensions (additional chord notes). Nowadays, several algorithms can correctly segment and label approximately 80 percent of a symbolic dataset (see for reviews [20] [16]). Within the audio domain, hidden Markov Models are frequently used for chord label assignment [17] [18]. It is widely accepted that chord information from symbolic or audio data yields a relevant and musicological valid abstraction that can aid in discerning the structure of a piece and making similarity judgments.

Both research areas previously touched upon are important matters when it comes to MIR. Although the value of chord descriptions is generally recognized, and various models about the cognition of tonality are available, surprisingly little research has focused on how similar chord sequences relate to each other. Attempts include string matching [12] as a measure of similarity, and analyzing a chord sequences on the basis of rewrite rules [15] [19]. Mauch [13] analyzed the frequencies of chord classes, chord progression patterns within the Real Book data [3] that is used in this study as well. Still, we argue that similarity of chord sequences is underexposed within the MIR field, which is also evident from the fact that there is no MIREX track for chord progression similarity.

3 TONAL PITCH SPACE

The TPS is a model of tonality that fits human intuitions and is supported by empirical data from psychology [9] [1] . The TPS model can be used to calculate the distances between all possible chords and to predict corresponding tension and release patterns. Although the TPS can be used for defining relationships between chords in different keys, it is more suitable for calculating distances within local harmonies [4]. Therefore our here presented distance measure only utilizes the parts of TPS needed for calculating the chordal distances within a given key (this is motivated in section 4).

The basis of the TPS model is the *basic space* (see Table 1) which comprises five hierarchical levels consisting of pitch class subsets ordered from stable to unstable. Pitch classes are categories that contain all pitches one or more octaves apart. The first and most stable level (a) is the root level, containing only the root of the analyzed chord. The next level (b) adds the fifth of the chord. The third level (c) is the triadic level containing all pitch classes of the chord. The fourth (d) level is the diatonic level consisting of all pitch classes of the diatonic scale of the current key. The last and least stable level (e) is the chromatic level containing all pitch classes. The shape of the basic space of C major strongly resembles Krumhansl and Kessler's [9] C major-key profile. For every chord change, the levels (a-c) must be adapted properly and for a change of key, level d must be adapted. The basic space is hierarchical: if a pitch class is present at a certain level, it is also present at subsequent levels.

The basic spaces of chords can be used to calculate distances between these chords. First, the basic space is set to match the key of the piece (level d). Then the levels (a-c) can be adapted to match the chords to be compared. The distance between two chords is calculated by applying the Chord distance rule. Some examples of calculation are given in Tables 2 and 3.

The proposed distance measure uses a Chord distance rule that is slightly different from the Chord distance rule defined in TPS [10] and is defined as follows:

> CHORD DISTANCE RULE: $d(x, y) = j + k$, where $d(x, y)$ is the distance between chord x and chord y. j is the minimal number of applications of the Circle-of-fifths rule in one direction needed to shift x into y. k is the number of non-common pitch classes in the levels (a-d) within the basic spaces of x and y together divided by two [2] . A pitch class is non-common if it is present in x or y but not in both chords.
>
> CIRCLE-OF-FIFTHS RULE: move the levels (a-c) four steps to the right or four steps to the left (modulo 7) on level d [3] .

When plotted geometrically, the distances exhibit a regular pattern combining the diatonic circle of fifths horizontally and the common tone circle vertically (see Figure 1). If the chordal space is extended, it forms a toroidal structure. Because we are interested in the metrical properties of the chord distance rule it is good to observe that it has a maximum of 13 (see Table 4) [4] . This maximum can be obtained, for instance, by calculating the distance between a C major chord and an E chord containing all notes of the chromatic scale.

[1] The TPS is an elaborate model; due to space limitation we have to refer to [10], chapter 2, pages 47 to 59, for a more detailed explanation and additional examples.

[2] This calculation of k deviates from the calculation described in [10]. Lerdahl defined k as the number of non-common pitch classes in the levels (a-d) within the basic space of y compared to those in the basic space of x. This definition has the undesirable side effect of making the distance function non-symmetrical. Our proposed calculation preserves the symmetry of the distance function and yields equal or similar scores.

[3] If the chord root is non-diatonic j receives the maximum penalty of 3.

[4] Chords with a TPS score of 13 are not musically realistic, but it is useful from a computational point of view to observe that the TPS, and hence the TPSD, has a maximum score.

0							7				
0		2					**7**				
0		2		**4**	5		**7**				11
0		2		4	5		7		9		11
0	1	2	3	4	5	6	7	8	9	10	11

Table 2. The distance between the C and G7 in the context of a C major key. The bold numbers in the basic space are specific for the C major chord and the underlined numbers are specific for the G7 chord. The only common pitch class is G (7). All other 9 pitch classes in the levels (a-c) are non-common, therefore $k = \frac{9}{2}$. $j = 1$ because the Circle-of-fifths rule is applied once. The total score is: $1 + 4.5 = 5.5$.

		2									
		2						**9**			
		2			5	**6**		**9**			
	1	2		4	5	6	7		9		11
0	1	2	3	4	5	6	7	8	9	10	11

Table 3. The distance of a D and a Dm chord in the context of a D major key. The non-common pitch classes are the minor and major thirds (5 and 6), therefore $k = \frac{3}{2} = 1.5$. Note that the minor third does not belong to the D major scale and adds 2 non-common pitch classes. There is no application of the Circle-of-fifths rule ($j = 0$), hence the total score is $1.5 + 0 = 1.5$.

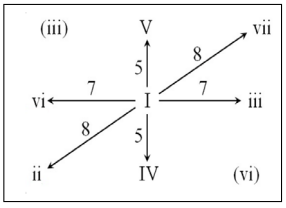

Figure 1. A portion of Lerdahl's chordal space with the roman numbers denoting the seven basic chords in an arbitrary key. The arrows denote the distances as calculated by the Chord distance rule.

0				4							
0				4			**7**				11
0	1	2	3	**4**	5	6	**7**	8	9	10	11
0	1	2	3	4	5	6	7	8	9	10	11
0	1	2	3	4	5	6	7	8	9	10	11

Table 4. Maximum chordal distance, demonstrated by comparing a C major chord with a chord on E chord containing all pitch classes. The bold numbers are specific for the C major chord and the underlined numbers are specific for the chord on E. The numbers that are underlined and bold are the common pitch classes, the remaining pitch classes are non-common, hence $k = 20$. $j = 3$ because the Circle-of-fifths rule is applied three times and yields its maximal value. The total score is $d(x, y) = \frac{20}{2} + 3 = 13$.

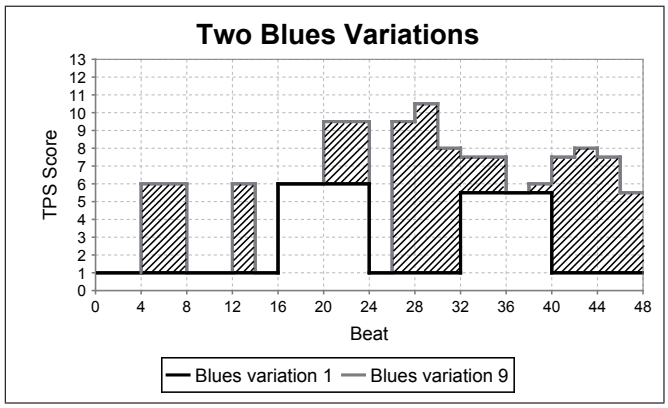

Figure 2. A plot of the step functions of Blues variation 1 and 9 (see Table 5). The TPSD is defined as the hatched area between Blues variation 1 and 9 divided by the length of the shortest step function (which can be either one in this example).

4 TONAL PITCH STEP DISTANCE

On the basis of the Chord distance rule, we define a distance function for chord sequences, called the Tonal Pitch Step Distance (TPSD). The TPSD compares two chord sequences and outputs a number between 0 and the maximum chordal distance 13. A low score indicates two very similar chord sequences and a high score indicates large harmonic differences between two sequences.

The central idea behind the TPSD is to compare the change of chordal distance to the tonic over time. This is done by calculating the chordal distance between each chord of the song and the triadic tonic chord of the key of the song. The reason for doing so, is that if the distance function is based on comparing subsequent chords, the chord distance depends on the exact progression by which that chord was reached. This is undesirable because very similar but not identical chord sequences can then produce radically different scores.

Plotting the chordal distance against the time results in a step function. The difference between two chord sequences can then be defined as the minimal area between the two step functions f and g over all possible horizontal shifts t of f over g (see Figure 2). These shifts are cyclic and to prevent longer sequences from yielding higher scores, the score is normalized by dividing it by the length of the shortest step function. Trivially, the TSPD can handle step functions of different length since the area between non-overlapping parts is always zero.

The calculation of the area between f and g is straightforward. It can be calculated by summing all rectangular strips between f and g, and trivially takes $O(n+m)$ time where n and m are the number of chords in f and g, respectively. An important observation is that if f is shifted along g, a min-

imum is always obtained when two vertical edges coincide. Consequently, only the shifts of t where two edges coincide have to be considered, yielding $O(nm)$ shifts and a total running time of $O(nm(n+m))$. For the results presented here we used this simple algorithm, but the running time can be further improved to $O(nm \log(n+m))$ by applying an algorithm proposed by Aloupis et al. [2]. They present an algorithm that minimizes the area between two step functions by shifting it horizontally as well as vertically.

We do not use the functionality of TPS to calculate distances between chords in different keys. We choose to do so for two reasons. First, modulation information is rarely present in chord sequence data. Second, we are interested in the harmonic similarity of chord sequences regardless of their keys. This implies that to analyze a chord sequence, the key of the sequence must be provided. The TSPD is not a metric, it does not satisfy the property of *identity of indiscernibles*; since two different chords can have the same Lerdahl distance (see Figure 1), it is possible to construct two different chord sequences with the same TPSD.

5 EXAMPLES

To illustrate how our measure behaves in practice, the distances are calculated for a number of blues progressions. Table 5 shows seventeen variations on a twelve bar blues (the last 12 columns) by Dan Hearle found in [1]. The progressions read from left to right with the numbers in the header denoting the bar. The progression in the top row is a very simple blues and as one moves down the progressions become more complex, more altered, and more difficult to play. The second column displays the distance between the progression in first row and the progression in the current row. The third column shows the distance between two subsequent progressions.

We can make the following observations. The scores correspond well to our intuitive idea of similarity between chord progressions. The scores between similar progressions are small and as progressions become more complex the calculated distance with respect to the most simple blues progression becomes higher.

6 EXPERIMENT

We tested the efficacy of the TPSD for retrieval purposes in an experiment. We used a collection of 388 sequences of chord labels that describe the chords of 242 jazz standards found in the Real Book [3]. The chord label data comes from a collection of user-generated Band-in-a-Box files; Band-in-a-Box is a commercial software package that can be used for generating musical accompaniment. The authors manually checked the files for consistency, quality and correct key. Within this collection, 85 songs contain two or more similar versions, forming 85 classes of songs. These songs have the same title and share a similar melody, but can differ in a number of ways. They can, for instance, differ in key and form, they may differ in the number of repetitions, or have a special introduction or ending. The richness of the chords descriptions can also diverge, i.e. a C7b9b13 may be written instead of a C7, and common substitutions frequently occur. Examples of the latter are relative substitution, i.e. Am instead of C, or tritone substitution, i.e. F#7 instead of C7.

Although additional information about timing and tempo in jazz can contain valuable cues that could be helpful for MIR [6], we only used the chord-per-beat information in our step functions. All songs with multiple versions are used as queries and all other 387 songs are ranked on their TPSD score. We then evaluate the ranking of the other versions.

7 RESULTS

Table 6 shows the results of the experiment. The second and seventh columns display the average first tier per song class. The first tier is the number of correctly retrieved songs within the best $(C-1)$ matches divided by $(C-1)$, where C is the size of the song class. Trivially, by using all songs within a class as a query, C first tiers are calculated and averaged for every class of songs. The third and eighth columns show the average second tier. The second tier is the number of correctly retrieved songs within the best $(2C-1)$ matches, divided by $(2C-1)$.

The grand averages [5] of the average first and second tiers are 74 and 77 percent, respectively. This implies that in 74 percent of the song classes the songs searched for are on top of the ranking. Seven song classes in Table 6 contain one or more matches with a TPSD score of 0.00; such a perfect match occurs when two step functions are identical for at least the length of the shortest chord sequence. If these matches are removed from the results, the grand averages of the first and second tiers become 71 and 75 percent, respectively.

8 CONCLUSION

We introduced a new distance function, the Tonal Pitch Step Distance, for chord progressions on the basis of harmonic similarity. The distance of a chord to the tonic triad of its key was determined by using a variant of Lerdahl's Tonal Pitch Space. This cognitive model correlates with empirical data from psychology and matches music-theoretical intuitions. A step function was used to represent the change of chordal distance to its tonic over time. The distance between two chord progressions was defined as the minimal area between two step functions.

[5] Song classes are not weighted on the basis of their size in the calculation of the grand average.

i	d(1,i)	d(i,i-1)	1	2	3	4	5	6	7	8	9	10	11	12
1	0.00	0.00	F7				Bb7		F7		C7		F7	
2	0.42	0.42	F7				Bb7		F7		C7	Bb7	F7	C7
3	1.00	0.67	F7	Bb7	F7		Bb7		F7		G7	C7	F7	C7
4	1.58	0.58	F7	Bb7	F7		Bb7		F7	D7	G7	C7	F7	C7
5	1.62	0.12	F7	Bb7	F7		Bb7		F7	D7	Gm7	C7	F7	Gm7 C7
6	2.31	0.69	F7	Bb7	F7		Bb7	Eb7	F7	D7	Db7	C7	F7	Db7 C7
7	2.75	1.10	F7	Bb7	F7	Cm7 F7	Bb7	Eb7	F7	Am7 D7	Gm7	C7	Am7 D7	Gm7 C7
8	3.31	0.56	F7	Bb7	F7	Cm7 F7	Bb7	Eb7	Am7	D7	Gm7	C7	Am7 D7	Gm7 C7
9	3.17	0.56	F7	Bb7	F7	Cm7 F7	Bb7	Bm7 E7	F7 E7	Eb7 D7	Gm7	C7 Bb7	Am7 D7	Gm7 C7
10	4.29	2.12	FM7	Em7 A7	Dm7 G7	Cm7 F7	Bb7	Bdim7	Am7 D7	Abm7 Db7	Gm7 C7	Dbm7 Gb7	F7 D7	Gm7 C7
11	5.12	2.08	FM7	Em7 Ebm7	Dm7 Dbm7	Cm7 Cb7	BbM7	Bbm7	Am7	Abm7	Gm7	C7	Am7 Abm7	Gm7 Gb
12	4.88	1.50	FM7	BbM7	Am7 Gm7	Gbm7 Cb7	BbM7	Bbm7	Am7	Abm7	Gm7	Gb7	FM7 Abm7	Gm7 Gb
13	5.23	1.48	FM7	BbM7	Am7 Gm7	Gbm7 Cb7	BbM7	Bbm7 Eb7	AbM7	Abm7 Db7	GbM7	Gm7 C7	Am7 D7	Dbm7 Gb
14	4.40	1.79	FM7	Em7 A7	Dm7 G7	Cm7 F7	BbM7	Bbm7 Eb7	Am7	Abm7 Db7	Gm7	C7	Am7 D7	Gm7 C7
15	4.98	0.75	FM7	Em7 A7	Dm7 G7	Gbm7 Cb7	BbM7	Bm7 E7	Am7	Abm7 Db7	Gm7	C7 Bb7	Am7 D7	Gm7 C7
16	5.42	1.94	F#m7 B7	Em7 A7	Dm7 G7	Cm7 F7	BbM7	Bbm7 Eb7	AbM7	Abm7 Db7	GbM7	Gm7 C7	Am7 D7	Gm7 C7
17	5.71	2.88	FM7	F#m7 B7	EM7 EbM7	DbM7 BM7	BbM7	Bm7 E7	AM7	Am7 D7	GM7	GbM7	FM7 AbM7	GM7 Gb

Table 5. Seventeen blues variations and the TPSD scores between each progression and the first one (second column), and between each progression and the preceding one (third column).

nr.	1st Tier	2nd Tier	Class size	Title of the song class	nr.	1st Tier	2nd Tier	Class size	Title of the song class
1	1.00	1.00	2	A Child is Born	44	1.00	1.00	2	Miyako
2	0.00	0.00	2	A Fine Romance	45	0.50	0.50	2	Moment Notice
3	1.00	1.00	2	A Night In Tunisia	46	1.00	1.00	3	Mood Indigo
4	0.50	0.50	2	All Blues	47	0.73	0.73	6	More I See You, The
5	0.67	0.83	3	All Of Me	48	0.33	0.33	3	My Favorite Things
6	0.50	0.50	2	Angel Eyes	49	1.00	1.00	4	My Funny Valentine
7	0.50	1.00	2	April In Paris	50	1.00	1.00	3	My Romance
8	0.00	0.00	2	Blue in Green	51	0.50	0.50	2	Nefertiti
9	0.50	0.50	4	Corcovado (Quiet Nights of Quiet Stars)	52	0.67	0.83	3	Nica' Dream
10	1.00	1.00	2	Days of Wine and Roses, The	53	1.00	1.00	2	Night Dreamer
11	1.00	1.00	2	Dearly Beloved	54	0.00	0.00	2	Night Has a Thousand Eyes, The
12	0.00	0.00	2	Desafinado	55	0.33	0.33	3	Oleo
13	1.00	1.00	2	Don't Get Around Much Anymore	56	1.00	1.00	3	Once I Loved
14	0.33	0.33	3	Easy to Love	57	1.00	1.00	3	One Note Samba
15	1.00	1.00	2	E.S.P.	58	0.50	0.50	2	Ornithology
16	0.60	0.60	5	Girl from Ipanema	59	1.00	1.00	2	Peace
17	1.00	1.00	2	Green Dolphin Street	60	1.00	1.00	2	Pensativa
18	1.00	1.00	2	Have You Met Miss Jones	61	0.00	0.00	2	Peri's Scope
19	0.70	0.80	5	Here's That Rainy Day	62	1.00	1.00	4	Satin Doll
20	1.00	1.00	4	Hey There	63	1.00	1.00	2	Scrapple from the Apple
21	0.63	0.67	6	How High the Moon	64	0.67	1.00	3	Shadow of Your Smile, The
22	1.00	1.00	3	How Insensitive	65	1.00	1.00	3	Solar
23	1.00	1.00	3	If You Never Come To Me	66	0.33	0.50	3	Some Day My Prince Will Come
24	0.00	0.00	2	I Love You	67	0.33	0.33	3	Song is You, The
25	1.00	1.00	2	I Mean You	68	1.00	1.00	3	Sophisticated Lady
26	1.00	1.00	2	In A Sentimental Mood	69	0.67	1.00	3	So What
27	0.00	0.00	2	Isotope	70	1.00	1.00	3	Stella by Starlight
28	1.00	1.00	3	Jordu	71	1.00	1.00	3	Stompin' at the Savoy
29	1.00	1.00	2	Joy Spring	72	0.67	1.00	3	Straight, No Chaser
30	1.00	1.00	2	Just Friends	73	1.00	1.00	2	Take the "A" Train
31	1.00	1.00	2	Lament	74	1.00	1.00	3	There is No Greater Love
32	1.00	1.00	2	Like Someone In Love	75	0.50	0.50	2	They Can't Take That Away From Me
33	1.00	1.00	3	Limehouse Blues	76	1.00	1.00	2	Triste
34	1.00	1.00	2	Little Waltz	77	1.00	1.00	2	Tune Up
35	0.83	1.00	4	Long Ago and Far Away	78	0.33	0.33	3	Wave
36	1.00	1.00	2	Look to the Sky	79	1.00	1.00	3	We'll Be Together Again
37	0.33	0.33	3	Lucky Southern	80	1.00	1.00	3	Well You Needn't
38	1.00	1.00	3	Lullaby of Birdland	81	1.00	1.00	3	When I Fall In Love
39	1.00	1.00	2	Maiden Voyage	82	0.17	0.25	4	Yesterdays
40	0.33	0.33	3	Meditation	83	0.45	0.60	5	You Are the Sunshine of My Life
41	1.00	1.00	2	Memories of Tommorow	84	1.00	1.00	3	You Are Too Beautiful
42	0.00	0.17	3	Michelle	85	1.00	1.00	2	You Don't Know What Love Is
43	1.00	1.00	2	Misty	**avg**.	**0.74**	**0.77**		

Table 6. 78 song classes of jazz standards found in the Real Book. The first and second tier results show the retrieval efficacy of the TPSD.

We showed that the TPSD as a retrieval method yields promising results. Similar versions of the same jazz standard found in the Real Book can be successfully retrieved on the basis of their chord progressions. The soundness of the TPSD was demonstrated on a series of blues variations.

9 FUTURE WORK

The problem in analyzing the retrieval quality of the TPSD is the lack of good ground truth data. We do not know of any collection of chord sequences that contain user generated similarity labels. It would be interesting to acquire such similarity data and further explore the performance of the TPSD. This could very well be done in a MIREX track. Another issue concerns the used dataset. The Real Book collection is a small collection and only contains songs of a specific musical genre: jazz standards. It would be interesting to explore the performance of the TPSD on a larger dataset and in other musical domains.

We can suggest several improvements of the TPSD as well. Currently, the TPSD does not treat modulations in a musical way. If a modulation occurs within a piece, the scores of the TPSD become very high, and although this enables the TPSD to recognize these modulations, the step function loses nuance. Applying a key finding algorithm locally might yield more subtle results. Another idea is to further exploit the fact that TPSD is very suitable for partial matching by using it for tracing repetitions within a query or matching meaningful segments of a query instead of the query as a whole.

We believe that further improvement of chord labeling algorithms and the development of tools that analyze these labels should be high on the research agenda because chord labels form an abstraction of musical content with substantial explanatory power. In the future we expect TPSD based methods to help users find songs in large databases on the Internet, or in their personal collections. We believe that retrieval on basis of harmonic structure is crucial for the next generation of content based MIR systems.

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A Study on Feature Selection and Classification Techniques for Automatic Genre Classification of Traditional Malay Music

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ABSTRACT

Machine learning techniques for automated musical genre classification is currently widely studied. With large collections of digital musical files, one approach to classification is to classify by musical genres such as pop, rock and classical in Western music. Beat, pitch and temporal related features are extracted from audio signals and various machine learning algorithms are applied for classification. Features that resulted in better classification accuracies for Traditional Malay Music (TMM), in comparison to western music, in a previous study were beat related features. However, only the J48 classifier was used and in this study we perform a more comprehensive investigation on improving the classification of TMM. In addition, feature selection was performed for dimensionality reduction. Classification accuracies using classifiers of varying paradigms on a dataset comprising ten TMM genres were obtained. Results identify potentially useful classifiers and show the impact of adding a feature selection phase for TMM genre classification.

1. INTRODUCTION

Interest on music information retrieval systems for the storage, retrieval and classification of large collections of digital musical files has grown in recent years. Metadata such as filename, author, file size, date and genres are commonly used to classify and retrieve these documents. Such manual classification is highly labour-intensive and costly both in terms of time and money [1].

An automatic classification system that is able to analyse and extract implicit knowledge of the musical files is therefore highly sought. One approach to musical classification that is currently being widely studied is classification by musical genres. Musical genres are labels created and used by humans for categorizing and describing music [2].

Examples of a few Western musical genres are such as Pop, Rock, Hip-hop, and Classical. Several systems for automated genre classification and retrieval of musical files have been researched and developed [2,3]. However, most of these studies were conducted using only western dataset and we focus on non-Western musical genres, and more specifically Traditional Malay Music (TMM).

TMM encompasses all traditional music from Malaysia, both West Malaysia (on the Peninsular) and the states on the Borneo Island (Sabah and Sarawak) [4]. Genre examples include Dikir Barat, Joget, Wayang Kulit, Gamelan, Etnik Sabah and Inang. In general, these musical genres have a strong sense of rhythm, partly due to the fact that TMM is traditionally played by ear as opposed to reading from written musical scores. Having the beat or rhythm clearly audible helps when the musical piece is being passed down orally through generations in the villages such as having clear gong hits. The significance of beat features for TMM genre classification in comparison to Western musical genres was investigated in a previous study using a two-phase methodology – feature extraction and classification [5]. In this paper we study the impact of adding a feature selection phase for TMM genre classification.

Feature extraction is a process where a segment of an audio is characterized into a compact numerical representation. Due to the high dimensionality of these feature sets, feature selection can be performed to reduce the dimensionality of the data as a preprocessing step prior to classification. With audio data, several studies have investigated the significance of this phase and the performance of several feature selection methods [21]. A comprehensive discussion on feature selection is available in Saeys et. al [6].

A large number of the studies performed on music classification have looked more closely at the feature extraction step, and in this study we investigate the classification phase more closely. The rest of this paper is organised as follows: Section 2 presents background information about music genre classification and TMM. An overview of feature selection is presented in Section 3. Section 4 presents the experimental framework and results are discussed in Section 5.

2. BACKGROUND

2.1. Music Genre Classification

Digital audio in general is categorized as speech, music and noise. Wold, et al[3] analyse and compare audio features such as rhythms, pitch, duration, loudness and instrument identification to classify various groups of audio such as speech, gender, animal sounds and sound

effects. However, music classification was not emphasized in their study.

There has been great interest of classification of audio based on musical genres in recent years [2, 7,8]. Tzanetakis and Cook [2] and Aucoturier and Pachet[8] categorized audio features into three categories; timbral related features, rhythmic related features, and pitch related features. Audio data that are to be classified cannot be represented as raw audio data, such as samples of amplitude value of digital audio signals. Hence, some form of parameterisation is required. Parameterisation of audio data is based on audio analysis, which can be done using several methods such as Fourier transform, wavelet transform, statistical methods, etc. [9].

Timbral related features are based on the Short Time Fourier Transform (STFT), which is used to determine the phase content of short local sections in a signal as it changes over time. The features are used in music-speech discrimination and speech recognition. Examples of timbral related features are such as spectral centroid, spectral roll off and time domain zero crossing, which measure the spectral shape, the changes in spectral shape and the noisiness of a signal respectively. Another feature, Mel-Frequency Cepstral Coefficients (MFCC), is also based on the STFT, but is typically used to provide compact representation of the spectral envelope, especially in speech recognition. Tzanetakis and Cook [2] provide a detailed discussion on timbral related features and the account of their experiment.

Beat features analyses the signals that calculate the rhythmic structure of music based on their wavelet transform [2]. It involves time-frequency analysis, which is useful to music classification as its algorithm is similar to human hearing. The main beat can be defined as the regular periodic sequence of pulses corresponding to where a human would tap his foot whilst listening to music. Although achievable, extraction of beat features is very difficult. Whilst it is trivial for human to do so to a music, it is not so with machines. Kosina[10] and Dixon[11] give good overviews on beat tracking methods. Li and Tzanetakis[12] investigate the effects of different feature set combinations for optimum classification performance. Features incorporated in the study were: FFT, MFCC, Pitch and Beat. Although suitable feature set combinations from this study was obtained, it was also suggested that they might not be generic to all genres but applicable only to western genres that was used in their study.

With classification, classifiers vary in terms of robustness, speed, memory usage and complexity. Several studies investigate the use of existing classifiers for musical genre classification [2,3,12]. For instance, OneR is a primitive form of classifier as it produces simple rule based on one attribute only, but it is useful in determining a baseline performance as a benchmark for other learning schemes [13]. Emphasis on the importance of understanding different classifiers is also discussed at length by [7,14].

2.2. Traditional Malay Music

Traditional Malay music is mainly derivative, influenced by the initial overall Indian and Middle Eastern music during the trade era and later from colonial powers such as Thailand, Indonesia, Portuguese and British who introduced their own culture including dance and music. A thorough overview on the origin and history of TMM can be found in [17]. The taxonomy of TMM depends on the nature of the theatre forms they serve and their instrumentations. Categorization of TMM genres has been studied extensively by Ang[18]. Music of these genres is usually disseminated non-commercially, usually performed by persons who are not highly trained musical specialists, undergoes change arising from creative impulses and exists in many forms. The musical ensembles usually include gendangs or drums that are used to provide constant rhythmic beat of the songs and gongs to mark the end of a temporal cycle at specific part of the song [19].

One common attribute that is shared by most TMM genres is that they are generally repetitive in nature and exist in ‘gongan’-like cycle. ‘Gongan’ is defined as a temporal cycle marked internally at specific points by specific gongs and at the end by the lowest-pitched gong of an ensemble [17]. It is an important structural function as it divides the musical pieces into temporal sections. Once every measure has been played, musicians continue playing in a looping motion by repeating the cycle from the beginning again until one of the lead percussionists signals the end of the song by varying their rhythms noticeably. Traditional Malay music does not have a chorus that plays differently than other parts of the songs, which is the usual occurrence in western music. Its repetitiveness and constant rhythms are two aspects that are taken into account to facilitate classification by genre later.

Very little study has been conducted on automatic traditional Malay music genre classification in the literature. Norowi, et al[20] study the effects of various factors and audio feature set combinations towards the classification of TMM genres. Results from experiments conducted in several phases show that factors such as dataset size, track length and location¸ together with various combinations of audio feature sets comprising Short Time Fourier Transform (STFT), Mel-Frequency Cepstral Coefficients (MFCCs) and Beat Features affect classification. Based on parameters optimized for TMM genres, classification performances were evaluated against three groups of human subjects:

experts, trained and untrained. Based on the result of this study performances of both machine and human were shown to be comparable. However, only the J48 classifier was used with 66.3% classification accuracy [5]. In this study, we assess the practical usefulness of a wide range of classifiers and identify potential classifiers that would improve the performance of TMM genre classification. We confine our study to the classifiers within WEKA which is discussed further in section 4.2.

3. FEATURE SELECTION

Feature selection is the process of removing features from the data set that are irrelevant with respect to the task that is to be performed. Feature selection can be extremely useful in reducing the dimensionality of the data to be processed by the classifier, reducing execution time and improving predictive accuracy (inclusion of irrelevant features can introduce noise into the data, thus obscuring relevant features). It is worth noting that even though some machine learning algorithms perform some degree of feature selection themselves (such as classification trees), feature space reduction can be useful even for these algorithms. Reducing the dimensionality of the data reduces the size of the hypothesis space and thus results in faster execution time.

In general, feature selection techniques can be split into two categories - filter methods and wrapper methods. Wrapper methods generally result in better performance than filter methods because the feature selection process is optimized for the classification algorithm to be used. However, they are generally far too expensive to be used if the number of features is large because each feature set considered must be evaluated with the trained classifier. For this reason, wrapper methods will not be considered in this study. Filter methods are much faster than wrapper methods and therefore are better suited to high dimensional data sets. Diverse feature ranking and feature selection techniques have been proposed in the machine learning literature, Such as:

- Correlation-based Feature Selection (CFS) [21]
- Principal Component Analysis (PCA) [21]
- Gain Ratio (GR) attribute evaluation [21]
- Chi-square Feature Evaluation [21]
- Support Vector Machine Feature Elimination (SVM-RFE) [22]

Some of these methods does not perform feature selection but only feature ranking, they are usually combined with another method when one needs to find out the appropriate number of attributes. Forward selection, backward elimination, bi-directional search, best-first search [13], genetic search [23], and other methods are often used on this task.

Fiebrink et. al [23] investigated the significance of the addition of feature selection with classification of Western musical genres. The results showed an almost similar classification accuracy using forward selection and PCA, a wrapper and filter method respectively. Classification utilized a fraction of time with PCA. We evaluate several filter methods for TMM genre classification in this study to achieve to our purpose: choose the best combination of feature selection and classification to classify the TMM genre.

4. EXPERIMENTS

The aim of this study is to investigate the impact of the addition of feature selection towards TMM classification. For this purpose some experiments were designed and conducted and explained in this section.

4.1. Data Set

The data collection and pre-processing stages of this study are described in these following sub-sections.

4.1.1. Data Collection

Ten TMM genres were involved in this study. The breakdown for each genre and its number of musical files are listed in Table 1. A relatively small dataset was used in this experiment due to the difficulty in obtaining digital files of TMM, as traditional Malay musical culture is fast corroding with little preservation in digital format. Whilst it was much easier to obtain dataset for western music, the number was also kept small to match the size of TMM dataset.

NO	Genre	Number
1	Dikir Barat	31
2	Etnik Sabah	12
3	Gamelan	23
4	Ghazal	17
5	Inang	10
6	Joget	15
7	Keroncong	43
8	Tumbuk Kalang	13
9	Wayang Kulit	17
10	Zapin	10

Table 1. Overall number of musical files for each genre

Musical files for this experiment were obtained from the Malaysia Arts Academy, Sultan Salahuddin Abdul Aziz Shah's Cultural and Arts Centre at Universiti Putra Malaysia, Student's Cultural Centre at Universiti Malaya and also personal collections of audio CDs from many individuals. The dataset became available in both digital and analog format. Quite a number of musical data for TMM genres were in analog format and were digitized manually. All of the digital music files were then

converted into wav files; the only audio format supported by the existing feature extraction tool used at the time of study. The whole dataset was later trimmed to specific length and location in the file by executing certain audio commands through batch processing before extraction began.

4.2. Genre Classification Component

This section discusses feature extraction and classification using Musical Research System for Analysis ad Synthesis (MARSYAS) [2] and Waikato Environment for Knowledge Analysis (WEKA) [13]. We use MARSYAS for feature extraction and WEKA for feature selection and classification.

4.2.1. Feature Extraction

The features were extracted from the music files through MARSYAS-0.2.2; a free framework that enables the evaluation of computer audition applications. MARSYAS is a semi-automatic music classification system that is developed as an alternative solution for the existing audio tools that are incapable of handling the increasing amount of computer data [2]. It enables the three feature sets for representing the timbral texture, rhythmic content and pitch content of the music signals and uses trained statistical pattern recognition classifiers for evaluation. The feature extractor will produce numerical outputs in the form of Attribute Related File Format (ARFF) files. In this study we extracted STFT + MFCC + Beat Feature because this combination of features had been achieved best accuracy for TMM genre classification in [5] and also includes the complete set of features (73 features).

Scenario	Description
S1	**No Feature selection**
S2	**Correlation-based Feature Selection with best first search strategy**
S3	**Correlation-based Feature Selection with genetic search strategy**
S4	**Correlation-based Feature Selection with greedy stepwise search strategy**
S5	**Principal Component Analysis**
S6	**Chi-square Feature Evaluation**
S7	**Gain Ratio Feature Evaluation**
S8	**SVM based Feature Evaluation**

Table 2. Description of scenarios

4.2.2 Feature Selection

We used seven feature selection methods in the experiments of study. There are eight scenarios at the experiments in which one scenario does not incorporate a feature selection steps. The description of scenario is shown in Table 2.

4.2.3 Classification

To compare the performance of classification algorithms, the list of classifiers chosen included a wide range of paradigms. The code written was based on the WEKA data mining package and the default parameters used for each algorithm. All experiments were carried out using a ten-fold cross validation approach and to control the validity of experiments. The list of classifiers and results of the experiments are shown in Table 3. These include AIRS (a classifier based on the Artificial Immune System (AIS) paradigm [25,26]), Bagging, Bayesian Network, Cart, Conjunctive rule learner (Conj-Rules), Decision Stump, Decision Table, IB1, J48 (an implementation of C4.5), Kstar, Logistic, LogitBoost, Multi-layer neural network with back propagation (MLP), Naïve Bayesian, Nbtree, PART (a decision list [27]), RBF Network, and SMO (a support vector machine [28]).

Of these classifiers, AIRS is discussed a little further. It's performance for musical genre classification has not been widely investigated. AIS is a computational method inspired by the biology immune system. It is progressing slowly and steadily as a new branch of computational intelligence and soft computing [29]. One of the AIS based algorithms is the Artificial Immune Recognition System (AIRS). AIRS is a supervised immune-inspired classification system capable of assigning data items unseen during training to one of any number of classes based on previous training experience. AIRS is probably the first and best known AIS for classification, having been developed in 2001. [29]. This study also investigates the performance of this algorithm for musical genre classification.

5. RESULTS

Table 5 lists the classifiers that obtained highest classification accuracies for each of the described scenarios. The descriptive details of the classification are shown in Table 4.

The highest accuracy was obtained with MLP using the Correlation-based Features Selection and a genetic search strategy. Although none of the evaluated classifiers and feature selection methods provided us with a combination that outperforms a particular combination, useful knowledge was gained regarding combinations that do not contribute significantly to the task. The functional-based classifiers, MLP and SMO prove to be superior to the other classifiers. The combination of MLP and Correlation-based Feature Selection with genetic search strategies has achieved best accuracy of 88.6%.

	min	max	Avg
S1	32.64	86	68.13
S2	33.68	87	71.96
S3	33.68	88.6	71.36
S4	33.68	87	72.25
S5	35.23	83.42	60.88
S6	31.6	87.05	71.45
S7	31.6	87.05	72.37
S8	31.6	84.46	69.89

Table 4. Descriptive statistics of scenarios

Scenario	Max Accuracy (%)	Classifier
S1	86	MLP
S2	87	MLP
S3	88.6	MLP
S4	87	MLP
S5	83.42	SMO
S6	87.05	AIRS-SMO
S7	87.05	AIRS
S8	84.46	SMO

Table 5. Max accuracy achieved by classifiers in each scenario

	Accuracy (%)								
Classifier	**S1**	**S2**	**S3**	**S4**	**S5**	**S6**	**S7**	**S8**	**Avg**
AIRS	51.81	84.97	82.90	84.97	59.10	87.05	87.05	83.41	77.66
Bagging	73.57	74.61	75.13	75.65	61.66	75.13	76.68	68.91	72.67
Bayesian Network	78.24	78.76	77.20	78.76	49.22	78.24	80.83	75.13	74.55
Cart	58.55	60.10	62.18	63.73	60	60.62	61.67	61.14	61
Conj-Rules	32.64	33.68	33.68	33.68	35.23	31.60	31.60	31.60	32.96
Decision Stump	33.68	33.68	33.68	33.68	35.75	33.69	33.68	33.68	33.94
Decision Table	60	53.36	56.48	53.37	45.60	51.81	52.85	55.96	53.68
IB1	75.65	84.45	82.38	84.46	67.36	84.97	84.97	80.83	80.63
J48	68.39	66.84	71.50	68.91	56.48	72.53	73.06	66.84	68.07
Kstar	76.68	83.41	82.38	83.42	60.62	82.38	80.83	79.79	78.69
Logistic	83.41	76.16	75.67	76.16	76.68	81.87	86.01	79.79	79.47
LogitBoost	77.20	82.90	81.34	82.90	70.98	76.17	81.35	76.69	78.69
MLP	86	87	88.60	87	82.90	82.90	84.47	83.94	85.35
Naïve Bayesian	78.24	80.82	79.79	80.82	75.65	75.13	77.72	79.79	78.50
Nbtree	63.73	74.61	70.47	74.61	48.70	77.72	75.13	69.43	69.3
PART	66.84	73.10	70.47	71.50	58	68.91	68.39	67.87	68.14
RBF	76.68	80.83	78.24	80.82	68.40	78.24	80.31	78.76	77.79
SMO	84.97	86	82.38	86	83.42	87.05	86.01	84.46	85.04

Table 3. The classifier's accuracies

Another significant observation that can be made is that the addition of the feature selection step has significantly improved the performance accuracy of the AIRS accuracy, an immune-based system. An improvement of almost 30% is obtained. However, the use of PCA did not still improve the performance of AIRS.

6. CONCLUSION

A comprehensive study evaluating the performance of a wide range of classifiers for automating TMM genre classification was performed. Features that were found to be useful for TMM genre classification in comparison to Western genres continued to be used in this study. Results obtained clearly identify potentially useful classifiers -- the functional-based classifiers MLP and SMO. The impact of including a feature selection phase for TMM genre classification was investigated. The results show that the addition of this phase did improve the performance of most classifiers by at least 1%. This addition however improved the performance of the immune-based classifier, AIRS very much as discussed above. Future work will include further experiments to investigate these findings on improved musical genre classification with AIRS and a comparative study to Western Musical genres.

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MOODSWINGS: A COLLABORATIVE GAME FOR MUSIC MOOD LABEL COLLECTION

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ABSTRACT

There are many problems in the field of music information retrieval that are not only difficult for machines to solve, but that do not have well-defined answers. In labeling and detecting emotions within music, this lack of specificity makes it difficult to train systems that rely on quantified labels for supervised machine learning. The collection of such "ground truth" data for these subjectively perceived features necessarily requires human subjects. Traditional methods of data collection, such as the hiring of subjects, can be flawed, since labeling tasks are time-consuming, tedious, and expensive. Recently, there have been many initiatives to use customized online games to harness so-called "Human Computation" for the collection of label data, and several such games have been proposed to collect labels spanning an excerpt of music. We present a new game, MoodSwings (http://schubert.ece.drexel.edu/moodswings), which differs in that it records dynamic (per-second) labels of players' mood ratings of music, in keeping with the unique time-varying quality of musical mood. As in prior collaborative game approaches, players are partnered to verify each others' results, and the game is designed to maximize consensus-building between users. We present preliminary results from an initial set of game play data.

1 INTRODUCTION

The detection and labeling of the emotional content (mood) of music is one of many music information retrieval problems without a clear "ground truth" answer. The lack of easily obtained ground truth for these kinds of problems further complicates the development of automated solutions, since classification methods often employ a supervised learning approach relying on such ground truth labels. The collection of this data on subjectively perceived features, such as musical mood, necessarily requires human subjects. But traditional methods of data collection, such as the hiring of subjects, have their share of difficulties since labeling tasks can be time-consuming, tedious, error-prone and expensive.

Recently, a significant amount of attention has been placed on the use of collaborative online games to collect such ground truth labels for difficult problems, harnessing so-called "Human Computation". For example, von Ahn et al. have created several such games for image labeling: the *ESP Game*, *Peekaboom* [1], and *Phetch*. More recently, several such games have been been proposed for the collection of music data, such as *MajorMiner* [2], *Listen Game* [3], and *TagATune* [4]. These implementations have primarily focused on the collection of descriptive labels for a relatively short audio clip.

We present a new game, *MoodSwings*, designed to explore the unique time-varying nature of musical mood. Of course, one of the joys of music is that the mood of a piece may change over time, gradually or suddenly. According to Huron [5], this combination of anticipation and surprise may be at the core of our enjoyment of music. Thus, our game is targeted at collecting dynamic (per-second) labels of users' mood ratings, which are collected in real-time as a player hears the music using the two-dimensional grid of emotional components: valence and arousal. As in other collaborative games, players are partnered in order to verify each others' results, providing a strong incentive for producing high-quality labels that others can agree upon. Accordingly, game scoring is designed to maximize consensus-building between partners. In this paper, we present data from an initial pilot phase of the game and demonstrate the utility of this approach for the collection of high-quality, dynamic labels of musical affect.

2 BACKGROUND

Models of affect and the categorization and labeling of specific emotions has received significant attention from a variety of research areas including psychology, physiology, neuroscience, as well as musicology. With the advent of digital music and very large music collections, recent work has focused on the problem of automatic music mood detection. Next, we briefly summarize some of the related work.

2.1 Mood models

Early work on the quantification of musical affect focused on the formation of ontologies using clusters of common

emotional adjectives and labels (e.g., "bright", "gloomy", "contemplative", "angry"). Ontologies proposed by Hevner [6] and Farnsworth [7] proposed eight and ten such mood clusters, respectively. All Music Guide [8], a large edited music information database, also uses a descriptive approach with a total of 179 distinct (but not unrelated) mood labels.

An alternative approach to the modeling of human emotions views affect as a combination of orthogonal continuous sub-features. The most popular such representation of musical affect is Thayer's two-dimensional valence-arousal space [9], which itself is derived from Russell's general model of human emotions (pleasant-unpleasant vs. arousal-sleep) [10]. Thayer's model decomposes emotion in music according to two principal dimensions:

- *valence:* positive vs. negative (e.g., happy vs. sad)
- *arousal:* high- vs. low-energy (e.g., energetic vs. calm)

According to this model, music can be broadly categorized into one of four quadrants: high valence and arousal (joy, exuberance), high valence and low arousal (contentment), low valence and high arousal (anger), and low valence and arousal (depression). But the model also views the axes as continuous features, allowing for an unlimited combination of overall moods. The continuous valence-arousal model is at the core of MoodSwings.

2.2 Ground truth label collection

The tedium and expense of label collection has presented a significant obstacle for researchers seeking such labels in order to train automatic systems for mood classification. When employing subjects specifically for the collection of mood labels for music, prior systems have used alternatively expert and non-expert populations.

The Pandora service [11] is an example of a well-known expert labeling system that employs a large team of musicians to manually label tracks according to hundreds of features (their "music genome"), some of which relate to mood. The former MoodLogic service used questionnaires given to their users to collect mood metadata. The data from these services is, sadly, not available to the public. Other commercial tools allow users to tag their own collections, such as the Moody plug-in for iTunes, which uses a quantized 4x4 valence-arousal grid.

More recently, online services that allow users to input free-form (unconstrained) tags, such as Last.fm [12], have collected myriad tags (some of which represent mood labels) across very large music collections. The accessibility of this data has led to a recent trend towards using such free-form tags as the basis for mood labeling by aggregating results from many users. Hu, Bay, and Downie collected labels and tags from AMG, Last.fm, and epinions.com to form the ground-truth mood "clusters" used in the 2007 MIREX mood detection evaluation [13].

2.3 Automatic mood classification of music

The general approach to automatic mood detection from audio has been to use supervised machine learning to train statistical models of acoustic features. Li and Ogihara [14] used acoustic features related to timbre, rhythm, and pitch to train Support Vector Machines (SVMs) to classify music into 13 mood categories derived from Farnsworth's emotion groupings. Using a hand-labeled library of 499 music clips (30-seconds each), they achieved an accuracy of $\sim$45%, with 50% of the database used for training and testing, respectively.

Lu, Liu, and Zhang [15] pursued mood detection and tracking (following dynamic mood changes during a song) using a variety of acoustic features related to intensity, timbre, and rhythm. Their classifier used Gaussian Mixture Models (GMMs) for Thayer's four principal mood quadrants in the valence-arousal representation. The system was trained using a set of 800 classical music clips (from a data set of 250 pieces), each 20 seconds in duration, hand labeled to one of the 4 quadrants. Their system achieved an accuracy of $\sim$85% when trained on 75% of the clips and tested on the remaining 25%.

In 2007, the Music Information Research Evaluation eXchange (MIREX) first included a "beta" task on audio music mood classification with 8 systems submitted. The audio clips used for this task were assigned to one of 5 mood clusters, aggregated from AMG mood labels (adjectives), and 600 30-second hand-labeled clips distributed equally among the 5 mood clusters were used in the evaluations. All participants performed reasonably well (far higher than chance) with the highest performing system [16] achieving correct classifications slightly over 60% of the time. It should be noted that several of the systems were primarily designed for the genre classification and then appropriated to the mood classification task as well [17].

3 MOODSWINGS

MoodSwings is a collaborative, two-player game that incorporates each listener's subjective judgements of the emotional content (mood) of music into the game play. At the start of a match, a player is partnered with another player anonymously across the internet. The goal of the game is for the players to dynamically and continuously reach agreement on the mood of 5 short (30-second) music clips drawn from a database of popular music.

The MoodSwings game board (Figure 1) is a direct representation of the valence-arousal space. The board represents a continuum of possible mood ratings, with arousal on the horizontal axis and valence on the vertical axis. During gameplay, players simultaneously listen to identical short music clips. Each player positions their circular cursor dynamically on the game board, indicating their instantaneous

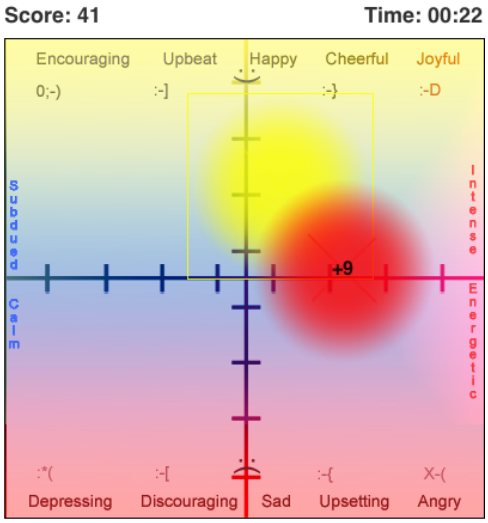

Figure 1. The MoodSwings game board

assessment of the mood of the music. A player's position is sampled once per second, indicated by the pulsing of their cursor. The partner's cursor is visible only intermittently, every few seconds. Scoring is based on the amount of overlap between the partners' circles, with greater congruency resulting in a greater number of points scored. The size of the players' cursors decreases over time as the clip plays, increasing the difficulty of scoring points as time elapses (the players must agree on their ratings more precisely to overlap and thus score points).

3.1 Game play sequence

A MoodSwings match consists of 5 rounds, each consisting of a different music clip with a duration of 30 seconds. Once a partner pairing is established and verified, each round commences after a short 3-second countdown. The partner pairing remains consistent for all 5 rounds. The game board remains inactive until the round begins.

1. Once the round begins, the player's cursor (colored yellow circle) becomes visible. The cursor "throbs" every second indicating the sampled position, but the player is free to continuously alter the cursor position between pulses.
2. After an initial period of 5 seconds, the partner's cursor becomes visible for the first time (again pulsing for one second). This allows a player to make an initial mood assessment independently, without influence from their partner.
3. Afterwards, the partner's cursor is visible once every 3 seconds. This interval is designed to prevent players from simply "chasing" their partners in order to accumulate more points.
4. The size of both cursors decreases continuously during the course of the round.

At the end of each round, the player is presented with their score for the round and their total points for the match thus far. Likewise, at the conclusion of a match, a list is presented with performing artist and title for each music clip (offering a link to a search engine for each piece), as well as the points for each round and the total score for the match (Figure 2).

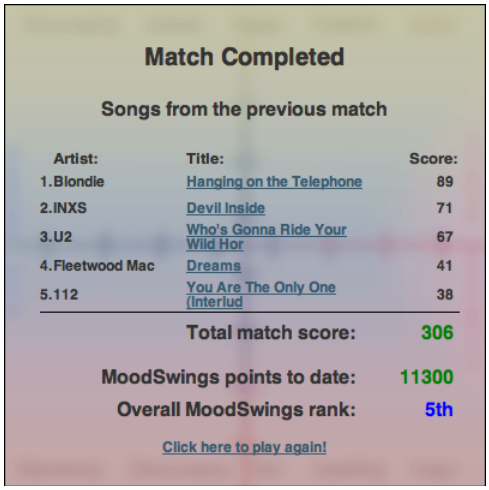

Figure 2. The MoodSwings post-match wrapup

3.2 Game scoring

The primary score calculation is based upon the amount of overlap between the player's cursor and that of their partner, which is designed to encourage maximum agreement in the mood assessments from both parties. This score is only accumulated whenever the partner's cursor is visible. When points accumulate, they are displayed on top of the partner's cursor (shown in Figure 1).

Players can also accumulate *bonus points* by "convincing" their partner to agree with a particular mood assessment (position). This provides an incentive for players to remain firm in a position and not to be capricious in their movements. It also encourages them to respond to a change in mood rapidly in order to "stake out" the new position before their partner. The rules for bonus points are as follows:

- A player is eligible to accumulate bonus points after remaining stationary for 1 second. This is indicated by a yellow square around the player's cursor. Bonus points will only be awarded while the player remains stationary.
- If a partner moves towards a player's location achieving overlap between the cursors, the stationary player

is awarded 5 bonus points. This is indicated by the cursor changing color to green (Figure 3).

- Bonus points may be awarded every second, even when the partner's cursor is not visible to the player.

The overall point system is designed so that a "good" score for a round is approximately 100 points, and a particularly good match total is 500 points. High scores for the top five players as well as the top five individual match scores are visible when first logging onto the MoodSwings website.

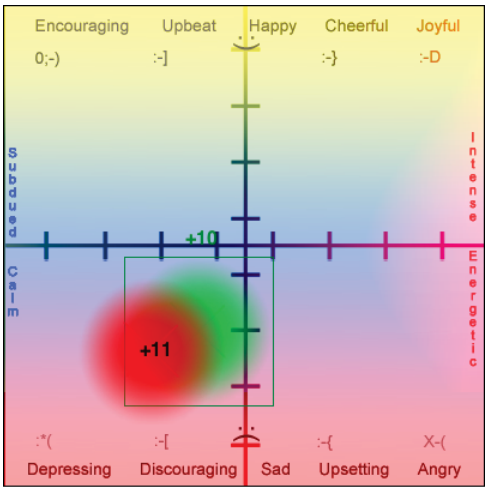

Figure 3. MoodSwings bonus scoring

3.3 Music database

The music for MoodSwings is drawn randomly from the well-known uspop2002 database, a collection of over 8000 popular music tracks from approximately 400 performing artists [18]. This database was chosen because of the composition of the corpus (popular music spanning several decades) and its potential appeal to a mass audience, the size of the database, and the fact that it includes a range of familiar and obscure tunes. In addition, since the corpus is well-known, a great deal of acoustic feature data and music metadata have already been calculated and compiled for the database, such as MFCCs [19]. This will allow other researchers to rapidly and easily deploy their algorithms using the labels collected from the game.

The uspop2002 database, however, is not without its issues. A fair number of the songs contain explicit lyrics, which may be objectionable to some players. Some of the clips randomly selected for the game will not include music at all because of extended applause from live recordings and other unusual spoken-word tracks from a few albums. Because such metadata (explicit lyrics, non-music sections) for tracks is not readily available, the game interface also offers players the opportunity to mark such tracks. In particular, beneath the game board for each round are the following options that a player may voluntarily select:

- *Clip does not contain music:* for selections that do not contain music (so that we can filter these tracks out in the future).
- *Song contains explicit lyrics:* this will help us create a future version of the game appropriate for all ages that excludes these songs.
- *Report a bug in this game:* for any other problem encountered during the current round.

3.4 Technical implementation

A primary goal during the development of MoodSwings was to allow access to the game through a standard web browser interface, so that the game would be widely accessible and would not require the installation of any additional software. In particular, our lab works closely with several K-12 school environments where the computers may be out of date and additional software is difficult to install.

The end-user interface for MoodSwings is principally coded using Asynchronous JavaScript and XML (AJAX), which also makes use of Dynamic HTML. Audio playback is handled through the Flash plug-in, which is included with all browsers. The browser portion communicates with a web server hosted by our laboratory that is used to synchronize players, record the collected mood assessments, and administer the game. A diagram of the overall architecture of the system is given in Figure 4.

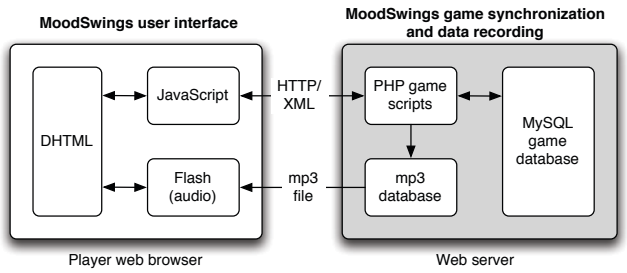

Figure 4. MoodSwings game architecture for a single player. Each player interacts with the server independently.

The web server uses scripts written in PHP to respond to client requests for game data, such as music clip information and partner mood coordinates, in XML format. The game data is stored in a MySQL database running on the server that tracks all of the following information:

- Valence and arousal assessments from each player for each second of the song.
- Game and match scores.

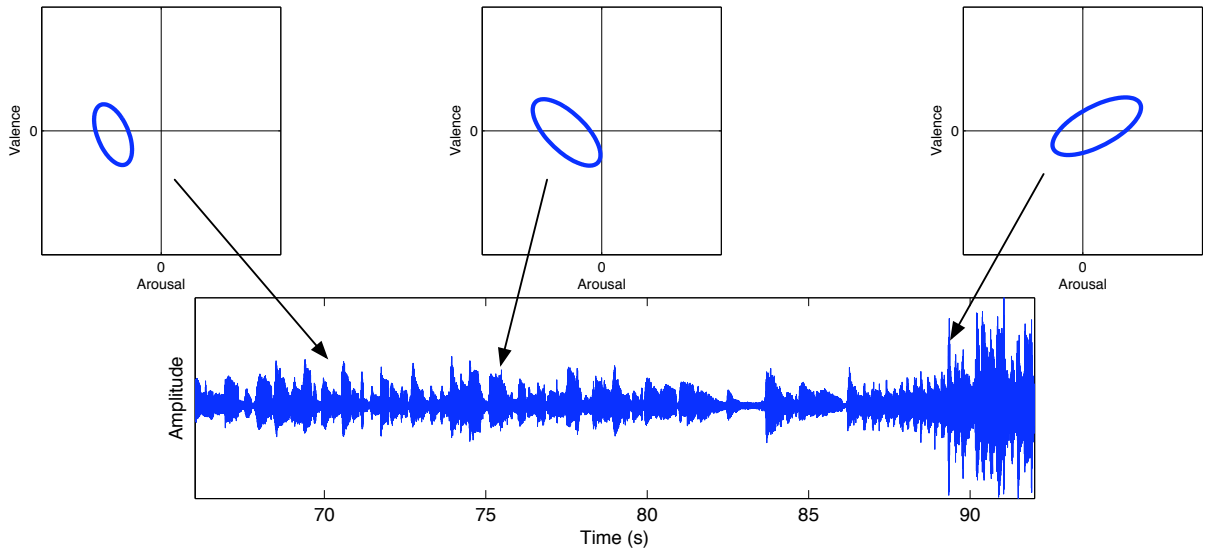

Figure 5. Progression of valence-arousal labels over a clip spanning the end of the first chorus and beginning of the second verse of "American Pie" (Don McLean). The ellipses represent the standard deviation across different players.

- Game and match parameters (song clips used, clip starting locations).
- Music metadata (artist, album, song title).
- User scores and high scores.

3.4.1 Technical limitations and single-player matches

Obviously, with simultaneous listening between the players there will be some latency in the data received from one's partner during a game. Since the mood assessments are only sampled once per second, it is likely that the partner's data received by a player will be at least one second out of date. In practice, this does not affect gameplay very much, since a player normally takes a second or two to react to sudden mood changes.

Even when other players are not available online, a player may still play a match against data previously recorded from another player's match. No distinction is made to the player to indicate a match against a live player vs. a pre-recorded match. In some ways, such single-player matches against recorded data may be preferred because the partner's labels can be pre-fetched so that synchronization between ratings and time samples is not an issue.

4 MOOD DATA COLLECTION

In our initial one week pilot phase, we attracted approximately 100 users and collected over 50,000 valence-arousal point labels spanning more than 1000 songs. Of course, given the collaborative structure of the game, many of the valence-arousal labels refer to the same locations in a song.

4.1 Example data from song clip in database

Figure 5 shows a sample of collected valence-arousal labels for a short section of the song "American Pie" by Don McLean, between the first chorus and second verse. The change in instrumentation and tempo within the segment (drums are added to voice and guitar) is generally marked by players as a change in intensity in the song, as well as a slight increase in valence.

4.2 Summary of labels collected to date

Figure 6 depicts the distribution of the collected points in the valence-arousal space. It is clear that the songs labeled thus far have bias towards high valence and arousal, which is consistent with a database of popular music. The plot also shows that players for the most part favor locations near the middle of each "quadrant" in the valence-arousal space, largely avoiding extreme values.

5 FUTURE WORK

The initial implementation of MoodSwings and the preliminary data collected thus far suggest the potential of such a collaborative system with many users in providing high-quality labels. We are currently investigating modifications to make the gameplay more fun, with the hope of drawing a greater number of repeat visits from users. A question arises in examining live matches vs. those played against recorded data. Qualitatively, the labels collected in single and two-player games appear equally valid, but we plan to verify their consistency by more closely examining the data.

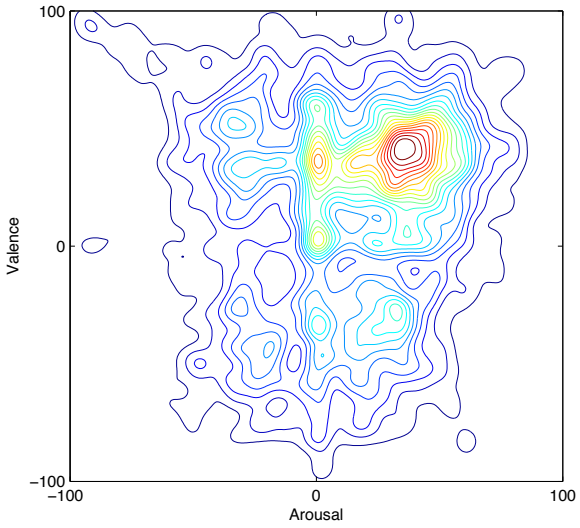

Figure 6. Contour plot of the distribution of ∼50,000 valence-arousal labels collected over one-week pilot period.

An issue in dealing with popular music is that the mood of most songs simply doesn't change very much or often, which can lead to rather static games (perhaps this is a comment on the state of popular music!). We plan to add an option for additional simultaneous players (3 or even more) that may produce more interesting group labeling dynamics. We also plan on augmenting the uspop2002 databse with a collection of classical music clips, which will have the added benefit of removing some of the lyric vs. audio confusion that arises in labeling some pop songs.

As our collection of labels grows, we intend to use the data to train a system for mood classification of short audio segments using a Hidden Markov Model to capture time-varying mood transitions. Once the entire database has been labeled, the data collected from MoodSwings will be made available to the music information retrieval research community through the game website.

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ON THE USE OF SPARSE TIME-RELATIVE AUDITORY CODES FOR MUSIC

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ABSTRACT

Many if not most audio features used in MIR research are inspired by work done in speech recognition and are variations on the spectrogram. Recently, much attention has been given to new representations of audio that are sparse and time-relative. These representations are efficient and able to avoid the time-frequency trade-off of a spectrogram. Yet little work with music streams has been conducted and these features remain mostly unused in the MIR community. In this paper we further explore the use of these features for musical signals. In particular, we investigate their use on realistic music examples (i.e. released commercial music) and their use as input features for supervised learning. Furthermore, we identify three specific issues related to these features which will need to be further addressed in order to obtain the full benefit for MIR applications.

1 INTRODUCTION

The majority of the features used in audio-related MIR research are based on Fourier analysis, which suffers from two weaknesses. The first is the trade-off in precision between time and frequency. The second, common to all block based representations, is a sensitivity to arbitrary alignment of the blocks with the musical events.

Sparse coding assumes a signal can be represented at a given point in time by a rather small number of basis functions taken from an overcomplete dictionary [9]. Recent work [2, 10, 13, 14] applies these ideas to audio streams. When set in the time domain, the result is a spikegram, an efficient representation of the signal that avoids both of the spectrogram's weaknesses. Figure 1 shows a given signal (an ascending and descending C-major scale played on a piano), its spectrogram and its spikegram. As can be seen, a spikegram is composed of a set of spikes, corresponding to the placement of a basis function (kernel) at a precise point in time and with a particular scaling coefficient. The spikegram encodes audio very efficiently and with arbitrary resolution along both axes.

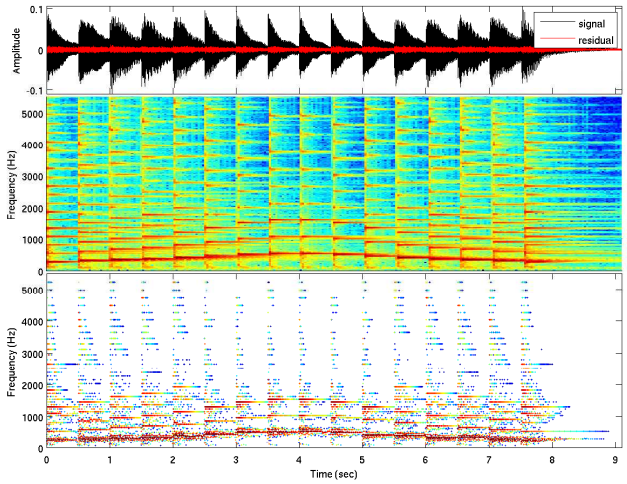

Figure 1. **Top:** original signal (an ascending and descending C-major scale played on the piano) and its residual (after encoding the signal to 20 dB SNR using the auditory codes). **Middle:** spectrogram. **Bottom:** spikegram. The horizontal axis represents time with the same resolution as that of the signal. The vertical axis corresponds to the frequencies of the n kernels used (Gammatones in this case). A dot represents a spike, a kernel placed at a specific point in time. The size and color of the dot are representative of the scaling coefficient of the kernel. This spikegram contains $13,000$ values, enough to encode the signal to 20dB SNR.

The organization of the paper is as follows. Section 2 reviews existing work on sparse coding for audio streams. Section 3 presents the sparse coding algorithm we use. Section 4 attempts to bring insight into the resulting codes. To this end, we encode the Tzanetakis genre dataset using predefined Gammatone kernels. In Section 5 the features are applied in a naive way to the task of genre recognition and are shown to work as well as other commonly used audio features. In section 6 we attempt to learn a set of kernels better suited to music than the general purpose Gammatones. The learned kernels are qualitatively different from those learned on other types of sounds. This suggests music poses specific coding challenges. Finally, in Section 7,

we conclude and identify three specific issues with sparse coding for further research.

2 RELATED WORK

In this section we present existing work on creating a sparse encoding of audio signals. First, we present work done in the frequency domain, and secondly, methods in the time domain.

2.1 Frequency Domain

Plumbley and Abdallah have many publications about sparse coding using a code book. See [10] for one of their recent articles. The main idea is to derive a dictionary of power spectra from a corpus of audio. One assumes that the spectra of the signal is a weighted sum of the dictionary elements. Weights can be derived in number of ways. For example in [10] a method employing non-negative matrix factorization and one employing variance estimation are used. In the same article, the authors apply their coding to the task of note detection. However, the most sparse representation they present is from a time domain method, discussed further in Section 2.2.

Another recent approach is introduced by Gardner and Magnesco [5]. The idea is to compute a transformation of the spectrogram that gives it higher precision. Thus it is possible to track a single signal efficiently.

2.2 Time Domain

The most used method in the time domain is based on a dictionary of kernels that are mapped onto a signal using a MAP estimate [2, 10, 13]. MAP gives the best reconstruction error, but is very computationally expensive. Thus, an approximation is developed in [2], and in [13] MAP is used as an optimization phase using only a subset of selected kernels.

Two other methods of encoding are described by Smith and Lewicki [13]. The first one uses a threshold on the correlation between the signal and the kernels; the second one uses matching pursuit. In experiments, matching pursuit comes out as the best compromise between computational complexity and Signal to Noise Ratio (SNR).

The kernels used can be either predefined or learned using a gradient ascent method. The first option is used by Smith and Lewicki in [13]. They use Gammatones, which are known to approximate the cochlear filters. The second option requires a corpus on which to learn the kernels. In [10], 57 kernels are learned on a recording of Beethoven's Bagatelle containing 57 played notes. Thus they expect each kernel to model a particular note. In [14], learning is done on environmental sounds (ambient and transient), mammalian vocalizations, speech and reversed speech.

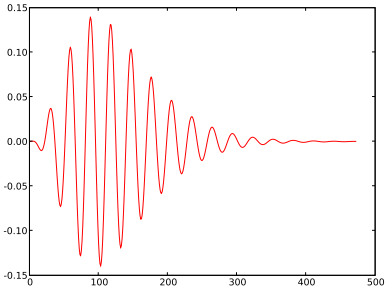

Figure 2. Example of a Gammatone kernel.

3 SPARSE TIME-RELATIVE CODING

In this section we describe the model we use for signal representation as well as the encoding algorithm. The model is the same as Smith and Lewicki's [13] and is used in conjunction with the matching pursuit encoding, as done in [13, 14].

3.1 Model for signal representation

We represent a signal using a sparse representation based on shiftable kernels [13]. This models the signal x as a linear superposition of kernels ϕ that are placed at precise time locations τ with a given scaling coefficient s. Formally:

$$x(t) = \sum_{m=1}^{M} \sum_{i=1}^{n_m} s_i^m \phi_m(t - \tau_i^m) + \epsilon(t),$$

where m runs over the different kernels, i over the different instances of a given kernel and ϵ represents additive noise. The length of the kernels is variable.

3.2 Choice of encoding algorithm

Encoding a signal amounts to determining the placement and scaling of kernels. This is a non-linear process and finding the optimal sparse representation using a generic dictionary of functions is NP-hard [3]. We choose to use matching pursuit [7], an iterative greedy algorithm, which is shown to offer a very good trade-off between computational resources and efficiency of representation [13]. The algorithm works iteratively in three steps. First, the signal is cross-correlated with the kernels. The best fitting projection is then selected and the corresponding kernel identity, placement and scaling are recorded. The projection is then subtracted from the signal and the procedure is repeated over the residual.

4 ENCODING USING GAMMATONE KERNELS

In this section, we attempt to bring insight into the encoding. Basic characteristics and properties are illustrated through

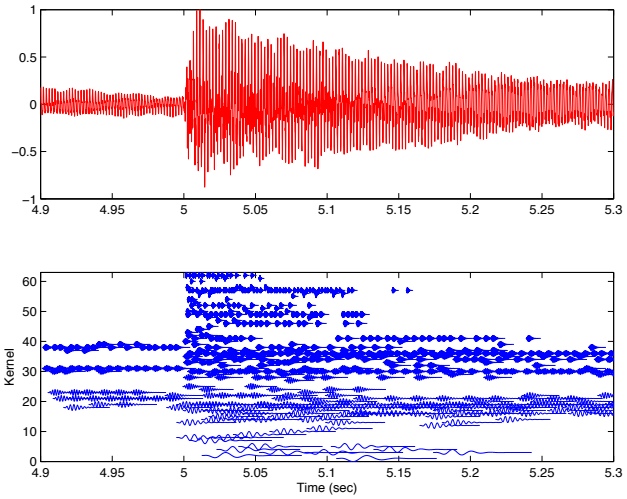

Figure 3. **Top:** original signal (close-up of a piano sounding an A4). **Bottom:** spikegram (using kernel shapes).

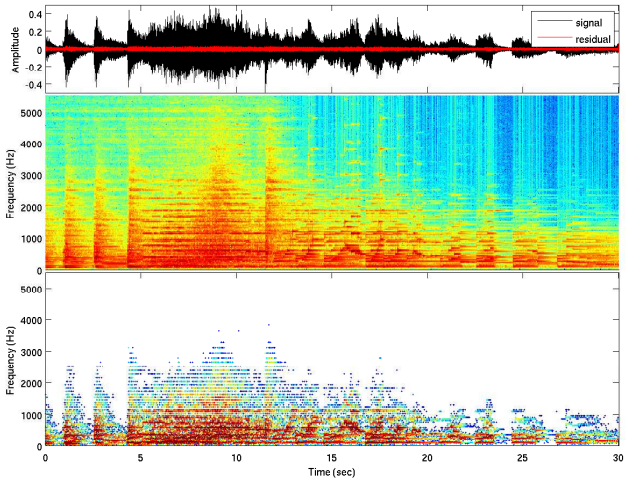

Figure 4. **Top:** original signal (30 seconds from a jazz song) and its residual (after encoding the signal to 20dB SNR using the auditory codes). **Middle:** spectrogram. **Bottom:** spikegram.

the encoding of the Tzanetakis genre dataset. We use Gammatone kernels (see Figure 2), which are motivated as biological general purpose kernels used in the cochlea. Their characteristic sharp rise and slow decay also intuitively fit many real phenomenons. The Gammatones are set according to an equivalent rectangular band (ERB) filter bank cochlear model using Slaney's auditory toolbox for Matlab [12], as done in [13]. Unless otherwise stated, we use 64 variable length normalized Gammatone kernels, with frequencies ranging from 20Hz to the Nyquist frequency.

An important characteristic of the coding is that kernels are placed at precise time locations. This characteristic allows for precise localisation of events. This is illustrated in Figure 3, which shows the encoding of a note played on a piano. The onset clearly stands out.

Another important property of the coding is the fact that spikes are only used on a per-need basis. In Figure 4 we see that most of the spikes are placed in the middle of the song where the energy is. This contrasts with the uniform distribution of encoding resources in the spectrogram and illustrates a characteristic of sparse coding: it is adaptive in the number of spikes for a given encoding ratio. Figure 5 shows the encoding of a metal song. Though the code is still sparse, more spikes are needed than for the jazz example, perhaps due to the constant presence of high-frequency guitar with heavy distortion.

We further investigate the previous result, i.e. that jazz seems easier to encode than metal. For this purpose we used Tzanetakis' genre dataset later presented in Section 5. In top of Figure 6 we show the average number of spikes per song needed to encode songs from different musical genres to a given SNR. Some genres seem to require more spikes, metal being the most needy. We placed an upper limit on the number of spikes computed (10^5), which is why the metal box is compressed. To make sure the difficulty lies in the high frequencies, we divide the 64 kernels in eight bins and count them separately. The results are at the bottom of Figure 6. The highest frequencies are on the leftt, the lowest on the right. As expected, many high frequency Gammatones were used for metal songs, whereas classical or jazz songs required few.

# of kernels	10^3	10^4	2.5×10^4	5×10^4
32	2.82 (0.97)	8.57 (2.17)	12.69 (2.87)	16.60 (2.88)
64	3.02 (1.03)	9.07 (2.29)	13.44 (2.98)	17.43 (2,70)
128	3.08 (1.04)	9.24 (2.31)	13.76 (2.99)	17.76 (2.52)

Table 1. Mean (standard deviation) of the SNR (dB) obtained over 100 songs (ten from each genre), as a function of the number of Gammatones in the codebook and of the maximum number of spikes allowed to reach a maximal value of 20 dB SNR.

It is also interesting to determine the effect on the signal to noise ratio of using different numbers of kernels in the ERB and different numbers of spikes in the spikegram. The results are presented in Table 1. Surprisingly, the number of kernels has relatively little impact for the tested range of values.

5 GENRE RECOGNITION

In order to explore the discriminative ability of the spikegram representation, we apply it to the classification task of genre recognition. As classifier, we use the AdaBoost meta-learning algorithm. This choice is motivated by the fact that source code [8] is available as well as published results for a set of genre recognition experiments [1]. Those experiments

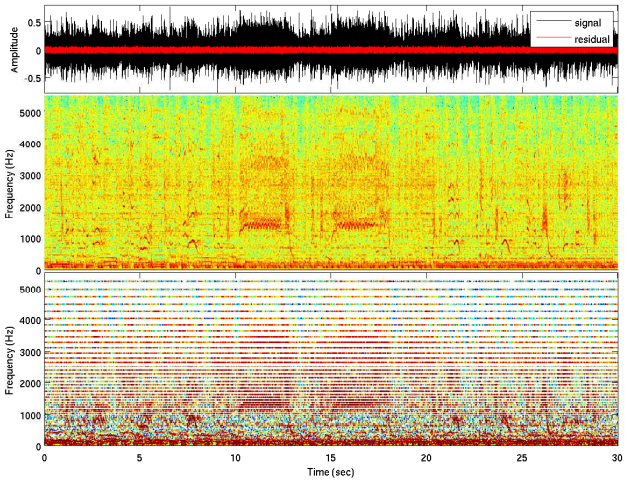

Figure 5. **Top:** original signal (30 seconds from a metal song) and its residual (after encoding the signal to 20dB SNR using the auditory codes). **Middle:** spectrogram. **Bottom:** spikegram.

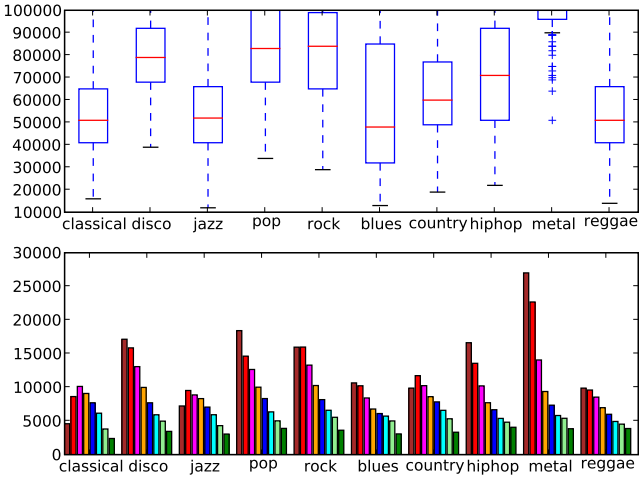

Figure 6. **Top:** box-and-whisker plot of the number of spikes (out of a maximum of 10^5) required to encode songs from various genres up to a SNR of 20dB. The red line represents the median and the box extends from the lower quartile to the upper one. **Bottom:** Bar plot of the mean number of kernels used per song depending on the genre. The 64 kernels are split into eight bins going from the highest frequency Gammatones (left) to the lowest ones (right).

were performed on a freely available dataset from Tzanetakis. (Similar datasets were used for previous MIREX contests. However those datasets are not available for download.)

The Tzanetakis genre dataset contains one thousand 30-second songs evenly distributed among 10 genres and is used in several papers [1, 16, 15, 6]. The dataset is very small and is annotated with only a single winner-take-all genre label. Though we believe the dataset is suitable for exploring the predictive power of the spikegram, it is clearly too small to reflect the challenges faced in finding structure in large datasets of audio.

5.1 Algorithm

AdaBoost [4] is a *meta-learning* method that constructs a *strong classifier* from a set of simpler classifiers, called *weak learners* in an iterative way. Originally intended for binary classification, there exist several ways to extend it to multiclass classification. We use AdaBoost.MH [11] which treats multiclass classification as a set of one-versus-all binary classification problems. In each iteration t, the algorithm selects the best classifier, called $h^{(t)}$ from a pool of *weak learners*, based on its performance on the training set, and assigns it a coefficient $\alpha^{(t)}$. The input of the *weak learner* is a d-dimensional observation vector $x \in \Re^d$ containing audio features for one segment of data (5 seconds in our experiments). The output of $h^{(t)}$ is a binary vector $y \in \{-1, 1\}^k$ over the k classes. $h_l^{(t)} = 1$ means a vote for class l by a *weak learner* while $h^{(t)}$, -1 is a vote against. After T iterations, the algorithm output is a vector-valued discriminant function:

$$g(x) = \sum_{t=1}^{T} \alpha^{(t)} h^{(y)}(x) \qquad (1)$$

We obtain a single label by taking the class with the "most votes" i.e $f(x) = \arg\max_l g_l(x)$. As *weak learner* we use *single stumps*, a simple threshold on one element of the feature vector.

We use an AdaBoost method for genre recognition based on a method detailed in Bergstra et al. [1].

5.2 Features from the spikegram

The input of the classifier needs to be a feature vector of fixed size for every segment of a song (5 seconds segments here). Of course, a sparse code is designed to adapt to specific audio signals. The number of spikes per segment changes, and the spikes can be positioned at any given music frame. Thus we have encode the information about the spikes differently in order to use it as input to a typical classifier.

Here we try two types of features based on the spikes. First, we simply count the number of times each spike is used in every segment, and we also sum their scaling coefficients s. We also give the same information, but normalized by the number of spikes in the segment. Finally, we give the classifier the total number of spikes in this segment. For 64 kernels, this yields an input vector of size 257 per segment.

feat. SC	feat. SC + MFCC	MFCC (B.)
63.0%	68.2%	63%

Table 2. Genre recognition results on Tzanetakis dataset with 5-fold cross-validation using the sparse code features (SC), the sparse code features and our implementation of MFCC (SC + MFCC) and result using MFCC from Bergstra et al. [1] (B.). Result from [1] is approximate (taken from a graph).

5.3 Results

Results are reported using a 5-fold cross-validation. For more details about reported results on this dataset in the literature, see Bergstra et al. [1].

Using AdaBoost.MH with single stumps as weak learners, and 5-second segments, Bergstra et al. [1] report an error per song of about 63% using MFCCs. Our results using the features derived from the spikegrams are presented in Table 2. We see that even with a naive use of the spikegram, results are comparable with commonly used audio features. The spikegrams also add information to the MFCC as can be seen in Column 2 of Table 2.

6 LEARNING KERNELS

The spikegrams used in the previous sections were based on the use of kernels of a specific predefined form (Gammatones). It is also possible to learn the kernels as done in [10, 14] over different audio corpora. Here we investigate the learning of kernels for western commercial music. There are at least three motivations for this. First, we hope to better encode music with learned kernels than with Gammatones. Secondly, these kernels should be more efficient as input for MIR tasks. Thirdly, perhaps these learned kernels can form meaningful blocks representing concepts like compression, note onsets or timber. This was the case in Plumbley et al. [10] where the authors could relate the learned kernels to piano notes. The difference here is that we train on more complex music and we put less constraint on the kernel lengths.

6.1 Algorithm

The learning is done by gradient ascent. The approximate gradient of a signal x, given a kernel ϕ, is (from [14]):

$$\frac{\partial}{\partial \phi} p(x|\phi) = \frac{1}{\sigma_\epsilon} \sum_i \hat{s}_i [x - \hat{x}]_{\tau_i} \qquad (2)$$

We can then perform gradient ascent on every dimension of the kernels using the residual signal at corresponding positions. The σ_ϵ is constant and can be ignored as we set the learning rate empirically.

In our experiments, 32 kernels are initialized using random values drawn from a Gaussian distribution. Throughout the learning, the kernels are kept normalized. We use a database containing tens of thousands of MP3s. Learning proceeds by iteratively randomly choosing a song from the database, encoding it using 1000 spikes, computing the residual signal and performing gradient ascent on the kernels. The learning rate was set to 0.01 and we did approximately 2000 iterations. Figure 7 shows samples from the 32 kernels we learned. The learned kernels are qualitatively different from Gammatones. They closely resemble those learned by [10]. However, our kernels are of variable length which may be better suited to encoding music with varying note durations.

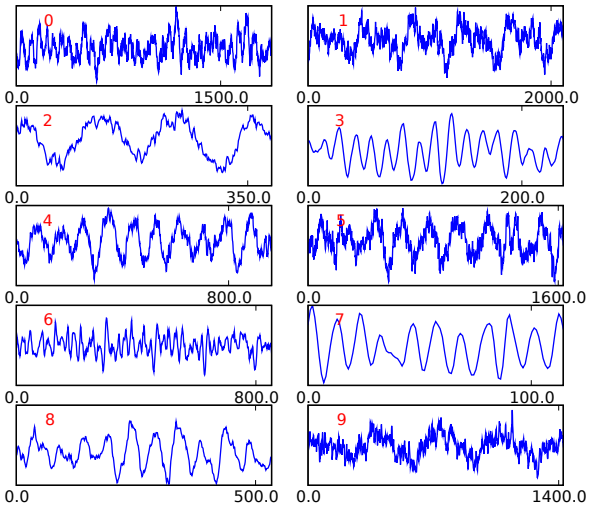

Figure 7. Examples of learned kernels

kernel	10^3	10^4	2.5×10^4	5×10^4
gammatones	2.82 (0.97)	8.57 (2.17)	12.69 (2.87)	16.60 (2.88)
learned	1.86 (0.57)	6.06 (1.49)	8.86 (2.00)	11.53 (2,37)

Table 3. Mean (standard deviation) of the SNR (dB) obtained over 100 songs (ten from each genre) with either 32 Gammatones or 32 learned kernels in the codebook.

6.2 Learned Kernels versus Gammatones

As a first experiment, we evaluate how well the learned kernels encode songs compared to the Gammatones. Results are shown in Table 3. Unfortunately, Gammatones perform significantly better at this task. This is a surprising result, as the learned kernels are trained to optimize this specific cost, suggesting that learning kernels for complex music poses specific challenges.

As a second experiment, we use the 32 learned kernels as input for genre recognition, and we compare the results with those obtained using 32 Gammatones. Results are shown in Table 4. We compare the results when encoding using 10^3

gamma (1K)	learn (1K)	gamma(10K)	learn (10K)
47.5 %	46.3 %	52.5 %	54.3 %

Table 4. Genre recognition results on Tzanetakis dataset with 5-fold cross-validation. Number in parentheses is the maximum number of spikes used to encode each song.

and 10^4 spikes for computational reasons. Learned kernels seem to perform similarly as Gammatones. This shows that the encoding capacity of kernels is different from their predictive property for a particular task.

7 DISCUSSION AND FUTURE WORK

Our goal with this work is to draw further attention to time-relative sparse codings. These codes manage to avoid the weaknesses of the spectrogram and encode an audio signal very efficiently. Yet there are three issues related to their use that will need to be addressed. The first is how to properly use them as input to MIR tasks. Indeed, in our genre recognition experiments, we summarize spike counts and scaling coefficients over a block. This destroys the useful time-relative configurations of the spikes and reintroduces a sensitivity to arbitrary block alignment. The fact that we are still able to get results comparable to MFCCs shows the potential of the features. The second issue relates to learning in the case of complex music. Our work seems to reveal specific difficulties. In particular, kernels trained over music tended to continually grow, and thus we had to enforce some length constraints. The disappointing results are also indicative that the learning problem becomes difficult with complex music. Solutions might lie in using weight decays, and starting from kernels trained on less complex signals. Finally, there is an issue with the computational complexity of the technique. Even with the use of matching pursuit, the computational requirements of encoding are above real time, which may not suit all applications. Answers may rely in performing approximate matching pursuit, or learning a function to do an approximate encoding using machine learning.

8 ACKNOWLEDGEMENTS

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THE 2007 MIREX AUDIO MOOD CLASSIFICATION TASK: LESSONS LEARNED

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ABSTRACT

Recent music information retrieval (MIR) research pays increasing attention to music classification based on moods expressed by music pieces. The first Audio Mood Classification (AMC) evaluation task was held in the 2007 running of the Music Information Retrieval Evaluation eXchange (MIREX). This paper describes important issues in setting up the task, including dataset construction and ground-truth labeling, and analyzes human assessments on the audio dataset, as well as system performances from various angles. Interesting findings include system performance differences with regard to mood clusters and the levels of agreement amongst human judgments regarding mood labeling. Based on these analyses, we summarize experiences learned from the first community scale evaluation of the AMC task and propose recommendations for future AMC and similar evaluation tasks.

1. INTRODUCTION

With the goal of systematically evaluating state-of-the-art algorithms for Music Information Retrieval (MIR) systems, the Annual Music Information Retrieval Evaluation eXchange (MIREX) included an Audio Mood Classification (AMC) task for the first time in 2007. It is inspired by MIR researchers' growing interest in classifying music by moods (e.g. [4][7][8]), and the difficulty in the evaluation of music mood classification caused by the subjective nature of mood. Most previous experiments on music mood classification used different mood categories and datasets, raising a great challenge in comparing systems. MIREX, as the largest evaluation event in the MIR community, is a good venue to build an available audio dataset and ground-truth for AMC and to facilitate collaborations among MIR researchers around the world.

The first AMC task in MIREX was a success. A ground-truth set of 600 tracks distributed across five mood categories was built based on metadata analysis and human assessments. A total of nine systems from Europe and North America participated in the task. Resultant accuracies ranged from 25.67% to 61.50%, with an average of 52.65%, a median of 55.83% and a standard deviation of 11.19%.

In this paper, we examine the evaluation process and explore the datasets in detail. We also analyze the system performances with a special focus on the possible effects that the data creation and evaluation process might have. The findings will help in organizing similar MIREX-style evaluations in the future.

The rest of the paper is organized as follows: Section 2 describes critical issues in preparing and carrying out the AMC evaluation task. Section 3 analyzes the human assessments made on candidate audio pieces for the purpose of building a ground-truth set. Statistical analyses on system performances from multiple angles are presented in Section 4, and Section 5 wraps up with discussions and recommendations based on the findings from the analyses.

2. EVALUATION SETUP

2.1. Dataset

A standard dataset used in evaluating a classification task should include a set of categories, a collection of samples distributed across these categories and ground-truth labels that ideally are given by agreements among multiple human judges. The AMC task adopted the set of five mood clusters proposed in [6] which effectively reduce the mood space into a manageable set. For clarity purposes, we present the mood clusters in Table 1. The words in each cluster collectively define the "mood spaces" associated with the cluster.

Cluster1	Cluster2	Cluster3	Cluster4	Cluster5
Rowdy	Amiable/	Literate	Witty	Volatile
Rousing	Good natured	Wistful	Humorous	Fiery
Confident	Sweet	Bittersweet	Whimsical	Visceral
Boisterous	Fun	Autumnal	Wry	Aggressive
Passionate	Rollicking	Brooding	Campy	Tense/anxious
	Cheerful	Poignant	Quirky	Intense
			Silly	

Table 1. Five mood clusters used in the AMC task

2.1.1. Audio Track Collection

Our previous study [6] shows that music mood, as a descriptive metadata type, is independent of music genre. Thus, it is desirable to evaluate mood classification algorithms against a collection of music pieces in a variety of genres. Furthermore, the ideal collection should have not been used in previous evaluations so as to avoid the overfitting problem caused by repetitive uses of the same collection. Keeping these criteria in mind, we chose the libraries of Associated Production Music (APM) as the candidate pool. The APM collection is "the world's leading production music library… offering every imaginable music genre from beautiful classical music recordings to vintage rock to current indie band sounds"[1]. This collection is made available to the MIR community under a contract of academic use between APM and the IMIRSEL lab. It contains 206,851 tracks pooled from 19 libraries produced by APM and covers 27 genres. Each track is annotated with rich, descriptive metadata including instrument, tempo, style, etc. One such metadata field is called "category" which contains a list of descriptors including 32 mood-related ones (e.g. "Moods-Aggressive", "Moods-Quirky"). Such descriptors are particularly useful in building the evaluation dataset using the following steps:

Step 1. Eliminate tracks without genre information, so that we can explicitly select tracks in diversified genres;

Step 2. A set of rules are manually designed according to statistics of the mood-related descriptors, and tracks matching the rules are selected and pre-labeled with the mood clusters specified by the rules. An exemplar rule is "if 'Moods-Quirky' ∈ Song1.category, then Song1 ∈ Cluster 4";

Step 3. Eliminate tracks shorter than 50 seconds, as a means to reduce the chances of including non-music content in the extracted 30 seconds clips (see below);

Step 4. Eliminate tracks from the same CDs or same libraries, for diversity purposes.

The result is a collection of 1250 tracks with 250 pieces in each mood cluster. These tracks were then truncated into 30-second clips and were taken as candidates to be judged by human evaluators.

The choice of using 30-second clips over whole tracks is due to several considerations. First, it lessens the burden on our human evaluators. Second, it reduces the runtime needed in audio processing. Finally, it alleviates variation caused by the changing nature of music mood, i.e., some pieces may start from one mood but end in another. The clips are extracted from the middle of the tracks which are assumed to be more representative than other parts.

2.1.2. Ground-truth and Human Assessment

Our goal was to build a ground-truth set of 600 clips with 120 in each mood cluster. These numbers were decided by polling opinions from potential participants via the AMC task wiki [2]. We planned to have multiple human assessments on each of the 1250 candidates selected by the metadata statistics. Based on a pilot experiment on selecting exemplar songs, we estimated at least half of the candidates would achieve majority agreements on their mood labels among human assessments. We used the Evalutron 6000 [5], a Web-based device designed for collecting human judgments for MIREX evaluations. As music mood is very subjective and judging music mood is a new task for human evaluation, we designed concise but explicit instructions to help control variations among the human assessors. The most important instructions include:

1) Ignoring lyrics. Many clips have lyrics which often express certain moods and thus could affect listeners' judgments. However, state-of-the-art audio music processing technology has not yet been developed to sufficiently transcribe lyrics. Hence, we should try our best to mitigate possible bias imposed by lyrics.

2) Mini-training on exemplar songs. In order to better articulate what the mood clusters mean, we prepared a set of exemplar songs in each mood cluster that were unanimously judged by 6 IMIRSEL members. We added special settings to the Evalutron to ensure an assessor cannot complete registration until finishing listening to the excerpts of the exemplar songs.

3) "Other" category. A human assessor can change the pre-selected label on a candidate piece if she does not agree with it. To better capture assessors' opinions, we designed an "Other" category for the cases when none of the five clusters seems appropriate to the assessors.

2.2. Evaluation Method

As in many other evaluations on classification tasks, the submitted systems were trained, tested and evaluated using cross-validation. In particular, the AMC task was evaluated using 3-fold cross-validation. As a starting point for evaluating on music mood classification, the AMC task was defined as a single-label classification problem, i.e., each song can only be classified into one mood cluster. The classification results were evaluated and compared using classification accuracy as the measure. There is also an alternative evaluation approach proposed during task preparation that is discussed in Section 5. The audio format used in the task was 22KHz mono-channel wav files, as voted on by the majority of the potential participants on the AMC task wiki.

[1] www.apmmusic.com/pages/aboutapm.html

[2] http://www.music-ir.org/mirex/2007/index.php/AMC

3. HUMAN ASSESSMENT ANALYSIS

3.1. Statistics on the Evalutron Data

Human assessment data were collected between 1 Aug. and 19 Aug., 2007, from volunteer assessors from the MIR/MDL research community, representing 5 different countries from Europe and the U.S. Among the 21 volunteers registered on the Evalutron, 15 of them provided at least one judgment and 8 of them finished assessing all assigned 250 clips while each of the remaining assessors completed 6 to 140 assessments.

Table 2 shows the number of human agreements for clips with at least two judgments for each of the mood clusters (denoted as "C1" to "C5"), produced by early September 2007. To build the ground-truth, we needed clips with at least two agreed judgments. As can be seen from the table, there were not enough such clips (< 120) for Cluster 1, 2 and 4. As a means to quickly compensate for the missing judgments, the IMIRSEL lab collected clips with only one judgment in these clusters and organized an in-house assessment on these clips.

	# of clips with 3 or 2 agreements	# of clips with no agreement	Total
C1	102 (57.6%)	75 (43.4%)	177
C2	105 (64.0%)	59 (36.0%)	164
C3	147 (79.9%)	37 (20.1%)	184
C4	95 (60.5%)	62 (39.5%)	157
C5	137 (75.3%)	45 (24.7%)	182
Total	586 (67.8%)	278 (32.2%)	864

Table 2. Agreements on clips with 3 or 2 judgments

The fact that Cluster 3 and 5 had enough agreed judgments and also allowed the highest agreement rate (> 70%) reflects that the clips pre-selected in these two clusters caused fewer confusions among human assessors. This possibly means the pre-labeling methods worked better for Cluster 3 and 5 than Cluster 1 (where agreement rate was the lowest) and/or that Cluster 3 and 5 were easier to assess than Cluster 1. In Section 4, we will see the participating systems performed better in Cluster 3 and 5 as well.

3.2. Effects on Reclassification

As human assessors were asked to change the pre-assigned labels whenever appropriate, it is interesting to see how aggregated human opinions differ from pre-assigned labels. We only consider pieces with at least two agreed human judgments. Table 3 shows how the agreed assessments concurred with or changed the pre-assigned labels. For example, among the clips pre-assigned to Cluster 1, 59.8% were kept in Cluster 1 by agreed human judgments while 40.2% of them were reclassified to other clusters including 17.6% to the "other" category (far higher than other mood clusters). For other clusters, more than 84% of the clips kept their pre-assigned labels. Cluster 1 seemed to have caused the greatest confusion with other clusters, according to the human assessments.

	C1	C2	C3	C4	C5	Other
C1	59.8%	11.8%	17.6%	6.9%	0.0%	17.6%
C2	1.0%	89.5%	5.7%	1.0%	0.0%	2.9%
C3	2.0%	1.4%	92.5%	0.7%	0.7%	2.7%
C4	0.0%	7.4%	1.1%	84.2%	2.1%	5.3%
C5	4.4%	0.0%	1.5%	1.5%	89.8%	2.9%

Table 3. Percentage distribution of agreed judgments (rows: pre-assigned labels, columns: human assessment)

3.3. Three Judgment Sets

Among the 600 audio pieces eventually used in the AMC task, 153 of them were agreed upon by 3 human judges (denoted as "Set 1"), 134 were assessed by 3 judges but only 2 of the judges reached agreement ("Set 2"), and 313 pieces were assessed by 2 judges who agreed on the mood label assignments ("Set 3"). Table 4 demonstrates the distribution of clips in these sets across mood clusters. In a later analysis (Section 4.4), we will investigate if the number of agreed judgments made a difference in system performances.

	C1	C2	C3	C4	C5	Total
Set 1	21	24	56	21	31	153
Set 2	41	35	18	26	14	134
Set 3	58	61	46	73	75	313
Total	120	120	120	120	120	600

Table 4. Number of audio pieces in different judgment sets across mood clusters

4. SYSTEM PERFORMANCE ANALYSIS

Nine systems participated in the AMC 2007 task. The average classification accuracy scores over a 3-fold cross-validation, as well as run time information were published on the MIREX results wiki[1]. Table 5 presents the accuracy results ($\in [0, 1]$) for each system broken down by fold, along with its overall average accuracy.

In this section, we examine the performance results from several different angles. We are interested in knowing the answers to the following questions:

1. Which system(s) performed better than the others in a statistically significant way? Is there any difference between systems a) across folds; or, b) across mood clusters?
2. Are there significant performance differences between the folds or between the mood clusters?

[1] http://www.music-ir.org/mirex/2007/index.php/MIREX2007_Results

3. Does the division of ground-truth into judgment Set 1, 2 and 3 (Section 3.1) have an effect on performance?

	GT	CL	TL	ME1	ME2	IM2	KL1	IM1	KL2	Avg.
Fold 1	0.70	0.66	0.67	0.63	0.62	0.58	0.52	0.51	0.22	0.57
Fold 2	0.56	0.59	0.58	0.57	0.51	0.55	0.52	0.45	0.26	0.51
Fold 3	0.58	0.57	0.56	0.53	0.55	0.54	0.46	0.46	0.29	0.50
Avg.	0.61	0.61	0.60	0.58	0.56	0.56	0.50	0.47	0.26	0.53

Legend: GT: George Tzanetakis; CL: Cyril Laurier and Perfecto Herrera; TL: Thomas Lidy, Andreas Rauber, Antonio Pertusa and José Manuel Iñesta; ME1, ME2: Michael I. Mandel and Daniel P. W. Ellis; IM1, IM2: IMIRSEL M2K; KL1, KL2: Kyogu Lee

Table 5. System accuracies across folds

4.1. Comments on Statistical Testing

Exploratory data analyses of the result datasets consistently indicated that these data failed to meet the necessary assumptions of parametric statistical tests in that they exhibit non-normal distributions and non-equal variances. Therefore, we adopted the non-parametric Friedman's ANOVA test [1] as our general tool to determine if significant differences were present among the groups of interest (e.g., among systems, among mood clusters, among judgment sets, etc.). If and only if a Friedman's overall test showed the presence of a statistically significant difference (at $p < 0.05$) somewhere among the groups, would we then turn to the Tukey-Kramer Honestly Significantly Different (TK-HSD) analyses [1] to determine where the significant differences actually existed among the groups. The TK-HSD is used rather than the commonly mis-used multiple *t*-tests because TK-HSD properly controls for the experiment-wise Type I error rate whereas the naïve adoption of multiple *t*-tests does not. Failure to adjust for this experiment-wise inflation in Type I error in situations such as ours, where many pair-wise comparisons are made, all but guarantees that the null hypotheses of no differences in means between pairs of interest (i.e., H_0: μ (rank x) = μ (rank y) will be falsely rejected somewhere within the set of comparisons being made.

4.2. System Performance Comparison

To explore question 1 outlined above, we looked at the results from two different viewpoints corresponding to questions 1.a (fold-based) and 1.b (cluster-based). In test 1.a, we ran the Friedman's test using the accuracies shown in Table 5. In test 1.b we used accuracy scores in Table 6. Both tests proved significant (1.a: χ^2 (8, 16) = 21.42, $p < 0.01$; 1.b: χ^2 (8, 32) = 15.61; $p = 0.048$).

We then conducted the follow-up TK-HSD analysis on the two sets, to see exactly how the system performances differed. The results are displayed in Table 7 (for set 1.a) and Table 8 (for set 1.b). The shaded areas in the tables represent the groups of systems which did not show significant differences within each group.

	C1	C2	C3	C4	C5	All
GT	0.43 (7)	0.53 (1)	0.80 (3)	0.52 (4)	0.80 (1)	0.62 (1)
CL	0.46 (5)	0.50 (2)	0.83 (2)	0.53 (2)	0.71 (4)	0.61(2)
TL	0.53 (2)	0.49 (3)	0.75 (4)	0.53 (3)	0.69 (6)	0.60 (3)
ME	0.52 (3)	0.46 (4)	0.70 (5)	0.55 (1)	0.67 (7)	0.58 (4)
IM_2	0.51 (4)	0.45 (5)	0.68 (7)	0.45 (6)	0.70 (5)	0.56 (5)
ME2	0.55 (1)	0.43 (6)	0.65 (8)	0.51(5)	0.66 (8)	0.56 (5)
KL1	0.25 (8)	0.37 (7)	0.84 (1)	0.25 (8)	0.78 (3)	0.50 (7)
IM_1	0.45 (6)	0.22 (9)	0.70 (5)	0.19 (9)	0.80 (1)	0.47 (8)
KL2	0.15 (9)	0.25(8)	0.33 (9)	0.27 (7)	0.28 (9)	0.26 (9)

Table 6. Accuracies cross mood clusters. Numbers in parenthesis are ranks within each cluster.

	GT	CL	TL	ME1	ME2	IM2	KL1	IM1	KL2
Group 1									
Group 2									

Table 7. System groups: fold-based performances

	CL	GT	TL	ME1	ME2	IM2	KL1	IM1	KL2
Group 1									
Group 2									

Table 8. System groups: cluster-based performances

The fact that KL2 is not grouped together with GT and CL for set 1.a (Table 7) indicates the two system pairs (GT, KL2) and (CL, KL2) are significantly different in the fold-based analysis. It is noteworthy that among all systems, CL has the best ranks across all mood clusters despite its average accuracy being the second highest. As the TK-HSD is based on ranks rather than the raw scores, CL is part of the only pair with difference in set 1.b, the cluster-based analysis.

4.3. Effects of Folds and Mood Clusters

In order to see whether there were any significant differences among folds or mood clusters, we transposed the datasets used in test 1.a and 1.b and conducted Friedmans' tests on the transposed sets. Again, both tests were significant: test 2.a: χ^2 (2, 16) = 6.22, $p < 0.01$; test 2.b: χ^2 (4, 32) = 27.91; $p < 0.01$).

The follow-up TK-HSD analysis showed that Fold 1 and Fold 3 were significantly different. In general, using more folds would help alleviate the impact of one fold on the overall performances. In regard to the five clusters, there were two pairs of difference: (Cluster 3, Cluster 1) and (Cluster 5, Cluster 1). This is consistent with what we saw in human assessment analysis: Cluster 3 and 5 reached the best agreements among assessors and best performances among systems while Cluster 1 caused the most confusion both among assessors and systems.

4.4. System Performance and Human Agreement

In this section, we investigate whether the number of agreed judgments on the audio pieces affects system performances. We calculated accuracy scores on each of the three judgment sets described in Section 3.1 and present them in Table 9.

	GT	CL	TL	ME1	ME2	IM2	KL1	IM1	KL2	Avg.
Set 1	0.72	0.67	0.67	0.67	0.65	0.66	0.54	0.46	0.28	0.59
Set 2	0.41	0.43	0.37	0.4	0.39	0.47	0.39	0.36	0.19	0.38
Set 3	0.64	0.62	0.65	0.59	0.56	0.55	0.49	0.51	0.26	0.54

Table 9. System accuracies across three judgment sets

From the table, we can see that across all systems, accuracies on Set 1 are consistently the highest and those on Set 2 are always the lowest. A Friedman's test on the scores in Table 9 indicated the existence of significant difference ($\chi 2$ (2, 16) = 16.22, $p < 0.01$). The follow-up TK-HSD analysis showed two pairs of significant difference: (Set 1, Set 2) and (Set 3, Set 2). This means that the systems performed significantly better on Set 1 and Set 3 than on Set 2. Set 2 consisted of audio pieces involving disagreement among human judges while the other two sets contained only agreed judgments. This suggests pieces with discrepant human judgments impose greater challenges to the systems. In order to reduce such inherent ambiguity in the dataset, our recommendation for future evaluations is to exclude pieces with mixed labels when three judgments are collected for each piece.

Now we investigate the interactions between the 3 judgment sets and the 3 folds in cross-validation. The breakdown of the sets and folds is shown in Table 10. We can see that Fold 1 was dominated by Set 1, the set with best system performances, while Fold 3 contained none of the Set 1 pieces[1]. Also, Fold 1 included fewer pieces from the problematic Set 2 than Fold 3 did. These factors at least partially explain the observation that systems performed significantly better on Fold 1 than Fold 3. Again, this shows that the unanimity among human assessors on the dataset had an important influence on system performances, and suggests that one way to reduce variations among folds is to stratify audio pieces with different numbers of human agreements when splitting the dataset into training and testing sets.

	Set 1	Set 2	Set 3	Total
Fold 1	116	22	62	200
Fold 2	37	47	116	200
Fold 3	0	65	135	200
Total	153	134	313	600

Table 10. Number of audio pieces across judgment sets and cross-validation folds

[1] The folds were split according to the result order of MySql queries on the collected human assessments. The impact of the levels of human agreements was not recognized until after the evaluation.

5. DISCUSSION AND RECOMMENDATIONS

5.1. Techniques Used in Systems

Comparing the abstracts of the submissions available on the task result wiki page, we found that most of the systems adopted Support Vector Machines (SVM) as the classifier[2], but with different implementations, including the libsvm library [2], SMO algorithm [9] implemented in the WEKA machine learning software [12], and DAG-SVM for efficient n-way classification [10]. Table 11 shows the systems grouped by the classifiers they used, as well as the average accuracy within each group.

	Weka SMO	LibSVM	DAG-SVM	KNN
System(s)	IM2, TL	CL, GT	ME1, ME2	IM1
Avg. Acc.	0.58	0.61	0.57	0.47

Table 11. System groups w.r.t. classifiers (KNN signifies a K-nearest neighbor classification model)

The systems utilized a variety of audio and symbolic features, but spectral features were included in all systems. Among them, GT and ME2 were exclusively based on spectral features. ME1 added temporal features to ME2 and improved the overall accuracy by 2%. CL utilized fairly rich music descriptors including temporal, tonal, loudness and high level features (e.g., danceability) while TL added symbolic features capturing pitches and note durations. According to the analysis in Section 4, there were no significant differences among these systems. Thus, the (high-level) features other than the basic spectral ones did not show any advantage in this evaluation. One possible reason is that the training set is too small (for each fold, only 80 training clips exist in each mood cluster) to optimize models with a large feature space. In future AMC tasks, a larger training set is desirable for allowing possible improvements by using large feature spaces.

5.2. Evaluation Approaches

During the evaluation setup process, two evaluation approaches were proposed. One was to evaluate on a closed dataset with ground-truth labels, as adopted in AMC 2007. The advantages of this approach include rigorous evaluation metrics and the ability to conduct cross-validation. However, since labeling ground-truth is human labor intensive, both the sizes of training and testing sets are limited in this approach. The other proposed approach was to train systems on a labeled dataset, and subsequently test them on an unlabeled audio pool. After each system returns a ranked list of song candidates for each mood cluster, human assessors make judgments only on the top ranked candidates. This

[2] As abstracts of KL1 and KL2 were not published on the task result wiki page, we do not know what techniques were used in these systems.

approach adopts the pooling mechanism first proposed in TREC, the annual evaluation event in the domain of text-focused information retrieval [11], and has been used in the Audio Similarity and Retrieval task in both MIREX 2006 and 2007. The obvious advantage is that the size of the testing set can be arbitrarily large because the number of judgments required is independent of the size of the test set. However, one downside of this approach is that traditional evaluation metrics (e.g. precision, recall, F-measure, accuracy) can only be used after careful modifications. For example, one cannot measure the absolute "recall" metrics, but can only compare systems using "relative recall" scores calculated by assuming all unjudged samples are irrelevant [11].

The AMC 2007 participants chose the first approach by voting on it: 8 voted for the first approach while only 1 voted for the second exclusively. However, 4 voters indicated they would prefer to have their systems evaluated using both approaches. Considering the second approach's advantage of being able to test systems on a large scale dataset, we recommend adopting this approach in future AMC tasks and perhaps in other classification tasks as well.

5.3. Recommendations on Task Organization

Based on the analysis on the three judgment sets, we recommend that future AMC tasks exclude pieces where judges disagree. This would reduce ambiguity in the dataset and help measure the systems more accurately.

The methods of acquiring more ground-truth of high quality need to be validated and improved in the future. Formal studies are needed to guide choices such as whether to provide human assessors with pre-selected labels and exemplar songs.

During the debates on how the evaluation should be done, the polls conducted through the AMC wiki drew votes from potential participants and helped make critical decisions. We recommend this polling method as it can collect opinions directly from participants in a timely manner, and can clearly present the distribution of all votes. In fact, 6 out of the 11 tasks in MIREX 2007 used polls to aid decision making.

6. ACKNOWLEDGEMENTS

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INSTRUMENT EQUALIZER FOR QUERY-BY-EXAMPLE RETRIEVAL: IMPROVING SOUND SOURCE SEPARATION BASED ON INTEGRATED HARMONIC AND INHARMONIC MODELS

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ABSTRACT

This paper describes a music remixing interface, called *Instrument Equalizer*, that allows users to control the volume of each instrument part within existing audio recordings in real time. Although query-by-example retrieval systems need a user to prepare favorite examples (songs) in general, our interface gives a user to generate examples from existing ones by cutting or boosting some instrument/vocal parts, resulting in a variety of retrieved results. To change the volume, all instrument parts are separated from the input sound mixture using the corresponding standard MIDI file. For the separation, we used an integrated tone (timbre) model consisting of harmonic and inharmonic models that are initialized with template sounds recorded from a MIDI sound generator. The remaining but critical problem here is to deal with various performance styles and instrument bodies that are not given in the template sounds. To solve this problem, we train probabilistic distributions of timbre features by using various sounds. By adding a new constraint of maximizing the likelihood of timbre features extracted from each tone model, we succeeded in estimating model parameters that better express actual timbre.

1 INTRODUCTION

One of promising approaches of music information retrieval is the query-by-example (QBE) retrieval [1, 2, 3, 4, 5, 6, 7] where a user can receive the list of musical pieces ranked by their similarity to a musical piece (example) that the user gives as a query. Although this approach is powerful and useful, a user has to prepare or find favorite examples and sometimes feels difficulty to control/change the retrieved pieces after seeing them because the user has to find another appropriate example to get better results. For example, if a user feels that vocal or drum sounds are too strong in the retrieved pieces, the user has to find another piece that has weaker vocal or drum sounds while keeping the basic mood and timbre of the piece. It is sometimes very difficult to find such a piece within a music collection.

We therefore propose yet another way of preparing an example for the QBE retrieval by using a music remixing interface. The interface enables a user to boost or cut the volume of each instrument part of an existing musical piece. With this interface, a user can easily give an alternative query with a different mixing balance to obtain refined results of the QBE retrieval. The issue in the above example of finding another piece with weaker vocal or drum sounds can thus be resolved. Note that existing graphic equalizers or tone controls on the market cannot control each individual instrument part in this way: they can adjust only frequency characteristics (e.g., boost or cut for bass and treble). Although remixing stereo audio signals [8] had reported previously, it had tackled to control only harmonic instrument sounds. Our goal is to control all instrument sounds including both harmonic and inharmonic ones.

This paper describes our music remixing interface, called *Instrument Equalizer*, in which a user can listen to and remix a musical piece in real time. It has sliders corresponding to different musical instruments and enables a user to manipulate the volume of each instrument part in polyphonic audio signals. Since this interface is independent of the succeeding QBE system, any QBE system can be used. In our current implementation, it leverages the standard MIDI file (SMF) corresponding to the audio signal of a musical piece to separate sound sources. We can assume that it is relatively easy to obtain such SMFs from the web, etc. (especially for classical music). Of course, given a SMF, it is quite easy to control the volume of instrument parts during the SMF playback, and readers might think that we can use it as a query. Its sound quality, however, is not good in general and users would lose their drive to use the QBE retrieval. Moreover, we believe it is important to start from an existing favorite musical piece of high quality and then refine the retrieved results.

2 INSTRUMENT EQUALIZER

The Instrument Equalizer enables a user to remix existing polyphonic musical signals. The screenshot of its interface is shown in Figure 1 and the overall system is shown in Figure 2. It has two features for remixing audio mixtures as follows:

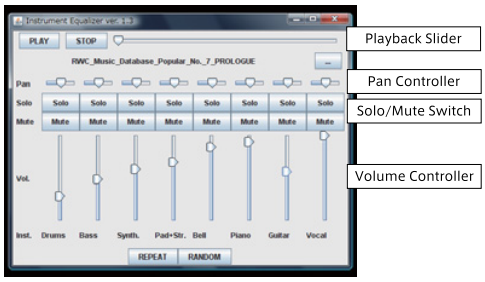

Figure 1. Screenshot of main window.

Figure 2. Instrument Equalizing System.

1. *Volume control function*. It provides the remixing function by boosting or cutting the volume of each instrument part, not by controlling the gain of a frequency band. A user can listen to the remixed sound mixture as soon as the user manipulates the volume.
2. *Interlocking with the hardware controller*. In addition to a typical mouse control on the screen, we allow a user to use a hardware controller shown in Figure 2 with multiple faders. It enables the user to manipulate the volume intuitively and quickly. This hardware controller makes it easy to manipulate the volume of multiple parts at the same time, while it is difficult on a mouse control.

To remix a polyphonic musical signal, the signal must be separated into each instrument part. We use an integrated weighted mixture model consisting harmonic-structure and inharmonic-structure tone models [9] for separating the signal, but improve the parameter estimation method of this model by introducing better prior distributions. This separation method needs a standard MIDI file (SMF) that is synchronized to the polyphonic signal. We assume that the SMF has already been synchronized with the input signal by using audio-to-score alignment methods such as [10, 11, 12]. For the separation using the integrated model, the parameters of the model are initialized by template sounds recorded from a MIDI sound generator. and gradually improved to represent actual sounds in the sound mixture.

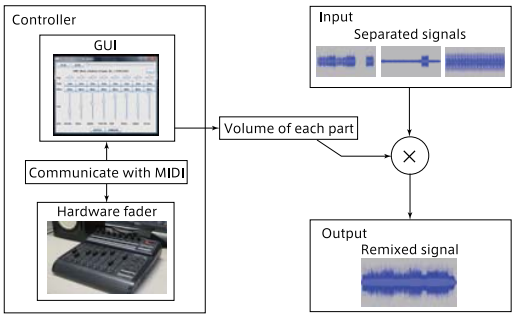

Figure 3. System architecture.

2.1 Internal architectures

This section describes the internal architectures of controlling the volume of each instrument part. The procedures described in this section are performed in real time under the assumption that the musical signals of each instrument part already have been obtained in advance from the target polyphonic musical signal, as described in Section 3. Let $x_k(c,t)$ and $y_k(c,t)$ be a separated signal and the volume of instrument k at channel c and time t, respectively. $y_k(c,t)$ satisfies the following condition:

$$\forall k, c, t : 0 \leq y_k(c,t) \leq 1,$$

and $y_k(c,t)$ is obtained as

$$y_k(c,t) = (\text{value of volume slider } k) \cdot (\text{value of the pan } c).$$

The overview of the architecture is shown in Figure 3.

1. *Volume control function*. The output signal, $x(c,t)$, is obtained as

$$x(c,t) = \sum_k y_k(c,t) \cdot x_k(c,t).$$

   Each $y_k(t)$ is obtained in real-time from the volume sliders in the GUI in Figure 1.
2. *Interlocking with the hardware controller*. The GUI and the hardware controller communicate by MIDI. If users control the hardware fader, a MIDI message which represents the new volume is sent to the GUI, and vice varsa. Since a motor is embeded in the fader, MIDI messages from the GUI move the fader to the position corresponding value of the volume.

3 SOUND SOURCE SEPARATION CONSIDERING TIMBRE VARIETIES

In this section, we first define our sound source separation problem and the integrated model. We then describe timbre varieties and timbre feature distributions for estimating parameters of the model.

3.1 Integrated model of harmonic and inharmonic models

The sound source separation problem is to decompose the input power spectrogram, $X(c,t,f)$, into the power spectrogram corresponding to each musical note, where c, t, and f are the channel (e.g., left and right), the time, and the frequency, respectively. We assume that $X(c,t,f)$ includes K musical instruments and the k-th instrument performs L_k musical notes. We use the tone model, $J_{kl}(c,t,f)$, to represent the power spectrogram of the l-th musical note performed by the k-th musical instrument (*(k,l)-th note*), and the power spectrogram of a template sound, $Y_{kl}(t,f)$, to initialize the parameters of $J_{kl}(c,t,f)$. Each musical note of the SMF is played back on a MIDI sound generator to record the corresponding template sound. $Y_{kl}(t,f)$ is monaural because SMFs may not include any sound localization (channel) information. $Y_{kl}(t,f)$ is normalized to satisfy the following relation, where C is the total number of the channels:

$$\sum_c \iint X(c,t,f)\,dt\,df = C \sum_{k,l} \iint Y_{kl}(t,f)\,dt\,df.$$

For this source separation, we define an integrated model, $J_{kl}(c,t,f)$, as the sum of harmonic-structure tone models, $H_{kl}(t,f)$, and inharmonic-structure tone models, $I_{kl}(t,f)$, multiplied by the whole amplitude of the model, $w_{kl}^{(J)}$, and the relative amplitude of each channel, $r_{kl}(c)$:

$$J_{kl}(c,t,f) = w_{kl}^{(J)} r_{kl}(c)\big(H_{kl}(t,f) + I_{kl}(t,f)\big),$$

where $w_{kl}^{(J)}$ and $r_{kl}(c)$ satisfy the following constraints:

$$\sum_{k,l} w_{kl}^{(J)} = \iint X(c,t,f)\,dt\,df,\ \forall k,l:\ \sum_c r_{kl}(c) = C.$$

All parameters of $J_{kl}(c,t,f)$ are listed in Table 1. The harmonic model, $H_{kl}(t,f)$, is defined as a constrained two-dimensional Gaussian mixture model (GMM), which is a product of two one-dimensional GMMs, $\sum E_{kl}^{(H)}(m,t)$ and $\sum F_{kl}^{(H)}(n,t,f)$, and is designed by referring to the harmonic-temporal-structured clustering (HTC) source model [13]. The inharmonic model, $I_{kl}(t,f)$, is defined as a product of two nonparametric functions. The definition of these models is as follows:

$$H_{kl}(t,f) = w_{kl}^{(H)} \sum_{m=1}^{M}\sum_{n=1}^{N} E_{kl}^{(H)}(m,t) F_{kl}^{(H)}(n,t,f),$$

$$E_{kl}^{(H)}(m,t) = \frac{u_{kl}(m)}{\sqrt{2\pi}\phi_{kl}} \exp\left(-\frac{(t-\tau_{kl}-m\phi_{kl})^2}{2\phi_{kl}^2}\right),$$

$$F_{kl}^{(H)}(n,t,f) = \frac{v_{kl}(n)}{\sqrt{2\pi}\sigma_{kl}} \exp\left(-\frac{(f-n\omega_{kl}(t))^2}{2\sigma_{kl}^2}\right), \text{and}$$

$$I_{kl}(t,f) = w_{kl}^{(I)} E_{kl}^{(I)}(t) F_{kl}^{(I)}(t,f),$$

Table 1. Parameters of the integrated model.

Symbol	Description
$w_{kl}^{(J)}$	overall amplitude
$r_{kl}(c)$	relative amplitude of each channel
$w_{kl}^{(H)}$, $w_{kl}^{(I)}$	relative amplitude of harmonic and inharmonic tone models
$u_{kl}(m)$	coefficient of the temporal power envelope
$v_{kl}(n)$	relative amplitude of n-th harmonic component
τ_{kl}	onset time
ϕ_{kl}	diffusion of a Gaussian of power envelope
$\omega_{kl}(t)$	F0 trajectory
σ_{kl}	diffusion of a harmonic component along the freq. axis
$E_{kl}^{(I)}(t)$	power envelope of inharmonic tone model
$F_{kl}^{(I)}(t,f)$	relative amplitude of frequency f at time t of inharmonic tone model

where M is the number of Gaussian kernels representing the temporal power envelope and N is the number of Gaussian kernels representing the harmonic components. $u_{kl}(m)$, $v_{kl}(n)$, $E_{kl}^{(I)}(t)$, $F_{kl}^{(I)}(t,f)$, $w_{kl}^{(H)}$, and $w_{kl}^{(I)}$ satisfy the following conditions:

$$\forall k,l:\ \sum_m u_{kl}(m) = 1, \quad \forall k,l:\ \sum_n v_{kl}(n) = 1,$$

$$\forall k,l:\ \int E_{kl}^{(I)}(t)\,dt = 1, \quad \forall k,l,t:\ \int F_{kl}^{(I)}(t,f)\,df = 1,$$

$$\text{and} \quad \forall k,l:\ w_{kl}^{(H)} + w_{kl}^{(I)} = 1.$$

The goal of this separation is to decompose $X(c,t,f)$ into $J_{kl}(c,t,f)$ by estimating a spectrogram distribution function, $\Delta^{(J)}(k,l;c,t,f)$, which satisfies

$$\forall k,l,c,t,f:\ 0 \le \Delta^{(J)}(k,l;c,t,f) \le 1 \quad \text{and}$$
$$\forall c,t,f:\ \sum_{k,l} \Delta^{(J)}(k,l;c,t,f) = 1.$$

With $\Delta^{(J)}(k,l;c,t,f)$, the separated power spectrogram, $X_{kl}^{(J)}(c,t,f)$, is obtained as

$$X_{kl}^{(J)}(c,t,f) = \Delta^{(J)}(k,l;c,t,f) X(c,t,f).$$

Furthermore, let $\Delta^{(H)}(m,n;k,l,t,f)$ and $\Delta^{(I)}(k,l,t,f)$ be spectrogram distribution functions which decompose $X_{kl}^{(J)}(c,t,f)$ into each Gaussian distribution of the harmonic model and the inharmonic model, respectively. These functions satisfy

$$\forall k,l,m,n,t,f:\ 0 \le \Delta^{(H)}(m,n;k,l,t,f) \le 1,$$
$$\forall k,l,t,f:\ 0 \le \Delta^{(I)}(k,l,t,f) \le 1, \quad \text{and}$$
$$\forall k,l,t,f:\ \sum_{m,n} \Delta^{(H)}(m,n;k,l,t,f) + \Delta^{(I)}(k,l,t,f) = 1.$$

To evaluate the 'effectiveness' of this separation, we can use a cost function defined as the Kullback-Leibler (KL) divergence from $X_{kl}^{(J)}(c,t,f)$ to $J_{kl}(c,t,f)$:

$$Q_{kl}^{(J)} = \sum_c \iint X_{kl}^{(J)}(c,t,f) \log \frac{X_{kl}^{(J)}(c,t,f)}{J_{kl}(c,t,f)}\, dt\, df.$$

By minimizing the sum of $Q_{kl}^{(J)}$ over (k,l) pertaining to $\Delta^{(J)}(k,l;c,t,f)$, we obtain the spectrogram distribution function and model parameters (i.e., the most 'effective' decomposition).

By minimizing the $Q_{kl}^{(J)}$ pertaining to each parameter of the integrated model, we obtain model parameters estimated from the distributed spectrogram. This parameter estimation is equivalent to a maximum likelihood estimation. The parameter update equations are described in appendix A.

3.2 Timbre varieties within each instrument

Even within the same instrument, different instrument bodies have different timbres, although its timbral difference is smaller than the difference among different musical instruments. Moreover, in live performances, each musical note could have slightly different timbre according to the performance styles. Instead of preparing a set of many template sounds to represent such timbre varieties within each instrument, we represent them by using a probabilistic distribution.

We use parameters of the integrated model, $u_{kl}(m)$, $v_{kl}(n)$, and $F_{kl}^{(I)}(t,f)$, to represent the timbre variety of instrument k by training a diagonal Gaussian distribution with mean $\mu_k^{(u)}(m)$, $\mu_k^{(v)}(m)$, $\mu_k^{(F)}(f)$ and variance $\Sigma_k^{(u)}(m)$, $\Sigma_k^{(v)}(n)$, $\Sigma_k^{(F)}(f)$, respectively. Note that other probability distributions, such as a Dirichlet distribution, are available in this case. The model parameters for training the prior distribution are extracted from instrument sound database [14] (i.e., the parameters are estimated without any prior distributions).

By minimizing the cost function,

$$\begin{aligned} Q_{kl}^{(p)} =& \sum_c \iint X_{kl}^{(J)}(c,t,f) \log \frac{X_{kl}^{(J)}(c,t,f)}{J_{kl}(c,t,f)}\, dt\, df \\ &+ \frac{1}{2}\sum_m (u_{kl}(m) - \mu_k^{(u)}(m))^2/\Sigma_k^{(u)}(m) \\ &+ \frac{1}{2}\sum_n (v_{kl}(n) - \mu_k^{(v)}(n))^2/\Sigma_k^{(v)}(n) \\ &+ \frac{1}{2}\iint (F_{kl}^{(I)}(t,f) - \mu_k^{(F)}(f))^2/\Sigma_k^{(F)}(f)\, dt\, df, \end{aligned}$$

where the last term is an additional cost by using the prior distribution, we obtain the parameters by taking into account the timbre varieties. This parameter estimation is equivalent to a maximum *A Posteriori* estimation.

3.3 Cost function without considering timbre feature distributions

In Itoyama's previous study [9], they used template sounds instead of timbre feature distributions to evaluate the 'goodness' of the feature vector. The cost function, $Q_{kl}^{(Y)}$, used in [9] can be obtained by replacing the negative log-likelihood, $Q_{kl}^{(p)}$, with the KL divergence, $Q_{kl}^{(Y)}$, from $Y_{kl}(t,f)$ (the power spectrogram of a template sound) to $J'_{kl}(t,f)$:

$$\begin{aligned} Q_{kl}^{(Y)} =& \sum_c \iint X_{kl}^{(J)}(c,t,f) \log \frac{X_{kl}^{(J)}(c,t,f)}{J_{kl}(c,t,f)}\, dt\, df \\ &+ \sum_c \iint r_{kl}(c) Y_{kl}(t,f) \log \frac{r_{kl}(c) Y_{kl}(t,f)}{J_{kl}(c,t,f)}\, dt\, df. \end{aligned}$$

4 EXPERIMENTAL EVALUATION

We conducted experiments to confirm whether the performance of the source separation using the prior distribution is equivalent to the one using the template sounds. In the first experiment, we separated the sound mixtures which were generated from a MIDI sound generator. In the other one, the sound mixtures were created from the signals with multiple tracks [15] which were before mixdown. In this experiment, we compared the following two conditions:

1. using the log-likelihood of timbre feature distributions (proposed method, section 3.2),
2. using the template sounds (previous method [9], section 3.3).

4.1 Experimental conditions

We used 5 SMFs from the RWC Music Database (RWC-MDB-P-2001 No. 1, 2, 3, 8, and 10) [16]. We recorded all musical notes of these SMFs by using two different MIDI sound generators made by different manufacturers. We used one of them for the test (evaluation) data and the other for obtaining the template sounds or training the timbre feature distributions in advance.

The experimental procedure is as follows:

1. initialize the integrated model of each musical note by using the corresponding template sound,
2. estimate all the model parameters from the input sound mixture, and
3. calculate the SNR in the frequency domain for the evaluation.

The SNR is defined as follows:

$$\mathrm{SNR} = \frac{1}{C(T_1 - T_0)} \sum_c \int \mathrm{SNR}_{kl}(c,t)\, dt,$$

$$\mathrm{SNR}_{kl}(c,t) = \log_{10} \int \frac{X_{kl}^{(J)}(c,t,f)^2}{\left(X_{kl}^{(J)}(c,t,f) - X_{kl}^{(R)}(c,t,f)\right)^2}\, df,$$

Table 2. Experimental conditions.

Frequency analysis	Sampling rate	44.1 kHz
	Analyzing method	STFT
	STFT window	2048 points Gaussian
	STFT shift	441 points
Parameters	C	2
	M	10
	N	30
MIDI sound generator	Test data	YAMAHA MU-2000
	Template sounds	Roland SD-90

Table 3. SNRs of the signals separated from sound mixtures generated from a MIDI tone generator. P1, P2, and P3 are the SNRs which are based on the prior distribution trained using 1, 2, and 3 instrument bodies, respectively. T is the SNR which is based on the template sounds.

	P1	P2	P3	T
P001	11.5	12.1	11.6	14.0
P002	12.3	12.3	12.5	12.3
P003	11.5	11.8	12.4	10.8
P008	8.1	7.8	8.3	4.9
P010	9.1	8.8	8.9	12.2
Ave.	10.5	10.6	10.8	10.8

Table 4. SNRs of the signals separated from sound mixtures generated from CD recordings.

	P1	P2	P3	T
P001	9.3	9.4	9.4	9.0
P002	11.4	11.5	11.7	11.6
P003	4.0	4.1	4.2	3.1
P008	7.1	7.2	7.3	6.1
P010	8.4	8.5	8.5	8.0
Ave.	8.1	8.2	8.3	7.6

where T_0 and T_1 are the beginning and ending times of the input power spectrogram, $X(c,t,f)$, and $X_{kl}^{(R)}(c,t,f)$ is the ground-truth power spectrogram corresponding to the (k,l)-th note (i.e., the spectrogram of an actual sound before mixing). We used a 40-parameters for the prior distributions (1–10 dimensions: $u_{kl}(1)$, ..., $u_{kl}(M)$, 11–40 dimensions: $v_{kl}(1)$, ..., $v_{kl}(N)$), where M = 10 and N = 30. Other experimental conditions are shown in Table 2.

4.2 Experimental results

The results are listed in Tables 3 and 4. In both experiments, the SNRs were improved by increasing the number of instrument bodies for training the prior distributions. Furthermore, the average SNR of P3 is equal to that of T in Table 3 and the average SNRs in Table 4 is improved from the average SNR of T. This means that although the SNRs decrease in some cases where the timbre difference between template sounds and input sounds is large, it can be resolved by using the better prior distributions.

5 CONCLUSION

In this paper, we have proposed the novel use of a music remixing interface for generating queries for the QBE retrieval, explained our Instrument Equalizer on the basis of sound source separation using an integrated model consisting of harmonic and inharmonic models, and described a new parameter estimation method for the integrated model by using the timbre feature distributions. We confirmed that this method increased separation performance for most instrument parts simply by using the basic timbre features.

Although the use of the timbre feature distributions is promising, it has not been fully exploited in our experiments. For example, we have not tried to use training data including various performance styles and instrument bodies. We plan to evaluate our method by using such various training data as well as more advanced timbre features. Some performance benchmark for audio source separation [17] will helpful to compare our separation method with other ones. Future work will also include the usability evaluation of the Instrument Equalizer for the use of the QBE retrieval.

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A DERIVATION OF THE PARAMETER UPDATE EQUATION

In this section, we describe the update equations of each parameter derived from the M-step of the EM algorithm. By differentiating the cost function about each parameter, the update equations were obtained. Let $X^{(H)}_{klmn}(c,t,f)$ and $X^{(I)}_{kl}(c,t,f)$ be the decomposed power:

$$X^{(H)}_{klmn}(c,t,f) = \Delta^{(H)}(m,n;k,l,t,f)X^{(J)}_{kl}(c,t,f)$$
$$\text{and} \quad X^{(I)}_{kl}(c,t,f) = \Delta^{(I)}(k,l,t,f)X^{(J)}_{kl}(c,t,f).$$

A.1 $w^{(J)}_{kl}$: overall amplitude

$$w^{(J)}_{kl} = \sum_c \iint \left(\sum_{m,n} X^{(H)}_{klmn}(c,t,f) + X^{(I)}_{kl}(c,t,f) \right) dt\, df.$$

A.2 $r_{kl}(c)$: relative amplitude of each channel

$$r_{kl}(c) = \frac{C \iint \left(\sum_{m,n} X^{(H)}_{klmn}(c,t,f) + X^{(I)}_{kl}(c,t,f) \right) dt\, df}{\sum_c \iint \left(\sum_{m,n} X^{(H)}_{klmn}(c,t,f) + X^{(I)}_{kl}(c,t,f) \right) dt\, df}.$$

A.3 $w^{(H)}_{kl}$, $w^{(I)}_{kl}$: amplitude of harmonic and inharmonic tone models

$$w^{(H)}_{kl} = \frac{\sum_{c,m,n} \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df}{\sum_c \iint \left(\sum_{m,n} X^{(H)}_{klmn}(c,t,f) + X^{(I)}_{kl}(c,t,f) \right) dt\, df}$$

and

$$w^{(I)}_{kl} = \frac{\sum_c \iint X^{(I)}_{kl}(c,t,f)\, dt\, df}{\sum_c \iint \left(\sum_{m,n} X^{(H)}_{klmn}(c,t,f) + X^{(I)}_{kl}(c,t,f) \right) dt\, df}.$$

A.4 $u_{kl}(m)$: coefficient of the temporal power envelope

$$u_{kl}(m) = \frac{\sum_{c,n} \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df + \mu^{(u)}_k(m)}{\sum_{c,m,n} \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df + 1}.$$

A.5 $v_{kl}(n)$: relative amplitude of n-th harmonic component

$$v_{kl}(n) = \frac{\sum_c \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df + \mu^{(v)}_k(m)}{\sum_{c,n} \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df + 1}.$$

A.6 τ_{kl}: onset time

$$\tau_{kl} = \frac{\sum_{c,m,n} \iint (t - m\phi_{kl}) X^{(H)}_{klmn}(c,t,f)\, dt\, df}{\sum_{c,m,n} \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df}.$$

A.7 $\omega_{kl}(t)$: F0 trajectory

$$\omega_{kl}(t) = \frac{\sum_{c,m,n} \iint nf X^{(H)}_{klmn}(c,t,f)\, df}{\sum_{c,m,n} \iint n^2 X^{(H)}_{klmn}(c,t,f)\, df}.$$

A.8 ϕ_{kl}: diffusion of a Gaussian of power envelope

$$\phi_{kl} = \frac{-A^{(\phi)}_1 + \sqrt{A^{(\phi)2}_1 + 4A^{(\phi)}_2 A^{(\phi)}_0}}{2A^{(\phi)}_1}, \quad \text{where}$$

$$A^{(\phi)}_2 = \sum_{c,m,n} \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df,$$

$$A^{(\phi)}_1 = \sum_{c,m,n} \iint m(t - \tau_{kl}) X^{(H)}_{klmn}(c,t,f)\, dt\, df, \text{ and}$$

$$A^{(\phi)}_0 = \sum_{c,m,n} \iint (t - \tau_{kl})^2 X^{(H)}_{klmn}(c,t,f)\, dt\, df.$$

A.9 σ_{kl}: diffusion of harmonic component along the frequency axis

$$\sigma_{kl} = \sqrt{\frac{\sum_{c,m,n} \iint (f - n\omega_{kl}(t))^2 X^{(H)}_{klmn}(c,t,f)\, dt\, df}{\sum_{c,m,n} \iint X^{(H)}_{klmn}(c,t,f)\, dt\, df}}.$$

A.10 $E^{(I)}_{kl}(t), F^{(I)}_{kl}(t,f)$: inharmonic tone model

$$E^{(I)}_{kl}(t) = \frac{\sum_c \int X^{(I)}_{kl}(c,t,f)\, df}{\sum_c \iint X^{(I)}_{kl}(c,t,f)\, dt\, df} \quad \text{and}$$

$$F^{(I)}_{kl}(t,f) = \frac{\sum_c X^{(I)}_{kl}(c,t,f) + \mu^{(F)}_k(f)}{\sum_c \int X^{(I)}_{kl}(c,t,f)\, df + 1}.$$

AUDIO COVER SONG IDENTIFICATION: MIREX 2006-2007 RESULTS AND ANALYSES

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ABSTRACT

This paper presents analyses of the 2006 and 2007 results of the Music Information Retrieval Evaluation eXchange (MIREX) Audio Cover Song Identification (ACS) tasks. The Music Information Retrieval Evaluation eXchange (MIREX) is a community-based endeavor to scientifically evaluate music information retrieval (MIR) algorithms and techniques. The ACS task was created to motivate MIR researchers to expand their notions of similarity beyond acoustic similarity to include the important idea that musical works retain their identity notwithstanding variations in style, genre, orchestration, rhythm or melodic ornamentation, etc. A series of statistical analyses were performed that indicate significant improvements in this domain have been made over the course of 2006-2007. Post-hoc analyses reveal distinct differences between individual systems and the effects of certain classes of queries on performance. This paper discusses some of the techniques that show promise in this research domain

1. INTRODUCTION

Founded in 2005, the annual Music Information Retrieval Evaluation eXchange (MIREX) is a community-based endeavor to scientifically evaluate music information retrieval (MIR) algorithms and techniques. Since its inception, over 300 music information retrieval (MIR) algorithms have been evaluated across 19 distinct tasks. These tasks were defined by community input and range from such low-level tasks such as Audio Onset Detection to higher-level tasks as Audio Music Similarity and Retrieval. More information about MIREX can be found at the MIREX wiki [8] where task descriptions and results are archived. This paper focuses on one specific task, namely Audio Cover Song Identification (ACS), which was first run in 2006 and repeated in 2007.

This paper is organized as follows: In Section 1.1, we discuss the motivation for conducting an ACS task. In Section 2, we introduce the task design and its evaluation dataset and the evaluation metrics used. In Section 3 we compare the results of the ACS 2006 and 2007 tasks. In Section 4, we focus on the ACS 2007 results and perform a set of statistical significance tests to investigate differences in system performance and the effects of the data on these performances. Section 5 summarizes what has been learned from the examining the different approaches to cover song identification. Section 6 contains the conclusion and future work.

1.1. Motivation

Aucouturier and Pachet's [1] seminal 2004 study identified the limitations of using audio-based timbral features to perform music similarity tasks. They performed more than one hundred machine learning experiments using spectral features and could only improve the performance 15% over a baseline. They called this problem the "glass ceiling". Whitman, et al. [11] investigated the "album effect", where they saw that the performances of artist identification systems were inflated by machine learning algorithms picking up on similarities in the production qualities of albums. The album effect has also been investigated by Kim, et al. [6]. Pampalk, et al. [9] addressed similar effects, where they evaluated genre classification systems on artist-filtered datasets and noted a marked reduction in performance.

The glass-ceiling, album and artist-filtering effects can also be seen throughout the MIREX 2005-2007 results. For example, comparing the best results for the Audio Genre Classification task of MIREX 2005, 82.34% (Bergstra, Casagrande and Eck), with the MIREX 2007 results, 68.29% (IMIRSEL (SVM)) [8] we see an apparent reduction in system performance across the two years. Similarly, the top Audio Artist Classification results for 2005, 72.45% (Mandel and Ellis) and 2007 48.14% (IMIRSEL (SVM)) also exhibit a seemingly large decline in performance. These performance drops can be partially explained by the fact that in 2005 there was no artist or album filtering of the test and training sets used these evaluations. In the 2007 Audio Genre Classification task, the data was filtered such that no track from the same artist could simultaneously exist in both the test and train sets in any cross-validation fold. Also, in the 2007 Audio Artist Identification task, the test collections did not include any track from the same album in test and training sets of any cross-validation folds.

The issues of an apparent glass ceiling, in conjunction with the absence of artist and album filtering in early MIR system evaluations overstating performance effectiveness,

indicated a need for the development and evaluation of methods using higher-order music descriptors in MIR similarity tasks. The ACS task was created to motivate MIR researchers to expand their notions of similarity beyond acoustic similarity to include the important idea that musical works retain their identity notwithstanding variations in style, genre, orchestration, rhythm or melodic ornamentation, etc. Because cover songs are known to span a range of styles, genres, and instrumentations, yet are often, in some sense, undeniably "similar," the evaluation of cover song identification performance can address the distinction between timbral similarity and "musical similarity". While identifying cover songs represents only a narrow scope of possible applications in regard to the use of higher-level features in MIR systems, it is an effective starting point in evaluating the usefulness of currently proposed "high-level musical descriptors", like those being investigated in [5][7].

2. ACS TASK DESCRIPTION

2.1. Evaluation Dataset

The ACS task dataset consists of 1000 tracks. Thirty different "cover song" groups each having 11 different versions for a total of 330 tracks are embedded within the ACS database. The original works in the cover song collection come from a variety of genres such as pop, rock, classical, baroque, folk, jazz, etc. Furthermore, within each group, the versions of each work similarly span a wide range of genres with different styles and orchestrations. The total length of the 330 cover songs is 21.2 hours with an average track length of 232 seconds (σ = 77 sec.). The remaining 670 tracks in the database are "noise". The noise tracks were chosen to be unrelated to any of the cover songs and their performing artists. The noise set also reflects a broad variety of genres and styles. The total length of the noise set is 45.8 hours with an average track length of 241 seconds (σ = 82 sec.). Thus the total length of the ACS dataset is 67 hours with an average track length of 242 seconds (σ = 72 sec.). Unlike many other MIREX tasks where 30 second clips were commonly used, the ACS task employed whole tracks to allow participants the opportunity to exploit the potentially important musical structure of the pieces.

All tracks in the dataset were encoded as 128 kbps MP3s and then decoded back to 22.05 kHz 16-bit WAV files using the LAME codec. Since the cover songs came from variety of sources with different encoding parameters, the MP3 encoding/decoding step was necessary to normalize the dataset to minimize the influence of coding effects on system performance.

2.2. Evaluation Methods and Metrics

The goal of the ACS task is to use each cover song track as a "seed/query" for identifying the 10 other versions of that piece in the dataset. All tracks in each cover song group are used as queries for a total of 330 queries. Since the underlying work of each individual cover song in a cover song group is known, the ground-truth for the ACS task is unambiguous and non-subjective. This distinguishes the ACS task from such other music similarity tasks as Audio Music Similarity, Audio Genre Classification, Audio Mood Classification, etc., which require the application of potentially subjective human judgments. The same dataset was used for both ACS 2006 and 2007. The identities of the pieces in the dataset have never been released to preclude the possibility of the *a priori* tuning of the submitted systems.

In ACS 2006, the overall evaluations were based on average performance and mean reciprocal rank (MRR). Average performance was defined as the mean number of covers identified within the top 10 returned items by the system. Rescaling the average performance score to the range of [0, 1] yields the precision at 10 (P@10) value for that system. Reciprocal rank was calculated as 1 over the rank of the first correctly identified cover song. In 2006 the systems only returned their top 10 candidates.

In ACS 2007, the participants introduced a new evaluation metric: mean average precision (MAP). For each query, average precision is calculated from the full returned list (i.e., 999 returned songs) as the average of precisions when the ranked list is cut off at each true item:

$$Ave.P = \frac{1}{10}\left(\sum_{r=1}^{999} p(r) \cdot I(r)\right) \quad (1)$$

where $p(r)$ is precision at rank r

$$p(r) = \sum_{j=1}^{r} \frac{I(j)}{r} \quad (2)$$

and $I(j)$ is a binary indicator function which is 1 if the j^{th} returned item in the list is a cover song, and 0 otherwise. The MAP is calculated as the mean of average precisions across all 330 queries. MAP is a commonly used metric in the text information retrieval domain [3]. Using MAP has the advantage of taking into account the whole returned list where correct items ranked closer to rank 1 receive the largest weights.

3. COMPARISON OF 2006-2007 RESULTS

Eight systems participated in ACS 2006 resulting in a task-wide P@10 of 0.08 (σ = 0.067), and a task-wide MRR of 0.19 (σ = 0.13). Table 1 (see Appendix A for legend) shows the P@10 values for each system. It is quite important to note that the systems labeled with '*' were not specifically designed for the ACS task. These systems, which were originally designed to participate in the Audio Music Similarity task, were graciously

volunteered to help the ACS organizers to determine whether standard music similarity algorithms would be adequate for the ACS task. Similarly, the task organizers included the IM system in the 2007 evaluations. The average P@10 of the top 4 (task-specific) systems in ACS 2006 was 0.13 ($\sigma = 0.073$). The average MRR of these 4 system was 0.28 ($\sigma = 0.14$).

Eight systems participated in ACS 2007 resulting in a task-wide MAP of 0.2062 ($\sigma = 0.1674$). The task-wide P@10 for ACS 2007 was 0.2057 ($\sigma = 0.1675$). The task-wide MRR was 0.38 ($\sigma = 0.27$). The average P@10 scores for the top 4 systems in 2007 were 0.34 with a standard deviation of 0.12.

Table 1 shows the P@10 values of the systems for both years. We can see a substantial improvement in ACS 2007 over 2006. The top 4 systems scores in 2007 are the same or better than the best score of 2006. After confirming that the top 4 mean P@10 values for both 2006 and 2007 are normally distributed using the Jarque-Bera (J-B) test [2] ($p < 0.05$), we ran a one-way ANOVA on the top 4 systems from each year to see if there is a statistically significant difference in performance between the years. The ANOVA indicated a significant difference between the P@10 means with $F(1,6) = 9.18$, $p = 0.023$. This result highlights that there has been a ~270% improvement of the top performing ACS systems in one year.

2006		2007	
DE	0.23	**SG**	0.50
KL1	0.11	**EC**	0.37
KL2	0.10	**JB**	0.26
CS	0.06	**JEC**	0.23
LR*	0.05	**KL1**	0.13
KWL*	0.04	**KL2**	0.09
TP*	0.04	**KP**	0.06
KWT*	0.03	**IM****	0.01

Table 1. Precision at 10 for ACS 2006 – 2007 results

It is also important to look at the non-task-specific systems. For example, the IM system which had the lowest score of all systems over both years was based on the same naïvely constructed spectral feature set (i.e., MFCC's, zero-crossing rates, spectral flux, spectral centroid, etc.) as the IM-SVM system that ranked amongst the top systems for the Audio Artist Identification, Audio Genre Classification and Audio Classical Composer Identification tasks in MIREX 2007. The weak performance of the non-task-specific systems strongly suggests that to capture the identity aspects of "music similarity," one should go beyond simple spectral features. Top-performing systems in ACS 2007 used higher-level features (e.g., rhythm, tonality, tonal sequences, etc.) so as to capture the musically important structures.

4. ANALYSIS OF 2007 RESULTS

In this section we will focus exclusively on the ACS 2007 results since the overall performance of the 2007 systems is significantly better than the 2006 group. Furthermore, in 2007 all but one system (i.e., IM) were specifically designed for the ACS task.

4.1. Descriptive Analyses of ACS 2007 Results

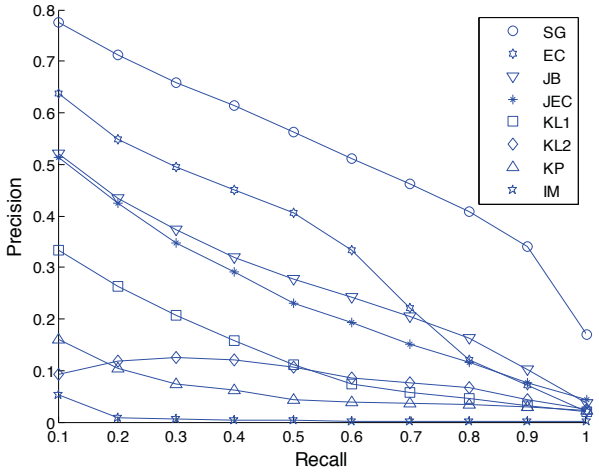

Figure 1. Precision-Recall curves for ACS 2007

The precision-recall graph in Figure 1 was generated by calculating the precision values at each recall level from 0.1 to 1 and averaging across all 330 queries for each system. Looking at the graph, we can see that SG and EC retrieved substantial numbers of relevant cover songs in early portions of their results list. In fact SG had 26 queries with perfect precision at recall equal to 1. SG had a further 10 queries where the entire relevant set was returned within the first 11 items. EC had 12 queries where 9 out of the 10 relevant cover songs were returned in the first 10 items. These results are quite remarkable because, given our dataset size (1000) and the number of relevant items per query (10), the probability of randomly returning the entire relevant set within the top 10 list once in 330 queries is only 1.26×10^{-21}! Also noteworthy, is the extraordinarily flat performance curve of the IM system.

Figure 2 shows the box-whisker plot of the distributions of MAP values for each system across the 30 cover song groups. The bottom, middle and top of each box represent the lower quartile, median and upper quartile values, respectively. The '+'s are the outliers for each distribution. The data were amalgamated with respect to their cover song groups because ACS task is primarily interested in system performance with regard to the identification of the 30 underlying works rather than the 330 individual pieces. Thus, a MAP score was calculated

for each cover song group as the mean of the average precision values for each group's 11 members.

In Figure 2, we see that there is a fair amount of variance across query groups with respect to system performance. This is especially noticeable in the top 4 performing systems (i.e., JB, JEC, EC, SG). This suggests that some query groups might have significant effects (positive or negative) on system performance.

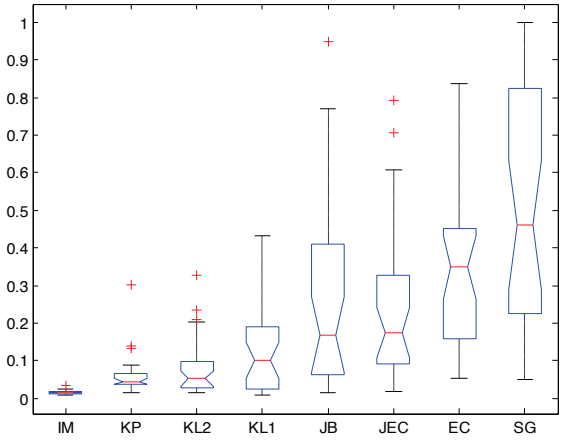

Figure 2. Box whisker plot of MAP distributions for each system across cover groups (query groups).

4.2. Inferential Significance Testing

In this section, we analyze the results in several different perspectives. We are interested in determining the answers to the following questions:

1. a) Is there any significant difference in performance means among the systems? and, b) If such differences exist, between which systems do they occur?
2. a) Is there any significance difference among performance means in query groups? and, b) If such differences exist, between which query groups do they occur?

Following the procedures outlined in [3], we determined whether the by-query-group MAP data discussed above were normally distributed across query groups using the J-B test. Most of those data did not conform to the normal distribution. However, after applying the arcsine square-root transformation:

$$y = arcsin(\sqrt{x}) \quad (3)$$

as recommended by [10], 7 out of 8 systems across query groups, and 28 out of 30 query groups across systems passed the J-B test ($p < 0.05$). Since the dataset is approximately normally distributed using arcsine square-root transformation, we selected a parametric test to investigate whether there are any significance differences among systems or query groups. Parametric tests are preferred, where appropriate, because they are more powerful than their non-parametric counterparts: they better detect differences that might be overlooked (i.e., they have lower Type II error rates).

A two-way ANOVA was chosen because it can provide answers to both of our system (Q.1) and our query group (Q.2) questions simultaneously. Table 2 shows the results of the two-way ANOVA on the transformed data. As one can see, there are significant differences among both systems and query groups.

Source	Sum Sq.	D.F.	Mean Sq.	F-stat	P
Systems	10.51	7	1.50	54.64	0.00
Query Groups	6.53	29	0.23	8.20	0.00
Error	5.58	203	0.027		
Total	22.62	239			

Table 2. Two-way ANOVA table

To further investigate the differences between individual system performance means (Q.1b), we performed the Tukey-Kramer Honestly Significantly Different (TK-HSD) analysis on the system data. TK-HSD was used because it can properly control the experiment-wise Type-I error rate unlike the commonly misused multiple *t*-tests [10]. Figure 3 shows the results of the TK-HSD on the transformed by-query-group MAP data for each system. The circled items refer to individual groupings based on the absence of significant differences within the grouping. The answer to Q.1b can be seen in Figure 3. It is evident that the SG system is significantly better than the other systems in this task. Also EC, JB and JEC have formed their own grouping. It is important to note that these results differ from those presented ACS 2007 results wiki [8] where the non-parametric Friedman's test was performed on the non-transformed data. The lower power of the Friedman's test appears to have missed the significantly better performance of the SG system.

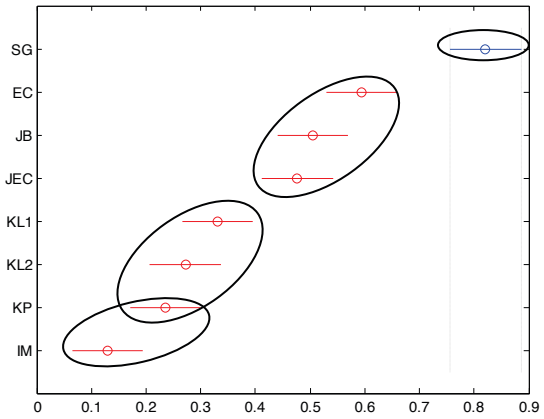

Figure 3. TK-HSD analysis on system effects based on the transformed by-query-group MAP data.

To answer Q.2b, We ran the TK-HSD test to determine where differences in performance occur with regard to the query groups. Table 3 displays the results. The first column represents the anonymized IDs of the query groups which are rank ordered with respect to the by-query-group MAP across the systems. The second column presents the by-query-group MAP. The shaded regions indicate sets of query groups which are not significantly different in how the algorithms performed. For example, the first column indicates that query groups 1 through 10 are not significantly different from one another.

Group no.	Avg. MAP
1	0.52
2	0.42
3	0.41
4	0.35
5	0.35
6	0.34
7	0.33
8	0.30
9	0.29
10	0.28
11	0.27
12	0.24
13	0.24
14	0.20
15	0.18
16	0.16
17	0.16
18	0.16
19	0.13
20	0.12
21	0.10
22	0.10
23	0.08
24	0.08
25	0.08
26	0.08
27	0.06
28	0.05
29	0.05
30	0.03

Table 3. Significance sets of cover song groups.

Table 3 clearly illustrates the wide range from 0.03 to 0.52 ($\sigma = 0.24$) of by-query-group MAP performances.

Since there is such a large discrepancy between the best query group and the worst, we investigated the two extremes to explain attempt to explain how systems behave in response to different query groups.

The best performing group (Group 1 in Table 3) is a late 17th-century canon with a regular structure and harmonic progression. All versions of Group 1 share the same rhythmic structure as they should, since canons are composed of replication of the rhythms and intervals of the same main theme. We surmise this makes it easier for algorithms to accurately compare the rhythmic and tonal structures of the songs. There is not much variance in orchestration or tempo in Group 1. This was one of the query groups, in which SG achieved its near-perfect MAP scores (0.998). The SG method uses sequences of tonal descriptors where songs are matched using dynamic programming for local alignment. The EC system also performed very well (MAP of 0.838) for this group. It uses correlation of beat-synchronous chroma features.

The worst performing group (Group 30 in Table 3) is an 18th century hymn set to its now-traditional 19th century melody. All the song versions in this cover group vary greatly in their harmonies, chord progressions, rhythms, tempi and dynamic ranges. The performances encompass many different styles such as country, blues, jazz, hip-hop, rock, etc. Group 30 songs exhibit a great deal of widely varying melodic ornamentation and several different languages. Virtually all task-specific systems use tempo and key-independent matching of the underlying tonal or harmonic structure of the pieces. Because the variations of the Group 30 songs contain a wide range of embellishments and changes to the harmonic structure, we believe the systems are sensitive to the varying nature of this group. SG scored a MAP of 0.052. EC scored the highest MAP of 0.06 for Group 30.

5. DISCUSSION

In the design of audio cover song algorithms, the top performing algorithms share a variety of attributes. A majority of the four top performing algorithms use chromagrams or pitch class profiles as the predominant feature representation, with methods for addressing possible changes in key and tempo in matching songs. Chromagrams represent the distribution of spectral energy quantized to the chromatic scale.

The EC algorithm addresses variations in tempo by performing a tempo induction stage and producing a beat-synchronous chromagram that contains a single chroma vector per beat, and uses cross correlation of the chromagrams for song matching. The SG system uses dynamic programming to align and match harmonic pitch class profiles. JEC also uses an underlying chromagram representation, but filters the logarithm of each of the 12 chroma channels into 25 logarithmically spaced bands. This 12×25 feature matrix captures the variation of each of the 12 chroma channel on scales of 1.5 to 60 seconds. Because a logarithm is used prior to filtering, large changes in tempo become apparent as simple shifts along the filter channel axis. Song matches are performed by calculating the Frobenius distance between feature matrices. JB also uses chromagrams, but these are used for performing HMM-based chord identification, with string alignment techniques being used to perform song matches on the chord transcription.

To address changes in key, the top performing algorithms perform circular shifts of their underlying representations to address possible transpositions. Therefore to calculate a possible song match, similarity scores are calculated multiple times for each transposition.

In contrast to the top performing algorithms, the worst performing algorithms across the two years are based predominantly on timbre features, which are highly effective for audio music similarity, genre identification, etc. However, for cover song identification, it is clear that

analysis of musical structure, and dealing with musical alterations to aspects such as key and tempo are necessary.

6. CONCLUSIONS AND FUTURE WORK

This paper presented an analysis of the evaluation of audio cover song identification systems. While the aim of identifying variations of musical pieces in audio collections is narrow in scope with regard to the overarching goals of MIR, it represents a necessary departure from a large portion of the MIR research done to date. In particular, we assert that cover song identification necessarily must explore "musical similarity" along structural dimensions, as opposed to those characterized merely by timbre. This is demonstrated by the poor performance of timbre-based audio similarity algorithms in identifying cover songs. However, we do not wish to imply that cover song identification is in some way superior to, or meant to serve as a replacement for related similarity and classification tasks (e.g. audio artist, audio genre, etc). Instead, it represents an interesting new direction of research because of its apparent need for analyzing underlying musical structure. The significant performance gains in a single year and the impressive performances of the top algorithms in 2007 suggest that some of the musical descriptors used by these algorithms are seemingly quite powerful. As discussed often in terms of the "glass ceiling" problem, it is our hope that such descriptors, in conjunction with all of the other research that has been carried out to date, can push the state of MIR research forward, and also allow musical similarity searches to be tailored along structural dimensions (e.g. "find me a song with a similar chord progression").

In the broader context of music similarity, an interesting future direction would be to test the ability of "cover song" systems to retrieve "relevant" songs (not necessarily cover versions) from an audio collection given a query. While we have seen algorithms intended to retrieve similar songs in the MIREX audio music similarity task performed poorly in cover song identification, it would be interesting to see if the reverse is true. That is, it could be beneficial to note whether these cover song systems, which rely more on matching tonal or chord sequences, would produce results that a human judge would deem "similar." This very topic is addressed by Ellis, Cotton and Mandel [4]. Although they found that traditional MFCC based approaches are superior to using only their beat synchronous chromagrams, the features used in cover song identification did perform significantly better than a random baseline.

7. ACKNOWLEDGEMENTS

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9. APPENDIX

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TP	Tim Pohle
EC	Daniel P. W. Ellis, Courtenay V. Cotton
IM	IMIRSEL M2K
JB	Juan Bello
JEC	Jesper Højvang Jensen, Daniel P. W. Ellis, Mads G. Christensen, Søren Holdt
KP	Youngmoo E. Kim, Daniel Perelstein
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HUBS AND HOMOGENEITY: IMPROVING CONTENT-BASED MUSIC MODELING

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ABSTRACT

We explore the origins of hubs in timbre-based song modeling in the context of content-based music recommendation and propose several remedies. Specifically, we find that a process of model homogenization, in which certain components of a mixture model are systematically removed, improves performance as measured against several ground-truth similarity metrics. Extending the work of Aucouturier, we introduce several new methods of homogenization. On a subset of the `uspop` data set, model homogenization improves artist R-precision by a maximum of 3.5% and agreement to user collection co-occurrence data by 7.4%. We also explore differences in the effectiveness of the various homogenization methods for hub reduction. Further, we extend the modeling of frame-based MFCC features by using a kernel density estimation approach to non-parametric modeling. We find that such an approach significantly reduces the number of hubs (by 2.6% of the dataset) while improving agreement to ground-truth by 5% and slightly improving artist R-precision as compared with the standard parametric model.

1 INTRODUCTION

Content-based music similarity is a promising but underdeveloped approach to automatic music recommendation. To date, most work in this area has been focused on calculating similarity through comparison of song-level statistical models. However, such systems have thus far yielded only limited results [2], regardless of modeling method. It is thought that this may be connected to the existence of "hubs", songs that are found to be inaccurately similar to a large number of songs in a database. The origin of these hubs has been conjectured, yet no clear strategy for combating them has been established.

We conjecture that some songs are modeled particularly poorly, in effect leaving them far from other songs in the database and thus unlikely to be recommended. These songs, which we call "anti-hubs", are shown to be identifiable from certain properties of their models. In this paper, we propose a method to systematically reduce the incidence of anti-hubs. Another modeling approach suggested by the goal of hub reduction was also explored.

2 METHODOLOGY

2.1 Prior Work

Gaussian mixture models (GMMs) of short-time MFCC frames have been explored extensively and are considered the state-of-the-art approach to content-based song modeling [10, 7, 4, 8, 1]. Typically, the symmetric Kullback-Leibler (KL) divergence is found between these models and is used as a similarity measure. Since there is no closed-form solution for mixture models, this distance must be approximated, usually by a Monte Carlo method [1] or the Earth Mover's distance (EMD) [11].

2.2 Modeling

Following this work, we also use the MFCC-GMM approach for song modeling. While Aucouturier [2] showed 50 components to be optimal, for speed we chose to use 32 components, and empirically deemed these to have sufficient modeling power.

For experiments using non-parametric modeling, we employed kernel density estimation (KDE) [9, 6] as implemented by MATLAB's `ksdensity` routine. Our models therefore consist of a sampled density function for each MFCC dimension, considering each to be independent. We empirically determined a resolution of 1,000 points per density function was sufficient to represent the distributions. The kernel bandwidth, which depends on the number of frames and their median absolute deviation, is scaled to control the smoothness of the density estimate, and this scaling can be varied to explore its effect on model performance.

2.3 Distance

Initial experiments showed the Monte Carlo-based distance to be prohibitively slow for comparing GMMs, and EMD was used instead. For KDE models, we adopted the Bhattacharyya distance [5], a common measure of similarity be-

tween two discrete probability distributions. Note that because each density is sampled over different ranges, we linearly interpolate over the maximum range of the two given models so that each density is defined and compared for common x values.

2.4 Data

These experiments used a subset of the `uspop` collection consisting of 617 songs from 40 artists. This set was intended to match the relative size of Berenzweig's subset [3], while not hand-picking tracks based on intra-class timbral homogeneity and inter-class heterogeneity as with Aucouturier's set [1].

2.5 Hubness

In measuring a kernel's hubness, we adopted the N-occurrences measure used by Aucouturier [1] and Berenzweig [3], choosing N to be 100. This measure is a count of the number of times a song appears in the top-N list of other tracks, in that a large value indicated a hub. Like Aucouturier, we considered a track a hub if its 100-occurrences are greater than 200 (2 times the mean) and an anti-hub if its 100-occurences is less than 20.

2.6 Ground-truth Agreement

In measuring agreement to ground-truth, we first measured each kernel's artist R-precision. This is the percentage of retrieved the R nearest neighbors with the same artist as the seed, where R is the number of the artist's songs in the data set. This corresponds to the common k-NN classifier with leave-one-out cross-validation, except that k is dependent on the size of each seed's class.

As another measure of ground-truth agreement, we used the OpenNap user collection co-occurrence data accompanying the `uspop` collection [4]. Using the top-N rank agreement score, we found how well our computed kernels' neighbor rankings matched the kernel computed from the OpenNap data.

3 HUBS OR ANTI-HUBS?

Berenzweig discovered that models with very few near neighbors, which we now refer to as anti-hubs (and classified by Pampalk as "always dissimilar" [8]), had certain characteristic properties [3]. It was hypothesized that perhaps the focus in the quest for understanding hubs was on the wrong side of the hub distribution: "... hubs may actually be the only songs behaving nicely, while non-hubs [are] pathologically far away from everything else." Because we base our recommendations and, in result, notions of hubness, on nearest neighbors in kernel space, anti-hubs could actually be considered as problematic as hubs. In other words, anti-hubs are absent from their rightful timbral neighborhoods, leaving their would-be neighbors near other songs that are perhaps not perceptually suitable.

	Correlation
Trace of single Gauss. covar.	-0.2432
Max. intra-comp. dist.	-0.3272
Max. comp. dist from centroid	-0.3156

Table 1. Pearson correlation coefficients between 100-occurences count and measures of model spread

We speculate that these anti-hubs originate not from what would be considered perceptually anomalous timbres, but from a relative small number of frames representing transient timbral sections. Because the algorithms used to train song models are musically agnostic (i.e. silence is as musically valid as a chorus), we have found several components of mixture models are spent modeling these unrepresentative timbral sections.

This section demonstrates that models of anti-hubs tend to contain outlier mixture components that can prove detrimental to their parent models' discriminative power. We also propose that anti-hubs are at least easier to identify through measuring attributes of these components and therefore more easily treatable.

3.1 Model Variance

By measuring the overall "variance" of his GMMs, Aucouturier found no correlation with hubness and this measure of model "spread" [1], disproving his hypothesis that hubs are well-connected to other models simply due to a relatively large distribution of frames. However, using three other measures of model spread, we found a negative correlation between model size and hubness, as seen in Table 1. This suggests hubs actually have small spreads compared to anti-hubs, likely indicating that anti-hubs have largely multi-modal distributions.

3.2 Outlier Components

But, as Berenzweig observed [3], the large spread of anti-hubs can be attributed to relatively few mixture components. Berenzweig observed anti-hubs contain components with very small variance, leading to models that are overly specific to a certain region in feature-space and thus making them less likely to match other models. He also found these components tend to have other common attributes: relatively significant prior probabilities so they cannot be ignored as "mathematical nuisances", large distance from the mixture's centroid meaning they are most likely to blame for anti-hubs' overall "wide diameter" models, and close proximity to the origin, suggesting these components are primarily

	Correlation
Min. log-det. of covar.	0.2247
Max. dist. from centroid	−0.3113
Min. dist. from origin	0.3253
Min. prior probability	0.0908

Table 2. Pearson correlation coefficients between 100-occurences count and measures of component attributes

modeling low-energy frames. We found that components of anti-hubs in general can be characterized with the same attributes. To verify, we calculated the Pearson correlation between each model's 100-occurrences count and measurements of the most extreme component according to these attributes. Table 2 shows these correlations, which were all found to be statistically significant.

4 HOMOGENIZATION

Aucouturier concurred that a significant amount of modeling power was being occupied by certain outlier frames, as seen through his experiments with "homogenizing" models [1]. His experiments were based on the idea that components with high prior probabilities model statistically important frames, so that we can, in effect, associate these component weights with component "importance". He then removed components whose prior probabilities fell below a given threshold, producing a "homogenized" version of the original model. Through this experiment, he claimed that most of the variance of a GMM is accounted for by the least 5-10% of the statistically weighted components. Also, he argued that the hubness of a song is based primarily on the least statistically "important" components, as the hubness of his collection increased by nearly a factor of 3 when the models were homogenized to just 90%.

Mixture models, however, typically contain components that are highly overlapped. In this way, the prior probability, or "weight", of a particular component may be low, but together with its neighbor components, could comprise a large mode in the overall density function. Therefore, the prior probabilities alone cannot be assumed to correlate with a component's "importance".

Therefore, we claim that component prior probabilities are not a reliable feature to effectively homogenize mixture models. We instead make use of the correlates to hubness highlighted in the previous section. In particular, we propose to base homogenization around procedures aimed at removing the components characterized by the above features. In each case, practically the same algorithm described by Aucouturier is used: components not meeting a certain defined threshold requirement are discarded and the component weights (prior probabilities) are re-normalized.

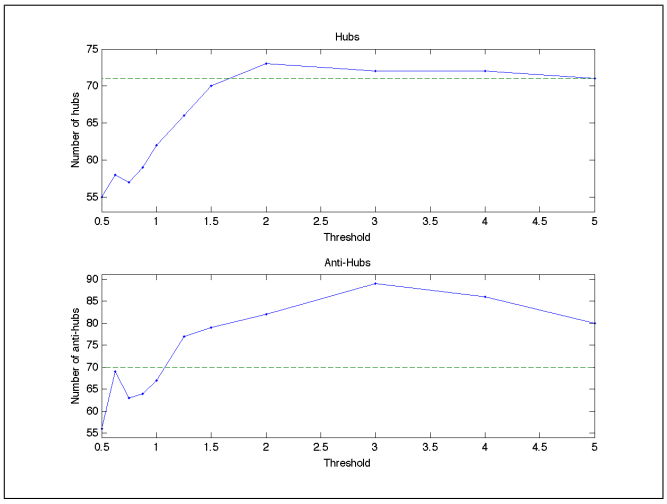

Figure 1. Influence of homogenization by distance from mixture centroid on number of hubs (top) and anti-hubs (bottom) for different thresholds. The un-homogenized amounts are plotted as horizontal lines for reference.

4.1 Homogenization by Distance from Centroid

The first method of homogenization explored was based on the observation that anti-hubs tend to have components that are distant from a main mode of the frame distribution. We therefore discarded components whose Euclidean distance from the component center to the mixture's centroid was greater than a given threshold. The threshold values were determined empirically by observing many activation sequences (showing the likelihood that each frame occurred from each GMM component) of models found from all sections of the hub distribution, as inspired by Berenzweig [3].

4.1.1 Effects on hubness

Figure 1 shows the effects of homogenization on the occurrence of hubs and anti-hubs. Note the symmetry of the un-homogenized distributions: there are approximately the same number of hubs and anti-hubs (71 and 70, respectively). It is clear that the number of hubs and anti-hubs decreased only for severe homogenization levels.

Interestingly, the number of anti-hubs greatly increased after mild homogenization. If anti-hubs become more centralized in distribution space after homogenization as intended, they should attain more neighbors. But would these neighbors be new or are anti-hubs simply getting closer to their previous nearest neighbors? To answer this, we observed where in the hub distribution each song's nearest neighbors existed. It was clear that anti-hubs' only near neighbors tended to be other anti-hubs. Because of this, if we treat some anti-hubs with homogenization, their former neighbors, who were generally unaffected by this procedure, become more severe anti-hubs. Therefore, while several

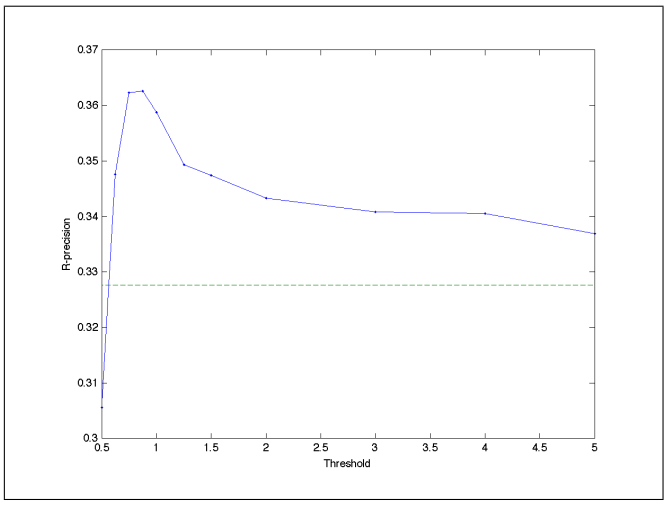

Figure 2. Influence of homogenization by distance from mixture centroid on artist R-precision. The R-precision level before homogenization is plotted as a horizontal line for reference.

anti-hubs are clearly being introduced into the song pool, increasing not only their similarity accuracy but also that of their new neighbors, we see many borderline anti-hubs dropping into the anti-hub region after homogenization.

4.1.2 Effects on agreement to ground-truth

Figure 2 shows the artist R-precision computed for each homogenization by distance from centroid threshold. R-precision increased monotonically until a threshold of 0.875 with a maximum increase of 3.50% (absolute) above the un-homogenized baseline (R-precision with a random kernel was found to be 0.03). This was likely due to anti-hubs being brought nearer to their appropriate perceptual neighbors, as songs by the same artist are generally similar timbrally. After this threshold, the R-precision drops dramatically. The decrease with severe homogenization was no doubt due to song models disintegrating into generic distributions with little discriminating information.

We then computed the top-39 (our dataset contains 40 artists) rank agreement scores against the OpenNap user co-occurrence kernel for each level of homogenization. Each score was averaged over 1,000 runs to smooth out inconsistencies resulting from ties. A Wilcoxon signed-rank test showed that agreement scores for homogenization distance thresholds from 3.0 to 0.875 were significantly higher than the un-homogenized score, where a maximum increase of 5.95% (absolute) over the GMM-EMD kernel was found.

4.2 Homogenization by Variance

We next examined homogenization by variance. With this method, we removed components whose log-determinant did not meet a minimum threshold. The determinant of a covariance matrix can be thought of as a measure of a component's volume, so we were in effect removing components that cover a small region of the timbral space. Again, the existence of such components was shown to be negatively correlated with a model's hubness, so we again expected homogenization to primarily affect models on the lower end of the hub distribution.

Note we were unable to affect all models with this approach without removing all of certain models' components. Therefore, our most severe homogenization level with this method affected only 70% of the models. This highlights an advantage in the use of relative component features (e.g. distance from centroid) in defining the homogenizing function, as opposed to absolute cut-offs (e.g. log-determinant).

4.2.1 Effects on hubness

Unlike the previously discussed homogenization method, this method did not show improvement in hubness at any level. The number of hubs increased by 2 for weak homogenization and remained unchanged for more severe thresholds. The number of anti-hubs, in fact, increased by 16 (2.6% of the dataset) for most homogenization levels. This was assumed to be a result of the aforementioned abandonment of borderline anti-hubs.

4.2.2 Effects on agreement to ground-truth

Despite its apparent detriment to hubness, this homogenization improved agreement to ground-truth. All levels except for the most severe were found to significantly increase over the un-homogenized level, with a maximum increase of 1.26% at the -110 threshold. Note this is only about half of the improvement seen with homogenization by distance from mixture centroid.

Significant improvement was also seen in agreement to the OpenNap user co-occurrence data, fairly consistently across homogenization levels. The maximum increase of 6.18% (absolute) was seen at a log-determinant threshold of -105.

4.3 Homogenization by Distance from Origin

The last homogenization method explored was based on the observation that anti-hubs tend to have components near the origin. These are likely modeling frames with low energy (e.g. silence) and can reasonably be considered not perceptually relevant in relation to the song's timbre. As before, several thresholds were found empirically, and components less than this distance away from the origin were discarded from the model. Like with homogenization by variance, we were only able to treat at most 60% without fulling collapsing certain models whose components were all fairly near the origin.

4.3.1 Effects on hubness

Hubness was not improved for any homogenization level with this method. In fact, like with homogenization by variance, we saw a slight increase in hubs and a considerable increase in anti-hubs, by more than 15 (2.4% of the dataset) for each threshold.

4.3.2 Effects on agreement to ground-truth

We saw that artist R-precision also improved with this homogenization, increasing monotonically with distance from origin threshold. The maximum increase of 3.09% (absolute) was found when discarding components less than 5.5 units from the origin (in MFCC space). All changes were found to be significant under the Wilcoxon signed-rank test.

Agreement with the OpenNap data significantly increased as well with this type of homogenization, increasing monotonically and reaching a maximum of 7.38% (absolute) above the un-homogenized baseline at a threshold of 5.5.

5 NON-PARAMETRIC MODELING

We discussed in Section 3 that using algorithms such as Expectation-Maximization to train parametric mixture models such as GMMs can result in mixture components that are devoted to modeling timbral frames that are not related to perceptually salient sections. This tends to result in models with poor representative power that in turn leads to inaccurately low similarity scores with other models. However, instead of iteratively training a parametric model to fit a given distribution, non-parametric approaches can be used to explicitly model these complex distributions of MFCC frames. In particular, using kernel density estimation (KDE), we are given some control over the effect spurious frames have on a model by increasing the kernel bandwidth. Wider bandwidths yield smoother density functions, effectively reducing the multi-modal behavior shown to be consistent with songs containing outlier frames (i.e. anti-hubs).

Aucouturier compares a type of non-parametric modeling to other modeling algorithms in his thesis [1]. Using three methods (independent histograms and two vector quantization methods), he shows that each performs much worse than the GMM approach, in both R-precision and hubness. Interestingly, our approach here is similar to his independent histograms modeling (which scored 24% lower in R-precision than GMM), in that we treat each MFCC dimension independently, but since we use estimated density functions, we use the Bhattacharyya distance or the Monte Carlo approximated divergence to compare these models instead of Euclidean distance.

We verified that our KDE models were consistent with the GMM models by computing the top-N rank agreement between kernels. We chose N to be 616 and slowly decaying rank weights to allow for a large set of neighbors to impact the scores. The agreement scores are shown in Table 3 and show KDE kernels from both distance metrics agree well with the GMM kernel.

	Agreement
Bhattacharyya	0.8940
Monte Carlo	0.8866
Random	0.3218

Table 3. Top-N rank agreement scores for KDE kernels and the standard GMM kernel using different distance metrics and unity bandwidth scaling

5.1 Hubness

Looking at hubness, however, here was a large discrepancy between the GMM kernel and the KDE-BD kernels. A large decrease was found in both the number of hubs and anti-hubs, as high as 16 and 29 respectively or 23% and 41% of the un-homogenized GMM levels. It seems there is no strong relationship between hubness and the smoothness of the density function. After examining this in more detail, it was shown that the anti-hub region is unaffected by smoothing of the density functions. This goes against our earlier hypothesis that anti-hubs are severely multi-modal, which first led to our experiments with homogenization. We speculated that the smoother the density functions (i.e. the more homogenized the underlying distribution), the more hub-like the model would become. We did see the amount of hubs decrease 9.8% (61 to 55) with increased smoothing, suggesting, if anything, we were decreasing hubs. This could be a result of more models from the middle of the hub distribution moving nearer hubs, thus splitting the former hubs' neighbors amongst the new hubs. In this way, a song simply occupying a centralized region in space (or as Aucouturier calls a "center of mass") does not make it a hub; the song must be relatively alone in this region.

It was also shown that there is a strong correlation (0.788) between the hubness values of GMM and KDE models. In other words, songs that appear as hubs in the GMM kernel are likely to appear as hubs in the KDE kernel. This is contrary to Aucouturier's experiment [1] where he finds a much weaker correlation between hubs appearing from GMMs and his non-parametric histogram models.

5.2 Agreement to ground-truth

No significant difference was found for artist R-precision scores on the KDE kernels as compared to the GMM kernels. However, KDE modeling improved agreement to the OpenNap data, where we saw an improvement of about 5% (absolute) for all bandwidth sizes. Similar results were seen

with the Monte Carlo distance, with a maximum increase over the GMM-EMD baseline of 6% (absolute).

5.3 Computation Time

Aside from apparently better discriminative power, KDE-BD models also showed advantages in necessary computation time. The total computation time to train the KDE models on all 617 songs of the `uspop` subset was found to decrease exponentially with kernel bandwidth. For very small bandwidths, we saw modeling time increase by over a factor of 15 over GMMs, with no apparent detriment to modeling power.

As far as distance computation time, the Bhattacharyya distance (with linear interpolation and 2,000-point density functions) took on average 83 ms. per pair of KDE models, compared to 30 ms. for finding the distance between two GMMs via the Earth Mover's distance [1]. This means computing a KDE-BD kernel took about 2.7 times longer (262 minutes) than the GMM-EMD kernel (95 minutes) on our 617-song `uspop` subset. Speed of the BD computation could of course be improved by employing lower order interpolation and more sparsely sampling the density functions.

The Monte Carlo distance, on the other hand, took significantly longer, averaging 586 ms. per pair (generating 2,000 samples per model), meaning the entire kernel took 31 hours to compute, which is entirely unacceptable in a real-world scenario. Granted, measures could be taken to increase its efficiency, but since the results on the above performance tasks were comparable to the BD kernel, no reason is seen to further use the computationally expensive Monte Carlo-based distance.

6 CONCLUSION

Homogenization of GMMs was shown to improve the hubness of several models, particularly anti-hubs. While the overall amounts of hubs and anti-hubs generally increased after this procedure, this was assumed to be a result of the abandonment of anti-hub neighbors who were themselves untreatable by the given homogenization method. Each method showed significant improvements, however, in agreement to ground-truth data, as shown in Table 4. This is encouraging, since the improved representative power of the models affected by homogenization seems to outweigh the expected loss in models left untreated.

Non-parametric modeling by kernel density estimation proved to offer not only significant reduction in hubness but considerable improvement in computation time and ground-truth agreement.

[1] Computed on a MacPro with 2 2.66 GHz Dual-Core Intel Xeon processors and 2 GB of RAM

Homogenization Method	Artist R-precision	OpenNap Agreement
Dist. from centroid	3.50%	5.95%
Variance	1.26%	6.15%
Dist. from origin	3.10%	7.40%

Table 4. Maximum percent improvement (absolute) of agreement to ground-truth data for different homogenization methods.

Overall, the work presented here suggests approaches to solutions to fundamental problems found in content-based music modeling.

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EXTENDING CONTENT-BASED RECOMMENDATION: THE CASE OF INDIAN CLASSICAL MUSIC

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ABSTRACT

We describe a series of experiments that attempt to create a content-based similarity model suitable for making recommendations about North Indian classical music (NICM). We introduce a dataset (nicm2008) consisting of 897 tracks of NICM along with substantial ground-truth annotations, including artist, predominant instrument, tonic pitch, *raag*, and parent scale (*thaat*). Using a timbre-based similarity model derived from short-time MFCCs we find that artist R-precision is 32.69% and that the predominant instrument is correctly classified 90.30% of the time. Consistent with previous work, we find that certain tracks ("hubs") appear falsely similar to many other tracks. We find that this problem can be attenuated by model homogenization. We also introduce the use of pitch-class distribution (PCD) features to measure melodic similarity. Its effectiveness is evaluated by *raag* R-precision (16.97%), *thaat* classification accuracy (75.83%), and comparison to reference similarity metrics. We propose that a hybrid timbral-melodic similarity model may be effective for Indian classical music recommendation. Further, this work suggests that "hubs" are a general features of such similarity modeling that may be partially alleviated by model homogenization

1 INTRODUCTION AND MOTIVATION

North Indian classical music (NICM) has for the past forty years become an increasingly international phenomenon. A great number of people have had some exposure but otherwise know little about the tradition. This presents an opportunity for a music discovery system that can suggest music based either on known artists or on simple descriptive terms. However, metadata for Indian classical music is often missing or inaccurate, and user-tagging of Indian classical music tracks is uncommon. These problems suggest the use of a content-based recommender. In addition to the goal of exploring models that can be used for content-based recommendation, we hope to explore whether issues observed with the standard timbre-based content-based recommendation (CBR) models, such as the prevalence of many false hits due to a few tracks ("hubs"), are artifacts of the data or appear more generally when novel music is considered. To date, published work on CBR [1, 2, 12] has focused on a few datasets that consist solely of a small slice of Western popular music. Finally we hope to show how properties of the musical genre can be exploited to improve CBR. Because NICM is significantly less polyphonic than most Western music, it relatively easy for us to extract melodic information which can be used in a similarity model.

2 BACKGROUND

NICM is one of the oldest continuous musical traditions in the world and it is an active contemporary performance practice. Since the 1960's, due to the emigration of Indians and the popularity of artists such as Ravi Shankar and Zakir Hussain, it has become widely known to international audiences. The repertoire of Indian classical music is extensive, consisting of dozens of styles and hundreds of significant performers. The most prevalent instruments, such as *sitar*, *sarod* and *tabla*, are timbrally quite different from popular Western instruments. NICM is an oral tradition and recordings therefore represent the primary materials.

The performance of Indian classical music typically involves a soloist, either a vocalist or instrumentalist, accompanied by a *tanpura* (drone) throughout and a *tabla* (percussion) in rhythmic sections. Most presentations begin with an extended ametric melodic improvisation (*alap*) and build in intensity throughout the performance. After this section, several compositions are usually presented with *tabla* accompaniment. Here too, the majority of the music is improvised. All NICM is based on *raag*, a melodic concept that defines the melodic parameters of a given piece. There are hundreds of *raags*, of which approximately 150 are widely performed. *Raags* are typically defined by a scale and a set of interrelated phrases. Although highly structured, they allow the performer tremendous scope for improvisation and elaboration. It is this development that forms the core of most NICM performances.

Another important, though quite distinct, performance tradition is solo *tabla*, in which the *tabla* player becomes the soloist. Here timbral and rhythmic patterns form the core material, and the melodic accompaniment is used primarily as a time-keeper.

3 DATABASE

The database was assembled from the author's personal music collection. Recordings encompass both commercial and non-commercial sources. Many of the non-commercial recordings are live concerts that are distributed informally amongst NICM listeners. A substantial number of the most historically important recordings are of this type. The recordings span a range from the early 20^{th} century to present with the vast majority of recordings being from the second half of the century. Low fidelity is common due to the quality of the initial recording or the number of intermediate analog copies. Common degradations include hiss and crackle, overloading, missing high and/or low frequency content, artifacts from excessive noise reduction processing, and wow due to tape speed fluctuation. The balance of the accompanying drone and *tabla* varies widely, in some cases barely audible, in other cases overwhelming.

The database consist of 897 tracks. Only the first five minutes of each track was used, leading to a total size of approximately seventy hours. This was done to reduce computation time since many of the tracks were over thirty minutes long. A total of 141 artists are contained in the database as well as 14 different instruments. The instruments include *sitar*, *sarod*, *tabla*, *shenai*, *flute*, *violin*, and *pakhawaj*, with the first three being the most common. There are 171 different *raags*. The distribution of tracks amongst the *raags* was uneven and 71 *raags* are represented by only one recording in the database.

All the features used in this study, including MFCCs and pitch-tracks, along with ground-truth annotation will be made available at `paragchordia.com/data/nicm08/`.

3.1 Annotation

For each track the main artist, instrument and *raag* was annotated. In duet tracks, of which there were only fourteen, both instruments were noted. A substantial difficulty in analyzing *raag* recordings is that there is no standard reference scale, and each performer is free to choose any pitch for the tonic. Thus, if frequency values are to be later interpreted as scale degrees, the tonic must be known for each recording. For each track an expert listener tuned an oscillator while listening to the performance. The tracks were divided amongst two experts, and tracks that were challenging or ambiguous were reviewed by both. This annotation was not done for solo *tabla* tracks, as they were excluded from the melodic similarity experiments.

4 METHOD

We describe the feature extraction and statistical modeling used to develop the timbral and melodic similarity models.

4.1 Timbre Modeling

Timbre modeling was done using MFCCs calculated on 20ms windows overlapped by 50%. Twenty coefficients were used excluding the 0^{th} coefficient. As mentioned earlier, features were only calculated on the first five minutes of each song.

A model was built for each track by assuming that each MFCC feature vector was a sample from an underlying distribution for the current track. Because a given track is likely to evolve over time, statistical models that are flexible enough to represent multiple clusters in the feature space are typically used. Following earlier approaches [1], a Gaussian mixture model (GMM) was trained for each track based on the frame-based MFCCs. The GMM was trained by initializing the means using the k-means clustering algorithm and then running the EM algorithm [10]. A total of thirty-two components were used for each GMM, each with a diagonal covariance matrix. The model parameters, namely the means and covariance matrices of the Gaussian components, become our model of each track.

The similarity of two tracks was judged by comparing the distributions that had been learned for each track [3, 12]. Although there are many intuitive ways to measure the distance of points in a feature space, it is less obvious how to compare distributions. Several methods have been proposed such as Kullback-Liebler (KL) divergence and Earth Movers Distance (EMD) [13]. KL divergence measures the relative entropy of two distributions: that is, the reduction in uncertainty for one distribution if the other is known. EMD has been widely applied to the comparison of GMMs. The algorithm considers the minimum cost to "move" the probability mass of the first distribution so it resembles the second. In one dimension it is easy to visualize: each GMM is a set of hills and the hills are moved and probability mass shifted from one to another until they are matched. Another approach that is perhaps the most natural is to compute the likelihood that the features vectors of one track are generated by the other tracks distribution. This last method, while intuitive, is rarely used because of the computational cost. Regardless of the method employed, the distance metric allows us to calculate a scalar value representing the measure of similarity between tracks. If we have n tracks then n^2 distances must be calculated, or $(n)(n+1)/2$ if the distance measure is symmetric, meaning that our computation time will increase as a square of the number of tracks we wish to analyze. The arrangement of all such distance pairs forms a similarity matrix. For any given seed song in the database the similarity matrix can then be used to retrieve the k nearest neighbors, which can then be used as candidates for recommendation.

4.2 Melody Modeling

NICM, as noted in the introduction, is based on *raag*. In addition to specifying melodic constraints, *raags* are tradition-

ally thought to elicit certain emotions in listeners. Chordia [7] empirically demonstrated that certain *raags* consistently elicit certain emotions, such as joy or sadness, even for listeners with little or no prior exposure to NICM. Thus *raag* identification is an important descriptor of both melodic and emotional content of a track. Chordia [6] demonstrated that pitch-class distributions (PCDs) could be used to recognize *raags* using a variety of classification algorithms. Further, PCDs might reveal connections or perceptual similarities between particular recordings beyond those suggested by *raag* name alone.

These insights are used in the current system to build a melodic similarity model. First, each piece was pitch tracked using the YIN [9] algorithm. Each pitch estimate was then converted to a scale degree using the manually annotated tonic. Given the tonic, the locations of the scale degrees were computed using the ratios that define the chromatic notes of a just intoned scale. The pitch estimate at each frame was compared to the ideal scale values in the log domain and assigned to the nearest scale degree. The octave information was then discarded, giving a sequence of pitch-classes. A histogram was then computed yielding a pitch-class distribution for the track. Because of the consistent presence of the drone, the tonic value usually overwhelms all other scale degrees without providing any useful discriminative power and was therefore discarded. Thus each track in the database was characterized by one eleven dimensional feature vector.

In addition to a pitch estimate, the YIN algorithm returns a pitch aperiodicity measure which was used to weight the pitch estimates. In one case, which we call linear pitch salience weighting, pitch aperiodicity was converted to a pitch salience measure as $1 - pitchAperiodicty$. In a second case, called ratio weighting, the pitch salience was defined as $1/pitchAperiodicity$. Previous work [8] showed improved performance on the *raag* classification task after weighting, particularly for ratio weighting. In practice, such weighting tends to eliminate or de-emphasize regions of the track where the soloist is silent and the pitch track is therefore noisy and uninformative.

A melodic similarity matrix was constructed by evaluating the distance between pairs of PCDs for each of the conditions. Two distance metrics were used, Pearson correlation and Euclidean distance, with each of these yielding a distinct similarity matrix. The similarity models were then used to retrieve the k nearest neighbors for each track.

5 EVALUATION

In the end, we are interested in how well such a similarity engine might perform for such tasks as making purchase recommendations, or suggesting songs in playlist. As such, the final evaluation requires building an application and judging its success in terms of user engagement or purchases.

model type	Accuracy Rates		
	k = 1	k = 5	k = 10
GMM	81.27%	71.28%	65.64%
GMM - homogenized	90.30%	80.57%	73.36%
non-parametric	87.96%	75.96%	67.97%

Table 1. Instrument classification accuracy using three different modeling techniques with a k-NN classifier. GMMs without homogenization using Earth Mover's Distance, homogenized with log determinant threshold of -150, and non-parametric modeling are shown.

However, before building such a system we would like to have some sense of whether our similarity judgments are appropriate. Various proxy tasks are used that hopefully correlate with the ultimate utility of the system. For example, Berenzweig [2] and Aucouturier [1] have used artist R-precision and classification tasks to judge the quality of the recommendations based on their timbre models.

Although it is true that an artist's sound may change from recording to recording, it is nevertheless likely that many artists will be timbrally consistent and thus distinguishable from each other . Therefore if our timbre model tends to return hits of the same artist we would tend to think it is doing better than if it does not. In addition to tasks based on artist name, we also evaluated R-precision and classification accuracy for the predominant instrument. Similarly, for melodic evaluation, we considered *raag* R-precision, *thaat* classification, and correlation with a ground-truth matrix expressing known relationships between *raags*.

5.1 Timbral Evaluation

Following the standard definition, we define R-precision to be the number of relevant hits for a given seed divided by the total possible relevant hits in the database. For example, in the instrument task, we consider the relevant hits to be the number of the R nearest neighbors that have the same instrument label as the seed, where R represents the total number of tracks with the same instrument as the seed. For the artist task, average R-precision was 26.96% using a GMM model and EMD compared with 2.08% for neighbors chosen randomly. A related task is classification, in which the k nearest neighbors are used to classify the seed track. Table 1 gives the classification performance for recognition of the predominant instrument. Nearest neighbor ($k = 1$) classification performs best with an accuracy of 81.27% for fourteen instrument targets. In the rare case (14 tracks) where there are two main instruments, we consider classification successful if either of the main instruments is matched.

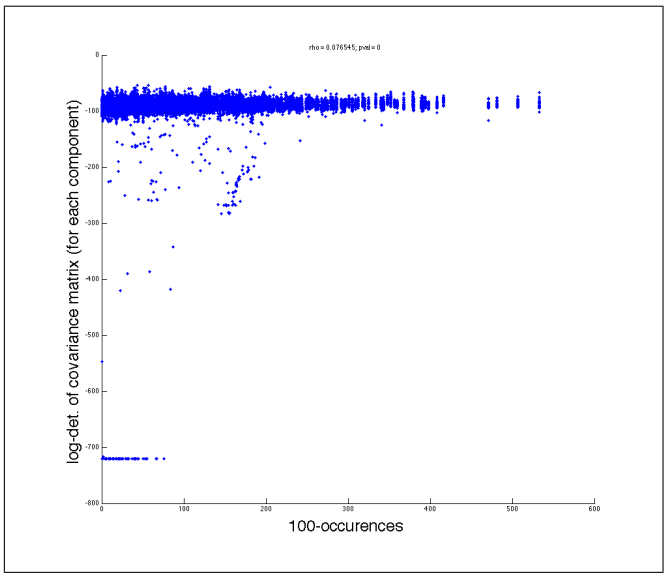

Figure 1. Log of the determinant for the components of GMMs. It can be seen that certain tracks have outlier components where the log-determinant is very small.

5.2 Hubness

It has been previously observed that a substantial problem with the standard timbre modeling strategies for CBR is the existence of hubs, certain tracks which inappropriately appear as hits for many different seeds [1]. More precisely, we define a hub to be a track that is a top one hundred hit for at least 200 of the seeds in the database (twice the average). In our dataset a total of 105 tracks, or 11.71% of the database, fit this definition. Recent work by Godfrey [11] has shown that these hubs may arise because certain tracks, termed "anti-hubs", are effectively taken out of the pool of possible neighbors due to poor modeling by the GMM. 121 tracks in our dataset (13.49%) were anti-hubs, which we defined as those that match less than twenty seeds. Figure 1 shows the log-determinant of the covariance matrix for each GMM component of each track. The determinant of the covariance matrix can be thought of as a measure of the volume of the Gaussian data cloud. We see that certain components have nearly zero volume. By looking at an activation matrix, which indicates when a GMM component is active in a track, Godfrey found that such components are often active only for very short segments of the track and are otherwise unused.

A possible solution to this problem is the removal of degenerate components of the GMM. This homogenized model may then have a more evenly distributed hub histogram. To see the effect of this, we reduced each GMM by removing components where the log of the determinant was less than a specified threshold. This was done for three levels as shown in Table 2. The similarity matrix was then recomputed using

Threshold	R-Precision
-300	0.3040
-200	0.3215
-150	0.3269

Table 2. Artist R-Precision results obtained using homogenization

these new models. We find that artist R-precision increases by 5.73 percentage points to 32.69% for the maximum level of homogenization, and instrument classification accuracy increases by 9.03 percentage points to 90.30%. This is consistent with results on the uspop2002 dataset, and suggests that the problems of hubs is general and may be ameliorated by model homogenization [11].

An alternative approach to the hub problem is to use alternative modeling strategies. One such approach is to model feature vectors non-parametrically rather than using a GMM. In kernel density estimation, the points in the feature space representing observations (e.g. MFCC feature vectors) are essentially convolved with a window such as a Gaussian, and summed to yield the density estimate [10]. The estimated densities are then compared using a metric such as the Bhattacharya distance [5], defined as

$$B(p,q) = -log \sum_{x \in X} \sqrt{p(x)q(x)}. \quad (1)$$

Using such an approach, we found that instrument R-precision increased by 3.14 percentage points to 30.10% and instrument classification accuracy improved 6.69 percentage points to 87.96% compared with the non-homogenized GMM model. Again, this is consistent with experiments performed on the uspop2002 dataset [11], suggesting that this modeling approach may be broadly applicable. A further advantage of this non-parametric approach is that computation time is greatly reduced, primarily because the iterative EM algorithm step can be skipped. The bandwidth of the kernel was varied but was not found to have a significant effect over a range of reasonable values.

5.3 Melody Evaluation

Raag, as noted above, is the most essential melodic description of NICM. Our first evaluation task was therefore *raag* R-precision. Accuracy was 16.97% compared to a random baseline of 1.18%. We excluded *tabla* and *pakhawaj* solos as well as a few semi-classical tracks that did not have a clearly defined *raag*. The sparseness of the data, with respect to certain *raags*, led us to additionally classify each track according to parent scale, or *thaat*. The system of abstract parent scale-types, developed by Bhatkande in the early 20^{th} century [4], consists of a mapping of the wide variety of scale-types used in *raags* to a small set of seven

note scales that were considered to represent basic melodic spaces. This step reduced the number of melodic categories to ten. Classification accuracy for *thaats* was 75.83% using the nearest neighbor. Surprisingly, neither of the pitch-salience weighted vectors performed better than the unweighted PCDs.

Although R-precision and classification tasks are informative, we would like to be able to compare the similarity matrix to some ideal ground-truth similarity matrix. Although no such data exist based on perceptual experiments, reasonable references may nevertheless be built.

A ground-truth similarity matrix was built by converting each *raag* to a binary PCD vector that indicated whether each of the twelve scale degrees was used in that *raag*. Distances between *raags* were then computed using the Hamming distance, which counts the number of mismatches between the vectors. It should be noted that a match is found when both *raags* use a certain scale degree and also when both omit a certain scale degree. We also experimented with the inner product, which counts the number of scale degrees that are shared between the *raags*. However, we found that the Hamming distance matched intuitive notions of distance better since the absence of scale degree in a *raag* is as important as its presence.

One approach to comparing similarity matrices is to generate lists of hits for each seed using both the empirical similarity matrix and the ground-truth similarity matrix. The average distance of the hits from the seed would be computed for all seeds in the empirical similarity matrix, for example based on the Euclidean distance between each retrieved track and the seed. Likewise the distances of the neighbors fetched according to the ground-truth matrix could be calculated. The averages over each model might then be compared. The problem with such an approach is that the distances cannot be directly be compared, since they use two different distance metrics. For example, in our case the ground-truth distances are based on Hamming distances between binary PCDs while the empirical distances are based on correlation between the continuous-valued PCDs. One approach that has been proposed to solve this problem is to use the rank order rather than the distance. Taking inspiration from measures used in text retrieval, Berenzweig et al. [3] define what they call the Top-N Ranking agreement:

$$s_i = \sum_{r=1}^{N} (\alpha_r)^r (\alpha_c)^{k^r}, \qquad (2)$$

where k_r is the ranking according to the empirical similarity matrix of the r^{th}-ranked hit using the ground-truth matrix. The score is computed for each seed and averaged to give the overall agreement. The α_c and α_r parameters determine the sensitivity of the score to ordering in each of the lists generated by the two metrics.

In order to compare the distances directly we used Hamming distance on the empirical results. This was done by converting each track into a binary PCD vector based on its *raag* label so that the Hamming distance could be applied. This made it easy to compare empirical performance to an upper bound, whereas the Top-N Ranking agreement can be difficult to interpret.

similarity matrix	Average Distances		
	k = 1	k = 5	k = 10
ground-truth	0.0331	0.1444	0.2770
empirical	1.831	2.3669	2.6235
random	4.682	4.668	4.6442

Table 3. Average distances from seeds calculated with binary PCDs for ground-truth, empirical, and random similarity matrices.

Results are substantially better than random. The rank list score was .1085 compared with .0087 for the random, with $N = 10$, $\alpha_c = .5^{1/3}$, and $\alpha_r = .5^{2/3}$. Using our method for direct distance comparison, the average distance for $k = 5$ was .1444 for the ground truth matrix and 4.668 for the random case. The value for the empirical similarity matrix fell in the middle of this range (2.367). These results are summarized in table 3.

We also found that hubness occurred with PCD vectors, although to a lesser extent than with timbral modeling. Hub and anti-hub percentages using the unweighted PCD feature and Hamming distance were 8.9% and 4.7% respectively, using the same definition as above. Not surprisingly this was less than for the timbre models, most likely due to the lower dimensionality of the feature space (eleven vs. twenty). Berenzweig demonstrated that hubness increases with the dimensionality of the feature space [2].

6 DISCUSSION

Although the artist R-precision is relatively low (32.69%), this is expected because in NICM instrumentation is quite similar for many artists. Without further annotation indicating higher-level artist clusters, we would expect relatively low precision values. That instrument classification accuracy was over 90% suggests that the timbre model is capturing essential information. The instrument classification result is particularly encouraging since it is likely that connecting tracks with the same main instrument would be important for a recommendation system. Similarly, although *raag* R-precision was low, *thaat* identification was correct in more than three out of four cases. Comparisons to random and best-case scenarios suggest that despite the difficulties of pitch tracking real recordings, PCDs are sufficiently robust to provide useful melodic information.

The results presented suggest that MFCC based timbre modeling is effective for NICM and generalizes beyond Western popular music. Further we find support for the idea that hubs may be a general problem when models are constructed using GMMs on frame-based MFCC features. This work also supports the observation that model homogenization may lead to improved retrieval precision and classification accuracy.

7 APPLICATION

We anticipate that the similarity modeling presented here could be used as the basis for a music recommendation system based on both timbral and melodic characteristics. One way of combining them would be to create a global distance metric. In the simplest case, one could weight timbre and melody equally and simply sum the distances in the respective similarity matrices. More likely we might imagine giving users control over the extent to which recommendations were controlled by one parameter or the other. This could be done explicitly, for example through a slider interface, or implicitly in a live application by tracking the perceived quality of the recommendations, for example by allowing users to rate the suggested tracks. We conjecture that this combined model approach may also allow adaptation to different patterns of user preferences; timbre might dominate the quality judgments of some users, while others might be more responsive to melodic content.

8 FUTURE WORK

Although encouraging, the current models are clearly quite simplistic and the results suggest there is ample room for improvement. On the timbral side, although including more features may improve performance incrementally, as is often noted, significant improvement will require a more perceptually grounded model. For melodic modeling we intend to generalize PCDs to include sequential structure though n-gram modeling. Earlier work has shown that this markedly improves *raag* classification performance [6]. Pitch tracking could also be improved by suppression of accompaniment.

We hope to use the techniques discussed here to create a CBR system for Indian classical music in which listeners will be able to generate playlists based either on artists or tracks, or alternatively based on simple emotional descriptors. This would allow us to replace evaluation based on measures such as R-precision and artificial ground-truth matrices with more objective measures of the success of these models in generating music streams.

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ANALYZING AFRO-CUBAN RHYTHM USING ROTATION-AWARE CLAVE TEMPLATE MATCHING WITH DYNAMIC PROGRAMMING

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ABSTRACT

The majority of existing research in Music Information Retrieval (MIR) has focused on either popular or classical music and frequently makes assumptions that do not generalize to other music cultures. We use the term Computational Ethnomusicology (CE) to describe the use of computer tools to assist the analysis and understanding of musics from around the world. Although existing MIR techniques can serve as a good starting point for CE, the design of effective tools can benefit from incorporating domain-specific knowledge about the musical style and culture of interest. In this paper we describe our realization of this approach in the context of studying Afro-Cuban rhythm. More specifically we show how computer analysis can help us characterize and appreciate the complexities of tracking tempo and analyzing micro-timing in these particular music styles. A novel template-based method for tempo tracking in rhythmically complex Afro-Cuban music is proposed. Although our approach is domain-specific, we believe that the concepts and ideas used could also be used for studying other music cultures after some adaptation.

1 INTRODUCTION

We present a set of techniques and tools designed for studying rhythm and timing in recordings of Afro-Cuban music with particular emphasis on "clave," a rhythmic pattern used for temporal organization. In order to visualize timing information we propose a novel graphical representation that can be generated by computer from signal analysis of audio recordings and from listeners' annotations collected in real time. The proposed visualization is based on the idea of Bar Wrapping, which is the breaking and stacking of a linear time axis at a fixed metric location.

The techniques proposed in this paper have their origins in Music Information Retrieval (MIR) but have been adapted and extended in order to analyze the particular music culture studied. Unlike much of existing work in MIR in which the target user is an "average" music listener, the focus of this work is people who are "experts" in a particular music culture. Examples of the type of questions they would like to explore include: how do expert players differ from each other, and also from competent musicians who are not familiar with the particular style; are there consistent timing deviations for notes at different metric positions; how does tempo change over the course of a recording etc. Such questions have been frequently out of reach because it is tedious or impossible to explore without computer assistance.

Creating automatic tools for analyzing micro-timing and tempo variations for Afro-Cuban music has been challenging. Existing beat-tracking tools either don't provide the required functionality (for example only perform tempo tracking but don't provide beat locations) or are simply not able to handle the rhythmic complexity of Afro-Cuban music because they make assumptions that are not always applicable, such as expecting more and louder notes on metrically "strong" beats. Finally the required precision for temporal analysis is much higher than typical MIR applications. These considerations have motivated the design of a beat tracker that utilizes domain-specific knowledge about Cuban rhythms.

The proposed techniques fall under the general rubric of what has been termed *Computational Ethnomusicology* (CE), which refers to the design and usage of computer tools that can assist ethnomusicological research [14] . Futrelle and Downie argued for MIR research to expand to other domains beyond Western pop and classical music [9]. Retrieval based on rhythmic information has been explored in the context of Greek and African traditional music [1].

Our focus here is the analysis of music in which percussion plays an important role, specifically, Afro-Cuban music. Schloss [13] and Bilmes [4] each studied timing nuances in Afro-Cuban music with computers. Beat tracking and tempo induction are active topics of research, although they have mostly focused on popular music styles [11]. Our work follows Collins' suggestion [5] to build beat trackers that embody knowledge of specific musical styles.

The *clave* is a small collection of rhythms embedded in virtually all Cuban music. Clave is a repeated syncopated rhythmic pattern that is often explicitly played, but often only implied; it is the essence of periodicity in Cuban music. An instrument also named "clave" (a pair of short sticks hit together) usually plays this repeating pattern. Clave is found mainly in two forms: *rumba clave* and *son clave*. (One way of notating clave is shown in Figure 1.)

Figure 1. Son (left) and rumba (right) clave

Our study of timing requires knowing the exact time of every note played by the clave. We can then decompose this data into an estimate of how tempo changes over time (what is called the *tempo curve*) and a measure of each individual note's deviation from the "ideal" time predicted by a metronomic rendition of the patterns shown in Figure 1.

Unfortunately, we do not know of any databases of Afro-Cuban music with an exact ground-truth time marked for every clave note or even for every downbeat. [1] Therefore we constructed a small four-song database [2] and gathered ground truth clave timing data by having an expert percussionist with Afro-Cuban experience tap along with the clave part. Custom sample-accurate tap detection/logging software automatically timestamps the taps.

Recordings of Afro-Cuban music challenge existing state-of-the-art beat-tracking algorithms because of the complex and dense rhythm and the lack of regular approximately isochronous pulses. Figure 2 shows how two recent state-of-the-art beat-tracking systems (BeatRoot [7] and a beat tracker using dynamic programming proposed by Ellis [8]) do not generate an accurate tempo curve for the recording *CB*. The plots in the figure are shown only in order to motivate the proposed approach. The comparison is not fair, as the other algorithms are more generally applicable and designed with different assumptions, but in any case it demonstrates the advantage of a domain-specific method to deal with these recordings: our method is specifically designed to take into account clave as the rhythmic backbone.

2 DATA PREPARATION

It is common for Afro-Cuban songs to begin with just the sound of the clave for one or two repetitions to establish the initial tempo. However as other instruments (both percussive and pitched) and voices enter the mix the sound of the clave tends to become masked. The first step of data preparation is to enhance the sound of the clave throughout the song using a matched filter approach. In addition onset detection is performed.

[1] Bilmes recorded about 23 minutes of Afro-Cuban percussion at MIT in 1992, and performed sophisticated analysis of the timing of the *guagua* and *conga* (but not clave) instruments [4]; unfortunately these analog recordings are not currently available to the research community.

[2] Here is the name, artist, and source recording for each song, along with the two-character ID used later in the paper: *LP*: *La Polemica*, Los Muñequitos de Matanzas, Rumba Caliente 88. *CB*: *Cantar Bueno*, Yoruba Andabo, El Callejon De Los Rumberos. *CH*: *Chacho*, Los Muñequitos de Matanzas, Cuba: I Am Time (Vol. 1). *PD*: *Popurrit de Sones Orientales*, Conjunto de Sones Orientales, Son de Cuba.

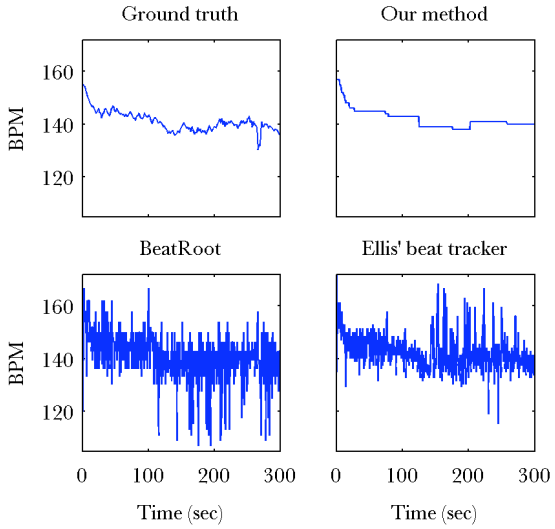

Figure 2. Four estimates of the tempo curve for our recording *CB*: Ground truth calculated from a human expert's tap times (upper left), curve from our method (top right), curve from BeatRoot (lower left), and curve from Ellis' dynamic programming approach (lower right).

2.1 Clave enhancement using Matched-Filtering

A matched filter detects or enhances the presence of an *a priori* known signal within an unknown signal. Its impulse response is a time-reversed copy of the known signal, which in our case is the beginning portion of one isolated clave note. The clave instrument affords little timbral variety and therefore every note of clave in a given recording sounds substantially like all the others, so a matched filter made from any single note (frequently easily obtained from the beginning of the song) will enhance the presence of the clave throughout the song and suppress the remaining signal. One free parameter is the filter order, i.e., the duration of the segment of the clave note; in each case we selected a "good" matched filter experimentally by listening to the output of different configurations. All the curves in Figure 2 and results in this paper have been calculated on audio signals output by matched filtering.

2.2 Onset detection

Onset detection aims at finding the starting time of musical events (e.g. notes, chords, drum events) in an audio signal; see [3],[6] for recent tutorials. We used *spectral flux* as the onset detection function, defined as:

$$SF(n) = \sum_{k=0}^{N/2} HWR(|X(n,k)| - |X(n-1,k)|) \quad (1)$$

where $HWR(x) = \frac{x+|x|}{2}$ is the half-wave rectifier function, $X(n,k)$ represents the k-th frequency bin of the n-th frame of the power magnitude (in dB) of the short time Fourier transform, and N is the corresponding Hamming window size. For the experiments performed in this work all data had a sampling rate $f_s = 44100$ Hz and we used a window size of 46 ms ($N = 2048$) and a hop size of about 11ms ($R = 512$). The onsets are subsequently detected from the spectral flux values by a causal peak-picking algorithm that finds local maxima as follows. A peak at time $t = \frac{nR}{f_s}$ (the time of the beginning of the nth frame) is selected as an onset if it fulfills the following conditions:

1. $SF(n) \geq SF(k) \ \ \forall k \ : n - w \leq k \leq n + w$
2. $SF(n) > \frac{\sum_{k=n-mw}^{n+w} SF(k)}{mw+w+1} \times thres + \delta$

where $w = 6$ is the size of the window used to find a local maximum, $m = 4$ is a multiplier so that the mean is calculated over a larger range before the peak, $thres = 2.0$ is a threshold relative to the local mean that a peak must reach in order to be sufficiently prominent to be selected as an onset, and $\delta = 10^{-20}$ is a residual value to avoid false detections on silent regions of the signal. All these parameter values were derived from preliminary experiments using a collection of music signals with varying onset characteristics.

In order to reduce the false detection rate, we smooth the detection function $SF(n)$ with a Butterworth filter to reduce the effect of spurious peaks:

$$H(z) = \frac{0.1173 + 0.2347z^{-1} + 0.1174z^{-2}}{1 - 0.8252z^{-1} + 0.2946z^{-2}} \qquad (2)$$

(These coefficients were found by experimentation based on the findings in [3],[6].) In order to avoid phase distortion (which would shift the detected onset time away from the $SF(n)$ peak) the signal is filtered in both the forward and reverse directions.

3 TEMPLATE-BASED TEMPO TRACKING

We propose a new method to deal with the challenges of beat tracking in Afro-Cuban music. The main idea is to use domain specific knowledge, in this case the clave pattern, directly to guide the tracking. The method consists of the following four basic steps: 1) Consider each detected onset time as a potential note of the clave pattern. 2) Exhaustively consider every possible tempo (and clave rotation) at each onset by cross-correlating each of a set of clave-pattern templates against an onset strength envelope signal beginning at each detected onset. 3) Interpret each cross-correlation result as a score for the corresponding tempo (and clave rotation) hypothesis. 4) Connect the local tempo and phase estimates to provide a smooth tempo curve and deal with errors in onset detection, using dynamic programming.

The idea of using dynamic programming for beat tracking was proposed by Laroche [10], where an onset function was compared to a predefined envelope spanning multiple beats that incorporated expectations concerning how a particular tempo is realized in terms of strong and weak beats; dynamic programming efficiently enforced continuity in both beat spacing and tempo. Peeters [12] developed this idea, again allowing for tempo variation and matching of envelope patterns against templates. An approach assuming constant tempo that allows a simpler formulation at the cost of more limited scope has been described by Ellis [8].

3.1 Clave pattern templates

At the core of our method is the idea of using entire rhythmic patterns (templates) for beat tracking rather than individual beats. First we construct a template for each possible tempo. We take the ideal note onset times in units of beats (e.g., for rumba clave, the list 0, 0.75, 1.75, 2.5, 3) and multiply them by the duration of a beat at each tempo, giving ideal note onset times in seconds. We center a Gaussian envelope on each ideal note onset time to form the template. The standard deviation (i.e., width) of these Gaussians is a free parameter of this method. Initial results with a constant width revealed a bias towards higher tempi, so widths are specified in units of beats, i.e., we scale the width linearly with tempo. Better results were obtained by making each template contain multiple repetitions of the clave, e.g., three complete patterns. Figure 3 shows a visual representation of the template rotations for all considered tempi.

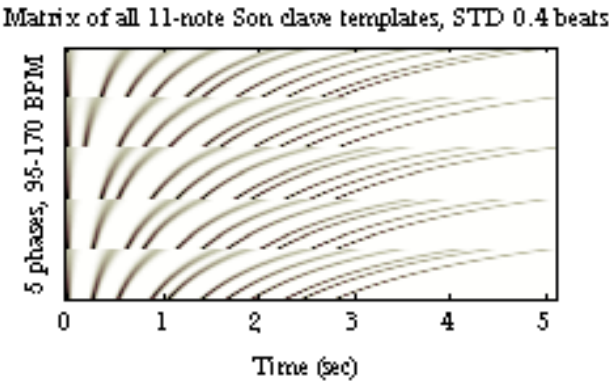

Figure 3. Clave Templates for all rotations and tempi

With a 5-note clave pattern, any given note played by the clave could be the 1st, 2nd, 3rd 4th or 5th note of the pattern. Therefore we make templates for all "rotations" of the clave, i.e., for the repeating pattern as started from any of the five notes. For example, rotation 0 of rumba clave is [0, 0.75, 1.75, 2.5, 3], and rotation 1 (starting from the second note) is [0.75 1.75 2.5 3 4] - 0.75 = [0 1 1.75 2.25 3.25]. Time 0 always refers to the onset time of the current note.

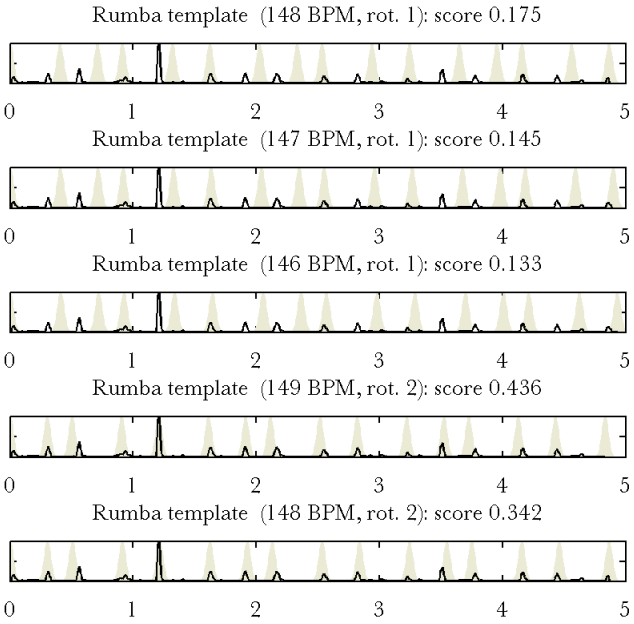

Figure 4. Matching different templates (filled grey) to the *CB* recording's energy envelope (black line). The X-axis is time (seconds), with zero the time of the onset under consideration. The score for each template match represents how well that template lines up with the energy envelope.

We cross-correlate (in other words, take the dot product of) these templates against segments of an onset strength envelope (in our case, simply the total energy in each 1024-sample window of the matched filter output) beginning at the time of each detected onset. We interpret the dot product between the onset strength signal $O(t)$ and a template $T_{j,k}(t)$ with tempo j and rotation k as the strength of the hypothesis that the given onset is the given note of clave at the given tempo. Figure 4 depicts this process for some tempi and rotations and the corresponding scores. We exhaustively compute these dot products for every candidate tempo j (e.g,. from 95 to 170 BPM in 1 BPM increments), for all five rotations of the clave pattern k , for every detected onset i at time t_i to produce a *score grid*:

$$score(i,j,k) = \sum_{t=0}^{LT_{j,k}-1} T_{j,k}(t)O(t_i+t) \qquad (3)$$

where $LT_{j,k}$ is the length of template $T_{j,k}$.

3.2 Rotation-blind dynamic programming

It is trivial to look at a given onset, pick the tempo and rotation with the highest score, and call that the short-term tempo estimate. However, due to the presence of noise, inevitable onset detection errors, and the matched filter's far-from-perfect powers of auditory source separation, simply connecting these short-term tempo estimates does not produce a usable estimate of the tempo curve. Better results can be achieved by explicitly discouraging large tempo changes. We use dynamic programming [2] as an efficient means to estimate the best tempo path (i.e., time-varying tempo). In the next section we will consider the rotations of the template; for now let the "rotation-blind" score be:

$$scoreRB(i,j) = max(score(i,j,k))\ k:1..5 \qquad (4)$$

We convert each score $scoreRB$ to a cost $C_{i,j}$ with a linear remapping so that the highest score maps to cost 0 and the lowest score maps to cost 1. We define a *path* P as a sequence of tempo estimates (one per onset), so that $P(i)$ is P's estimate of the tempo at time t_i. Our algorithm minimizes the *path cost* PC of the length n path P:

$$PC(P) = \sum_{i=0:n-1} C_{i,P(i)} + \sum_{i=0:n-2} F(P(i),P(i+1)) \quad (5)$$

where $F(tempo_1, tempo_2)$ is a "tempo discontinuity cost function" expressing the undesirability of sudden changes in tempo. F is simply a scalar times the absolute difference of the two tempi. Dynamic programming can efficiently find the lowest-cost path from the first onset to the last because the optimal path up to any tempo at time t_i depends only on the optimal paths up to time t_{i-1}. We record both the cost $PC(i,j)$ and the previous tempo $Previous(i,j)$ for the best path up to any given onset i and tempo j.

3.3 Rotation-aware dynamic programming

Now we will extend the above algorithm to consider rotation, i.e., our belief about which note of clave corresponds to each onset. Now our cost function $C_{i,j,k}$ is also a function of the rotation k. Our path tells us both the tempo $P_{tempo}(i)$ at time t_i and also the rotation $P_{rot}(i)$, so we must keep track of both previous $Previous_{tempo}(i,j)$ and $Previous_{rot}(i,j)$ (corresponding to the best path up to i and j). Furthermore, considering rotation will also give us a principled way for the path to skip over "bad" onsets, so instead of assuming that every path reaches onset i by way of onset $i-1$ we must also keep track of $Previous_{onset}(i,j)$.

The key improvement in this algorithm is the handling of rotation. Rotation (which indexes the notes in the clave pattern) is converted to *phase*, the proportion (from 0 to 1) of the distance from one downbeat to the next. (So the phases for the notes of rumba clave are [0, 0.1875, 0.4375, 0.625, 0.75]). The key idea is predicting what the phase of the next note "should be": Given phase ϕ_1 and tempo j_1 for onset i_1, a candidate tempo j_2 for onset i_2, and the time between onsets $\Delta T = t_2 - t_1$, and assuming linear interpolation of

	LP	CB	CH	PD	LPWT	PDWT
RB	40	26	22.9	63.1	39.96	62.7
RA	1.75	11	1.54	3.10	2.043	57.9

Table 1. RMS (in BPM) results for tempo curve estimation

tempo during the (short) time between these nearby onsets, we can use the fact that tempo (beat frequency) is the derivative of phase to estimate the phase $\hat{\phi}_2$:

$$\hat{\phi}_2 = \phi_1 + \Delta T((j_1 + j_2)/2)/(4 \times 60) \quad (6)$$

Dividing by 4×60 converts from BPM to bars per second.

Now we can add an extra term to our cost function to express the difference between the predicted phase $\hat{\phi}_2$ and the actual phase ϕ_2 corresponding to the rotation of whatever template we're considering for the onset at time t_2 (being careful to take this difference modulo 1, so that, e.g., the difference between 0.01 and .98 is only 0.03, not 0.97). We'll call this phase distance the "phase residual" R, and add the term $\alpha * R$ to our cost function.

Now let's consider how to handle "false" detected onsets, i.e., onsets that are not actually notes of clave. For onset n, we consider not just onset $n-1$ as the previous onset, but every onset i with $t_i > t_n - K$, i.e., every onset within K seconds before onset n, where K is set heuristically to 1.5 times the largest time between notes of clave (one beat) at the slowest tempo. We introduce a "skipped onset cost" β and include $\beta \times (n - i - 2)$ in the path cost when the path goes from onset i to onset n.

Table 1 shows the Root-mean-square (RMS) error between the ground truth tempocurve and the tempocurves estimated by the rotation-blind (RB) and rotation-aware (RA) configurations of our method. In all cases the rotation-aware significantly outperforms the rotation-blind method (which usually tracks correctly only parts of the tempo curve). The first three recordings (LP, CB, CH) have rumba-clave and the fourth piece (PD) has son-clave. The last two columns show the results when using the "wrong" template. Essentially when the template is not correct the matching cost of the beat path is much higher and the tempo curve estimation is wrong. Figure 5 shows the score grid for the rotation-blind (top) and rotation-aware (bottom) configurations overlaid with the estimated and ground truth tempocurves.

4 BAR-WRAPPING VISUALIZATION

A performance typically consists of about 625-1000 clave "notes". Simply plotting each point along a linear time axis would require either excessive width, or would make the figure too small to see anything; this motivates bar wrapping. Conceptually, we start by marking each event time (in this case, each detected onset) on a linear time axis. If we imagine this time axis as a strip of magnetic tape holding our recording, then metaphorically we cut the tape just before each downbeat, so that we have 200 short pieces of tape, which we then stack vertically, so that time reads from left to right along each row, and then down to the next row, like text in languages such as English. Each of these "strips" is then stretched horizontally to fill the figure width, adding a tempo curve along the right side to show the original duration of each bar. Figure 6 depicts the times of our detected onsets for *LP* with this technique. The straight lines show the theoretical clave locations. By looking at the figure one can notice that the 5th clave note is consistently slightly later than the theoretical location. This would be hard to notice without precise estimation of the tempocurve.

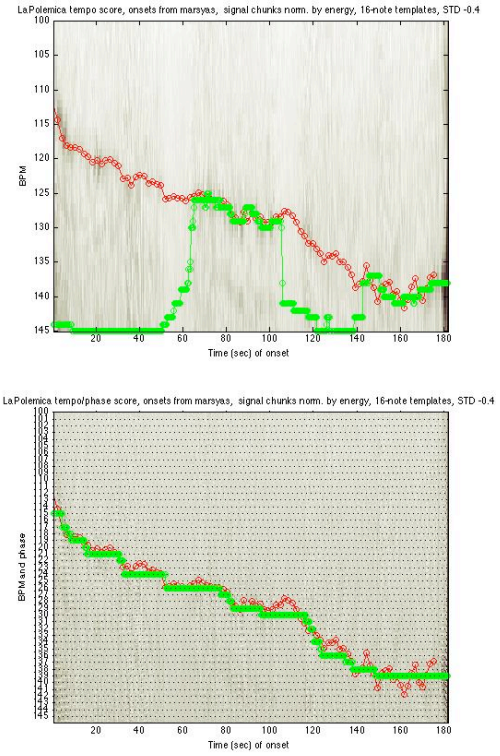

Figure 5. Rotation-blind (top) and rotation-aware (bottom) beat tracking

Rotation-aware dynamic programming is used to find the downbeat times. An explicit downbeat estimate occurs whenever the best path includes a template at rotation 0. But there might not be a detected onset at the time of a downbeat, so we must also consider implicit downbeats, where the current onset's rotation is not 0 but it is lower than the rotation of the previous onset in the best path. The phase is interpolated to estimate the downbeat time that "must have occurred" between the two onsets.

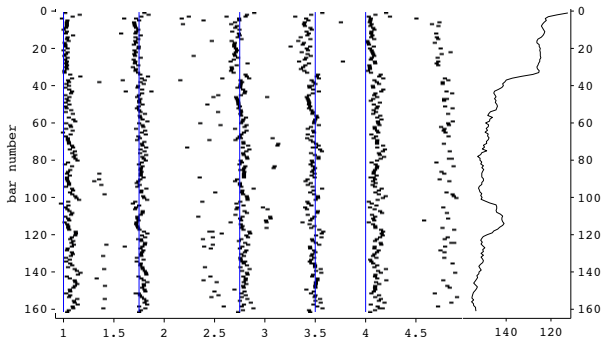

Figure 6. Bar-wrapping visualization

5 DISCUSSION AND CONCLUSIONS

Our beat-tracking method works particularly well for Afro-Cuban clave for many reasons: 1) The clave part almost never stops in traditional Afro-Cuban music (although it can be hard to hear when many other percussion instruments are playing).[3] 2) The clave pattern almost never changes in Afro-Cuban music.[4] 3) The clave instrument produces an extremely consistent timbre with every note, so matched filtering does a good job emphasizing it.[5] 4) Songs often begin with the clave alone, making it easy to construct our matched filter.[6] 5) The clave plays one of a few predetermined syncopated parts, favoring the use of predefined templates rather than assumptions of isochrony.

There are many future work directions. Rhythmic analysis can be used to categorize recordings into different styles and possibly identify particular artists or even percussionists. We also plan to apply the method to more recordings and continue working with ethnomusicologists and performers interested in exploring timing. It is our belief that our template-based rotation-aware formulation can also be applied to popular music by utilizing different standard drum patterns as templates. All the code implementing the method can be obtained by emailing the authors.

[3] Our method's phase- and tempo-continuity constraints allow it to stay on track in the face of extra or missing onsets and occasional unduly low template match scores, so we expect that it would still perform correctly across short gaps in the clave part.

[4] One subtlety of Afro-Cuban music is the notion of "3-2" versus "2-3" clave, which refers to a 180-degree phase shift of the clave part with respect to the ensemble's downbeat. Our method has no notion of the ensemble's downbeat and "doesn't care" about this distinction. Some songs change between 3-2 and 2-3 in the middle, but never by introducing a discontinuity in the clave part (which would be a problem for our algorithm); instead the other instruments generally play a phrase with two "extra" beats that shifts their relationship to the clave.

[5] In rare cases a different instrument carries the clave part; this should not be a problem for our method as long as a relatively isolated sample can be located.

[6] As future work we would like to explore the possibility of creating a "generic" clave enhancement filter that doesn't rely on having an isolated clave note in every recording, a weakness of the current method.

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MULTI-LABEL CLASSIFICATION OF MUSIC INTO EMOTIONS

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ABSTRACT

In this paper, the automated detection of emotion in music is modeled as a multilabel classification task, where a piece of music may belong to more than one class. Four algorithms are evaluated and compared in this task. Furthermore, the predictive power of several audio features is evaluated using a new multilabel feature selection method. Experiments are conducted on a set of 593 songs with 6 clusters of music emotions based on the Tellegen-Watson-Clark model. Results provide interesting insights into the quality of the discussed algorithms and features.

1 INTRODUCTION

Humans, by nature, are emotionally affected by music. Who can argue against the famous quote of the German philosopher Friedrich Nietzsche, who said that "*without music, life would be a mistake*". As music databases grow in size and number, the retrieval of music by emotion is becoming an important task for various applications, such as song selection in mobile devices [13], music recommendation systems [1], TV and radio programs [1] and music therapy.

Past approaches towards automated detection of emotions in music modeled the learning problem as a single-label classification [9, 20], regression [19], or multilabel classification [6, 7, 17] task. Music may evoke more than one different emotion at the same time. We would like to be able to retrieve a piece of music based on any of the associated (classes of) emotions. Single-label classification and regression cannot model this multiplicity. Therefore, the focus of this paper is on multilabel classification methods.

A secondary contribution of this paper is a new multilabel dataset with 72 music features for 593 songs categorized into one or more out of 6 classes of emotions. The dataset is released to the public [2], in order to allow comparative experiments by other researchers. Publicly available multilabel datasets are rare, hindering the progress of research in this area.

[1] http://www.musicovery.com/

[2] http://mlkd.csd.auth.gr/multilabel.html

The primary contribution of this paper is twofold:

- A comparative experimental evaluation of four multilabel classification algorithms on the aforementioned dataset using a variety of evaluation measures. Previous work experimented with just a single algorithm. We attempt to raise the awareness of the MIR community on some of the recent developments in multilabel classification and show which of those algorithms perform better for musical data.
- A new multilabel feature selection method. The proposed method is experimentally compared against two other methods of the literature. The results show that it can improve the performance of a multilabel classification algorithm that doesn't take feature importance into account.

The remaining of this paper is structured as follows. Sections 2 and 3 provide background material on multilabel classification and emotion modeling respectively. Section 4 presents the details of the dataset used in this paper. Section 5 presents experimental results comparing the four multilabel classification algorithms and Section 6 discusses the new multilabel feature selection method. Section 7 presents related work and finally, conclusions and future work are drawn in Section 8.

2 MULTILABEL CLASSIFICATION

Traditional *single-label* classification is concerned with learning from a set of examples that are associated with a single label λ from a set of disjoint labels L, $|L| > 1$. In *multilabel* classification, the examples are associated with a set of labels $Y \subseteq L$.

2.1 Learning Algorithms

Multilabel classification methods can be categorized into two different groups [14]: i) *problem transformation* methods, and ii) *algorithm adaptation* methods. The first group

contains methods that are algorithm independent. They transform the multilabel classification task into one or more single-label classification, regression or ranking tasks. The second group contains methods that extend specific learning algorithms in order to handle multilabel data directly.

2.2 Evaluation Measures

Multilabel classification requires different evaluation measures than traditional single-label classification. A taxonomy of multilabel classification evaluation measures is given in [15], which considers two main categories: *example-based* and *label-based measures*. A third category of measures, which is not directly related to multilabel classification, but is often used in the literature, is ranking-based measures, which are nicely presented in [21] among other publications.

3 MUSIC AND EMOTION

Hevner [4] was the first to study the relation between music and emotion. She discovered 8 clusters of adjective sets describing music emotion and created an emotion cycle of these categories. Hevner's adjectives were refined and regrouped into ten groups by Farnsworth [2].

Figure 1 shows another emotion model, called Thayer's model of mood [12], which consists of 2 axes. The horizontal axis described the amount of stress and the vertical axis the amount of energy.

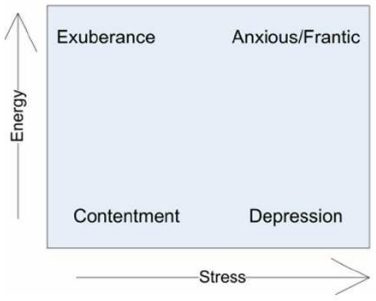

Figure 1. Thayer's model of mood

The model depicted in Figure 2 extends Thayer's model with a second system of axes, which is rotated by 45 degrees compared to the original axes [11]. The new axes describe (un)pleasantness versus (dis)engagement.

4 DATASET

The dataset used for this work consists of 100 songs from each of the following 7 different genres: Classical, Reggae, Rock, Pop, Hip-Hop, Techno and Jazz. The collection was created from 233 musical albums choosing three songs from each album. From each song a period of 30 seconds after the initial 30 seconds was extracted. The resulting sound clips were stored and converted into wave files of 22050 Hz sampling rate, 16-bit per sample and mono. The following subsections present the features that were extracted from each wave file and the emotion labeling process.

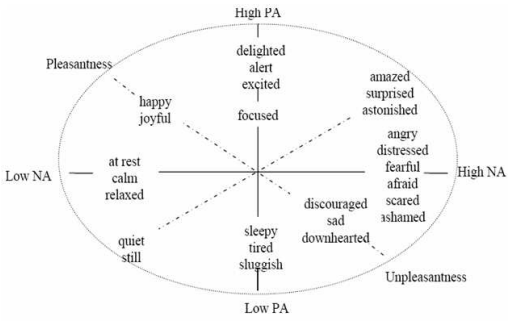

Figure 2. The Tellegen-Watson-Clark model of mood (figure reproduced from [18])

4.1 Feature Extraction

For the feature extraction process, the Marsyas tool [16] was used. The extracted features fall into two categories: rhythmic and timbre.

4.1.1 Rhythmic Features

The rhythmic features were derived by extracting periodic changes from a beat histogram. An algorithm that identifies peaks using autocorrelation was implemented. We selected the two highest peaks and computed their amplitudes, their BMPs (beats per minute) and the high-to-low ratio of their BPMs. In addition, 3 features were calculated by summing the histogram bins between 40-90, 90-140 and 140-250 BPMs respectively. The whole process led to a total of 8 rhythmic features.

4.1.2 Timbre Features

Mel Frequency Cepstral Coefficients (MFCCs) are used for speech recognition and music modeling [8]. To derive MFCCs features, the signal was divided into frames and the amplitude spectrum was calculated for each frame. Next, its logarithm was taken and converted to Mel scale. Finally, the discrete cosine transform was implemented. We selected the first 13 MFCCs.

Another set of 3 features that relate to timbre textures were extracted from the Short-Term Fourier Transform (FFT): Spectral centroid, spectral rolloff and spectral flux.

For each of the 16 aforementioned features (13 MFCCs, 3 FFT) we calculated the mean, standard deviation (std),

mean standard deviation (mean std) and standard deviation of standard deviation (std std) over all frames. This led to a total of 64 timbre features.

4.2 Emotion Labeling

The Tellegen-Watson-Clark model was employed for labeling the data with emotions. We decided to use this particular model because the emotional space of music is abstract with many emotions and a music application based on mood should combine a series of moods and emotions. To achieve this goal without using an excessive number of labels, we reached a compromise retaining only 6 main emotional clusters from this model. The corresponding labels are presented in Table 1.

Label	Description	# Examples
L1	amazed-surprised	173
L2	happy-pleased	166
L3	relaxing-calm	264
L4	quiet-still	148
L5	sad-lonely	168
L6	angry-fearful	189

Table 1. Description of emotion clusters

The sound clips were annotated by three male experts of age 20, 25 and 30 from the School of Music Studies in our University. Only the songs with completely identical labeling from all experts were kept for subsequent experimentation. This process led to a final annotated dataset of 593 songs. Potential reasons for this unexpectedly high agreement of the experts are the short track length and their common background. The last column of Table 1 shows the number of examples annotated with each label.

5 EMPIRICAL COMPARISON OF ALGORITHMS

5.1 Multilabel Classification Algorithms

We compared the following multilabel classification algorithms: binary relevance (BR), label powerset (LP), random k-labelsets (RAKEL) [15] and multilabel k-nearest neighbor (MLkNN) [21]. The first three are problem transformation methods, while the last one is an algorithm adaptation method. The first two approaches were selected as they are the most basic approaches for multilabel classification tasks. BR considers the prediction of each label as an independent binary classification task, while LP considers the multi-class problem of predicting each member of the powerset of L that exists in the training set (see [15] for a more extensive presentation of BR and LP). RAKEL was selected, as a recent method that has been shown to be more effective than the first two [15]. Finally, MLkNN was selected, as a recent high-performance representative of problem adaptation methods [21]. Apart from BR, none of the other algorithms have been evaluated on music data in the past, to the best of our knowledge.

5.2 Experimental Setup

LP, BR and RAKEL were run using a support vector machine (SVM) as the base classifier. The SVM was trained with a linear kernel and the complexity constant C equal to 1. The one-against-one strategy is used for dealing with multi-class tasks in the case of LP and RAKEL. The number of neighbors in MLkNN was set to 10.

RAKEL has three parameters that need to be selected prior to training the algorithm: a) the subset size, b) the number of models and c) the threshold for the final output. We used an internal 5-fold cross-validation on the training set, in order to automatically select these parameters. The subset size was varied from 2 to 5, the number of models from 1 to 100 and the threshold from 0.1 to 0.9 with a 0.1 step.

10 different 10-fold cross-validation experiments were run for evaluation. The results that follow are averages over these 100 runs of the different algorithms.

5.3 Results

Table 2 shows the predictive performance of the 4 competing multilabel classification algorithms using a variety of measures. We notice that RAKEL dominates the other algorithms in almost all measures.

	BR	LP	RAKEL	MLkNN
Hamming Loss	0.1943	0.1964	**0.1845**	0.2616
Micro F1	0.6526	0.6921	**0.7002**	0.4741
Micro AUC	0.7465	0.7781	**0.8237**	0.7540
Macro F1	0.6002	**0.6782**	0.6766	0.3716
Macro AUC	0.7344	0.7717	**0.8115**	0.7185
One-error	0.3038	0.2957	**0.2669**	0.3894
Coverage	2.4378	2.226	**1.9974**	2.2715
Ranking Loss	0.4517	0.3638	0.2635	**0.2603**
Avg. Precision	0.7378	0.7669	**0.7954**	0.7104

Table 2. Performance results

Table 3 shows the cpu time in seconds that was consumed during the training, parameter selection and testing phases of the algorithms. We notice that BR and MLkNN require very little training time, as their complexity is linear with respect to the number of labels. The complexity of LP depends on the number of distinct label subsets that exist in training set, which is typically larger than the number of labels. While the training complexity of RAKEL is bound by the subset size parameter, its increased time comes from

the multiple models that it builds, since it is an ensemble method. RAKEL further requires a comparatively significant amount of time for parameter selection. However, this time is still affordable (2.5 minutes), as it is only run offline.

	BR	LP	RAKEL	MLkNN
Training	0.77	3.07	6.66	0.51
Parameter selection	0	0	151.59	0
Testing	0.00	0.02	0.03	0.06

Table 3. Average training, parameter selection and testing cpu time in seconds

Concerning the test time, we notice that BR is the fastest algorithm, followed by LP and RAKEL. MLkNN is the most time-consuming algorithm during testing, as it must calculate the k nearest neighbors online after the query.

Table 4 shows the classification accuracy of the algorithms for each label (as if they were independently predicted), along with the average accuracy in the last column. We notice that based on the ease of predictions we can rank the labels in the following descending order L4, L6, L5, L1, L3, L2. L4 is the easiest with a mean accuracy of approximately 87%, followed by L6, L5 and L1 with mean accuracies of approximately 80%, 79% and 78% respectively. The hardest labels are L2 and L3 with a mean accuracy of approximately 73% and 76% respectively.

	BR	LP	RAKEL	MLkNN	Avg
L1	0.7900	0.7906	0.7982	0.7446	0.7809
L2	0.7115	0.7380	0.7587	0.7195	0.7319
L3	0.7720	0.7705	0.7854	0.7221	0.7625
L4	0.8997	0.8992	0.9031	0.7969	0.8747
L5	0.8287	0.8093	0.8236	0.7051	0.7917
L6	0.8322	0.8142	0.8238	0.7422	0.8031

Table 4. Accuracy per label

Based on the results, one can see that the classification model performs better for emotional labels such as label 4 (quiet) rather than label 2 (happy). This can be interpreted due to the fact that emotions such as quietness can be easily perceived and classified by humans in a musical context, as it is more objective than more difficult and abstract emotional labels such as happiness, which is more subjective.

6 EMPIRICAL FEATURE EVALUATION

Multilabel feature selection has been mainly applied to the domain of text categorization, due to the typically high dimensionality of textual data. The standard approach is to consider each label separately, use an attribute evaluation statistic (such as χ^2, gain ratio, etc) for each label, and then combine the results using an averaging approach. Two averaging approaches that appear in the literature are *max*, which considers the maximum score for each feature across all labels and *avg*, which considers the average of the score of each feature across all labels, weighted by the prior probability of each label. The disadvantage of these approaches, similarly to BR, is that they do not consider label correlations.

We propose a different approach in this paper, which has not been discussed in the literature before, to the best of our knowledge. At a first step, we apply the transformation of the LP method in order to produce a single-label classification dataset. Then, we apply a common attribute evaluation statistic. We argue that this approach could be more beneficial than the others, as it considers label correlations.

In order to evaluate our hypothesis we compare the Hamming loss of the MLkNN algorithm (known to suffer from the curse of dimensionality) using the best 1 to 71 features according to the three feature selection approaches using the χ^2 statistic. Figure 3 shows the results.

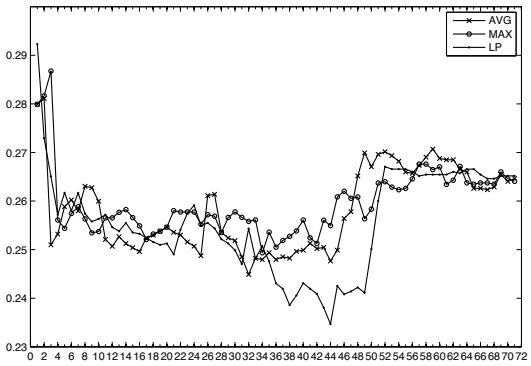

Figure 3. Hamming loss of the MLkNN classifier using the best 1 to 71 features as ordered by the χ^2 feature selection method, using the *max* and *avg* averaging approaches and the proposed method.

We notice that all approaches have similar performance for most of the horizontal axis (number of features retained), apart from the section from 36 to 50 features, where the proposed method leads to better results. It is in this area that the best result is achieved for MLkNN, which is a Hamming loss of approximately 0.235. This is an indication that taking label correlations may be fruitful, especially for selecting important features beyond those that are correlated directly with the labels.

7 RELATED WORK

We discuss past efforts on emotion detection in music, mainly in terms of emotion model, extracted features and the kind

of modeling of the learning problem: a) single label classification, b) regression, and c) multilabel classification.

7.1 Single-label Classification

The four main emotion classes of Thayer's model were used as the emotion model in [9]. Three different feature sets were adopted for music representation, namely intensity, timbre and rhythm. Gaussian mixture models were used to model each of the four classes. An interesting contribution of this work, was a hierarchical classification process, which first classifies a song into high/low energy (vertical axis of Thayer's model), and then into one of the two high/low stress classes.

The same emotion classes were used in [20]. The authors experimented with two fuzzy classifiers, using the 15 features proposed in [10]. They also experimented with a feature selection method, which improved the overall accuracy (around 78%), but they do not mention which features were selected.

The classification of songs into a single cluster of emotions was a new category in the 2007 MIREX (Music Information Retrieval Evaluation eXchange) competition. The top two submissions of the competition [3] were based on support vector machines. The model of mood that was used in the competition, was 5 clusters of moods proposed in [5], which was compiled based on a statistical analysis of the relationship of mood with genre, artist and usage metadata. Among the many interesting conclusion of the competition, was the difficulty to discern between certain clusters of moods, due to their semantic overlap. A multilabel classification approach could overcome this problem, by allowing the specification of multiple finer-grain emotion classes.

7.2 Regression

Emotion recognition is modeled as a regression task in [19]. Volunteers rated a training collection of songs in terms of arousal and valence in an ordinal scale of 11 values from -1 to 1 with a 0.2 step. The authors then trained regression models using a variety of algorithms (again SVMs perform best) and a variety of extracted features. Finally, a user could retrieve a song by selecting a point in the two-dimensional arousal and valence mood plane of Thayer.

Futhermore, the authors used a feature selection algorithm, leading to an increase of the predictive performance. However, it is not clear if the authors run the feature selection process on all input data or on each fold of the 10-fold cross-validation used to evaluate the regressors. If the former is true, then their results may be optimistic, as the feature selection algorithm had access to the test data. A similar pitfall of feature selection in music classification is discussed in [3].

[3] http://www.music-ir.org/mirex/2007

7.3 Multilabel Classification

Both regression and single-label classification methods suffer from the same problem: No two different (clusters of) emotions can be simultaneously predicted. Multilabel classification allows for a natural modeling of this issue.

Li and Ogihara [6] used two emotion models: a) the 10 adjective clusters of Farnsworth (extended with 3 clusters of adjectives proposed by the labeler) and b) a further clustering of those into 6 super-clusters. They only experimented with the BR multilabel classification method using SVMs as the underlying base single-label classifier. In terms of features, they used Marsyas [16] to extract 30 features related to the timbral texture, rhythm and pitch. The predictive performance was low for the clusters and better for the super-clusters. In addition, they found evidence that genre is correlated with emotions.

In an extension of their work, Li and Ogihara [7] considered 3 bipolar adjective pairs (Cheerful vs Depressing), (Relaxing vs Exciting), and (Comforting vs Disturbing). Each track was initially labeled using a scale ranging from -4 to +4 by two subjects and then converted to a binary (positive/negative) label. The learning approach was the same with [6]. The feature set was expanded with a new extraction method, called Daubechies Wavelet Coefficient Histograms. The authors report an accuracy of around 60%.

The same 13 clusters as in [6] were used in [17], where the authors modified the k Nearest Neighbors algorithm in order to handle multilabel data directly. They found that the predictive performance was low, too.

Compared to our work, none of the three aforementioned approaches discusses feature selection from multilabel data, compares different multilabel classification algorithms or uses a variety of multilabel evaluation measures in its empirical study.

8 CONCLUSIONS AND FUTURE WORK

The task of multi-label mapping of music into emotions was investigated. An evaluation of four multi-label classification algorithms was performed on a collection of 593 songs. Among these algorithms, RAKEL was the most effective and is proposed for emotion categorization. The overall predictive performance was high and encourages further investigation of multilabel methods. The performance per each different label varied. The subjectivity of the label may be influencing the performance of its prediction.

In addition, a new multilabel feature ranking method was proposed, which seems to perform better than existing methods in this domain. Feature ranking may assist researchers working on feature extraction by providing feedback on the predictive performance of current and newly designed individual features. It also improves the performance of multilabel classification algorithms, such as MLkNN, that don't

take feature importance into account.

Multilabel classifiers such as RAKEL could be used for the automated annotation of large musical collections with multiple emotions. This in turn would support the implementation of music information retrieval systems that query music collections by emotion. Such a querying capability would be useful for song selection in various applications.

Future work will explore the effectiveness of new features based on time frequency representation of music and lyrics, as well as the hierarchical multilabel classification approach, which we believe has great potential in this domain.

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