MelodyShape: a Library and Tool for Symbolic Melodic Similarity based on Shape Similarity Version 1.0

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Abstract

MelodyShape¹ is a Java library and tool to compute the melodic similarity between monophonic music pieces. It implements several algorithms that compute similarity based on the geometric shape that melodies describe in the pitch-time plane. All these algorithms have obtained the best results in the MIREX² Symbolic Melodic Similarity task in 2010, 2011, 2012 and 2013 editions, as well as the best results reported for the 2005 collection. This document describes the tool and its execution options; for background information the reader is referred to [3].

1 Dependencies

MelodyShape is available both as source code and as an executable JAR package. It requires two libraries found in the Apache Commons project, which are *not* included in the JAR package: commons-cli and commons-math. JAR packages for both libraries can be downloaded from the Apache Commons project website³. MelodyShape v1.0 was specifically implemented and tested with versions commons-cli-1.2 and commons-math3-3.2, also available from the MelodyShape website. Both libraries will have to be accessible from the Java classpath.

The easiest way to run the MelodyShape tool is to download all JAR files in the same directory, and then run Java from the command line:

```
commons-cli-1.2.jar commons-math3-3.2.jar melodyshape-1.0.jar data
$ java -jar melodyshape-1.0.jar
usage: melodyshape-1.0 -q <file/dir> -c <dir> -a <name> [-k <cutoff>] [-1]
                       [-t <num>] [-v] [-vv] [-h]
-q <file/dir> path to the query melody or melodies.
-c <dir>
             path to the collection of documents.
-a <name>
             algorithm to run:
             - 2010-domain, 2010-pitchderiv, 2010-shape
             - 2011-shape, 2011-pitch, 2011-time
             - 2012-shapeh, 2012-shapel, 2012-shapeg, 2012-time, 2012-shapetime
             - 2013-shapeh, 2013-time, 2013-shapetime
-k <cutoff>
             number of documents to retrieve.
-1
             show results in a single line (omits similarity scores).
-t <num>
             run a fixed number of threads.
-17
             verbose, to stderr.
-vv
             verbose a lot, to stderr.
             show this help message.
```

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¹For the latest version of the software and this document, please visit http://julian-urbano.info.

²http://music-ir.org/mirex/wiki/MIREX_HOME

³http://commons.apache.org

```
This program comes with ABSOLUTELY NO WARRANTY.

This is free software, and you are welcome to redistribute it under the terms of the GNU General Public License version 3.
```

Currently, MelodyShape v1.0 is compiled for Java 7. It can be redistributed and/or modified under the terms of the GNU General Public License version 3.

2 Tool Options

2.1 Basic Execution

MelodyShape requires one or more query melodies, a collection of melodies to sort according to their melodic similarity to the queries, and the name of the algorithm to compute those similarities. These three parameters must be indicated with arguments -q, -c and -a in the command line. For instance, we can compute the similarity of melody data/q01.mid with all melodies in directory data/coll/ using algorithm 2010-shape:

The output is a list of all files in data/coll/ sorted by melodic similarity to the query in descending order, along with the computed similarity score. Note that all files must have extension .mid or .midi. In addition, only one MIDI pitch can be on at any point in time.

2.2 Several Queries

MelodyShape can be run several times, once for each of the queries we have. Alternatively, it can be run once for all queries, resulting in a more efficient execution. The following would run MelodyShape will all queries in directory data/4:

```
$ java -jar melodyshape-1.0.jar -q data/ -d data/coll/ -a 2010-shape
q01.mid
          06C6.mid
                     0.48581398
q01.mid
          1065.mid
                      0.16719302
. . .
                     -0.18696516
q01.mid
          0C29.mid
          0859.mid
q02.mid
                     0.44809958
. . .
q06.mid
          OB63.mid
                      -0.19795232
                      -0.22072806
q06.mid
          OEA2.mid
```

Results are again sorted by melodic similarity to the queries, one query after another. The first column identifies the query file, the second column identifies the document and the third column shows the computed similarity score.

2.3 Cutoff: -k

The -k command line argument can be used to indicate how many results to show per query. For example, to show only the two most similar melodies for each query:

```
$ java -jar melodyshape-1.0.jar -q data/ -c data/coll/ -a 2010-shape -k 2
q01.mid
          06C6.mid
                     0.48581398
q01.mid
          1065.mid
                     0.16719302
q02.mid
          0859.mid
                     0.44809958
q02.mid
          03FA.mid
                     0.18398533
q03.mid
          13ED.mid
                     0.35113359
q03.mid
          015D.mid
                     0.32187245
```

⁴Throughout this document, underlined text highlights the changes with the previous executions.

```
q04.mid
          OD1E.mid
                     0.15161275
q04.mid
          OBEA.mid
                     0.14805929
q05.mid
          OB2C.mid
                     0.18844743
q05.mid
          OF60.mid
                     0.13771394
q06.mid
          0859.mid
                     0.26753167
q06.mid
          OB15.mid
                     0.24442237
```

2.4 Single Line: -1

The -1 command line argument indicates that results should be displayed in a single line, one line per query:

```
$ java -jar melodyshape-1.0.jar -q data/ -c data/coll/ -a 2010-shape <u>-k 5 -l</u>
                                 07D0.mid
q01.mid
          06C6.mid
                      1065.mid
                                             OA5A.mid
                                                         04D7.mid
                                                         OB2C.mid
          0859.mid
                      03FA.mid
                                 OB15.mid
                                             0454.mid
q02.mid
. . .
q06.mid
          0859.mid
                      OB15.mid
                                 OFOA.mid
                                             OB2C.mid
                                                         0454.mid
```

Similarly, the first column identifies the query, and the rest identifies the most similar melodies in the collection, from left to right.

2.5 Verbose: -v and -vv

The -v command line argument can be used to display the execution progress, simply showing when each query starts being run:

```
$ java -jar melodyshape-1.0.jar -q data/ -c data/coll/ -a 2010-shape -k 5 -l -v
(1/6) q01.mid...done.
                                07D0.mid
q01.mid 06C6.mid
                                           OA5A.mid
                                                      04D7.mid
                     1065.mid
(2/6) q02.mid...done.
q02.mid 0859.mid
                    03FA.mid
                                OB15.mid
                                           0454.mid
                                                      OB2C.mid
(6/6) q06.mid...done.
q06.mid
         0859.mid
                    OB15.mid
                                OFOA.mid
                                           OB2C.mid
                                                      0454.mid
```

The -vv command line argument can be used to display real-time execution progress, some information about the algorithm used and time spent running each query:

```
$ java -jar melodyshape-1.0.jar -q data/ -c data/coll/ -a 2010-shape -k 5 -l <u>-vv</u>
Reading queries...done (6 melodies).
Reading collection...done (2000 melodies).
Instantiating algorithm...done.
 Comparer: nGram(3, Hybrid(Cache(Freq(BSplineShape(8.0,1.0,0.5)))))
   Ranker: Untie(nGram(3,Hybrid(Eq)))
  Threads: 4
(1/6) q01.mid: [===========] 100% comparing...ranking...done (3 sec).
q01.mid 06C6.mid 1065.mid 07D0.mid
                                        OA5A.mid
                                                  04D7.mid
(2/6) q02.mid: [===========] 100% comparing...ranking...done (2 sec).
                                                 OB2C.mid
                             OB15.mid 0454.mid
q02.mid 0859.mid
                  03FA.mid
(6/6) q06.mid: [=======>
                                      ] 51% comparing...
```

All this extra information is output to the standard error stream, so that the actual results per query can be easily set apart with a pipeline:

```
$ java -jar melodyshape-1.0.jar -q data/ -c data/coll/ -a 2010-shape -l -vv > out
Reading queries...done (6 melodies).
Reading collection...done (2000 melodies).
Instantiating algorithm...done.
Comparer: nGram(3,Hybrid(Cache(Freq(BSplineShape(8.0,1.0,0.5)))))
   Ranker: Untie(nGram(3,Hybrid(Eq)))
   Threads: 4
```

```
(2/6) q02.mid: [===========] 100% comparing...ranking...done (2 sec).
(6/6) q06.mid: [============] 100% comparing...ranking...done (4 sec).
$ cat out
q01.mid
       06C6.mid
                1065.mid
                        07D0.mid
                                         04D7.mid ...
                                OA5A.mid
q02.mid
       0859.mid
                03FA.mid
                        OB15.mid
                                0454.mid
                                         OB2C.mid ...
q06.mid
       0859.mid
                OB15.mid
                        OFOA.mid
                                OB2C.mid
                                         0454.mid ...
```

2.6 Number of Threads: -t

The -t command line argument can be used to indicate the number of threads to use by the algorithms. By default, as many threads as CPU cores found in the machine will be used. We can, for instance, restrict CPU usage to just two threads:

```
\ java -jar melodyshape-1.0.jar -q data/ -c data/coll/ -a 2010-shape \underline{-t\ 2}
```

3 Algorithms

MelodyShape implements several melodic similarity algorithms based on a geometric model that represents melodies as spline curves in the pitch-time plane. The similarity between two melodies is then computed with a sequence alignment algorithm between sequences of spline spans: the more similar the shape of the curves, the more similar the melodies they represent. MelodyShape v1.0 implements algorithms based on this model that have been submitted to the MIREX Symbolic Melodic Similarity task in 2010, 2011, 2012 and 2013.

The following table shows the -a command line argument to be used for each algorithm, with the corresponding official MIREX submission name⁵. For instance, argument -a 2011-shape corresponds to algorithm UL1-Shape from MIREX 2011, which is the same as algorithms JU4-Shape, ULMS1-ShapeH and JU1-ShapeH from MIREX 2010, 2012 and 2013, respectively.

-a argument	MIREX 2010	MIREX 2011	MIREX 2012	MIREX 2013
2010-domain	JU1-Domain			
2010-pitchderiv	JU2-PitchDeriv			
2010-shape	JU4-Shape	UL1-Shape	ULMS1-ShapeH	JU1-ShapeH
2011-shape	JU4-Shape	UL1-Shape	ULMS1-ShapeH	JU1-ShapeH
2011-pitch		UL2-Pitch		
2011-time		UL3-Time	ULMS5-Time	JU3-Time
2012-shapeh	JU4-Shape	UL1-Shape	ULMS1-ShapeH	JU1-ShapeH
2012-shapel			ULMS2-ShapeL	
2012-shapeg			ULMS3-ShapeG	
2012-time		UL3-Time	ULMS5-Time	JU3-Time
2012-shapetime			ULMS4-ShapeTime	JU2-ShapeTime
2013-shapeh	JU4-Shape	UL1-Shape	ULMS1-ShapeH	JU1-ShapeH
2013-time		UL3-Time	ULMS5-Time	JU3-Time
2013-shapetime			ULMS4-ShapeTime	JU2-ShapeTime

The reader is referred to the corresponding MIREX participation report for a detailed description of each of these algorithms [2, 4, 5, 1]. For a general description of the geometric model, the reader is referred to [3].

Acknowledgments

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⁵Algorithm JU3-ParamDeriv from MIREX 2010 is not implemented in MelodyShape v1.0.

References

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