

Evaluation of Chord and Chroma Features and Dynamic Time Warping Scores on Cover Song Identification Task

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Music Information Retrieval

MUSICOLOGY

MUSIC INFORMATION RETRIEVAL

INFORMATICS

MUSIC THEORY

MATHEMATICS

MUSIC ACOUSTIC

PHYSICS

Music Information Retrieval

MUSICOLOGY

INFORMATICS

MUSIC TECHNOLOGY

AUDIO SIGNAL PROCESSING

MUSIC INFORMATION RETRIEVAL

OPTICAL MUSIC SCORE RECOGNITION

MUSIC COGNITION

MUSIC SYNTHESIS & COMPUTER MUSIC

Task – Cover Song Identification

- Cover song
 - An alternative version, performance, or recording of a previously published musical piece



Queen
Somebody To Love



Queen + George Michael
Somebody To Love

Motivation

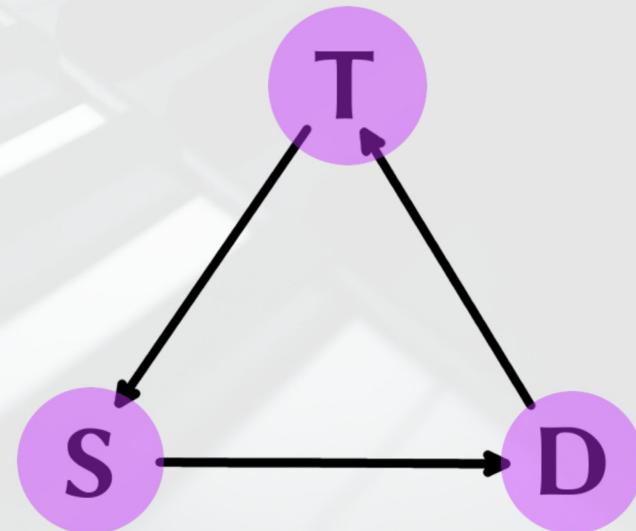
- Contrary to Audio Fingerprinting (e.g. Shazam), **Cover Song Identification** is not widely known and used yet
 - Complexity of the problem, cover song variability
 - State-of-the-art (Serra et al., MIREX 2007, 73.5 % acc.)
 - New impuls needed for CSI
 1. Features comparison
 - We compare harmony features for CSI
 2. Algorithms comparison
 - We compare (Dynamic Time Warping) DTW scores
 3. Large-scale research
 - We form harmony „fingerprints“
 4. New data sets
 - Future work (ISMIR)

Why Music Harmony?

„Most important in music is its harmony.“

Ilja Zeljenka, Slovak Composer

- Tonic
- Subdominant
- Dominant



Demo 1: Simple Harmony Movement

Folk song: Slovenské mamičky



Basic harmonic functions

T - D - T - D

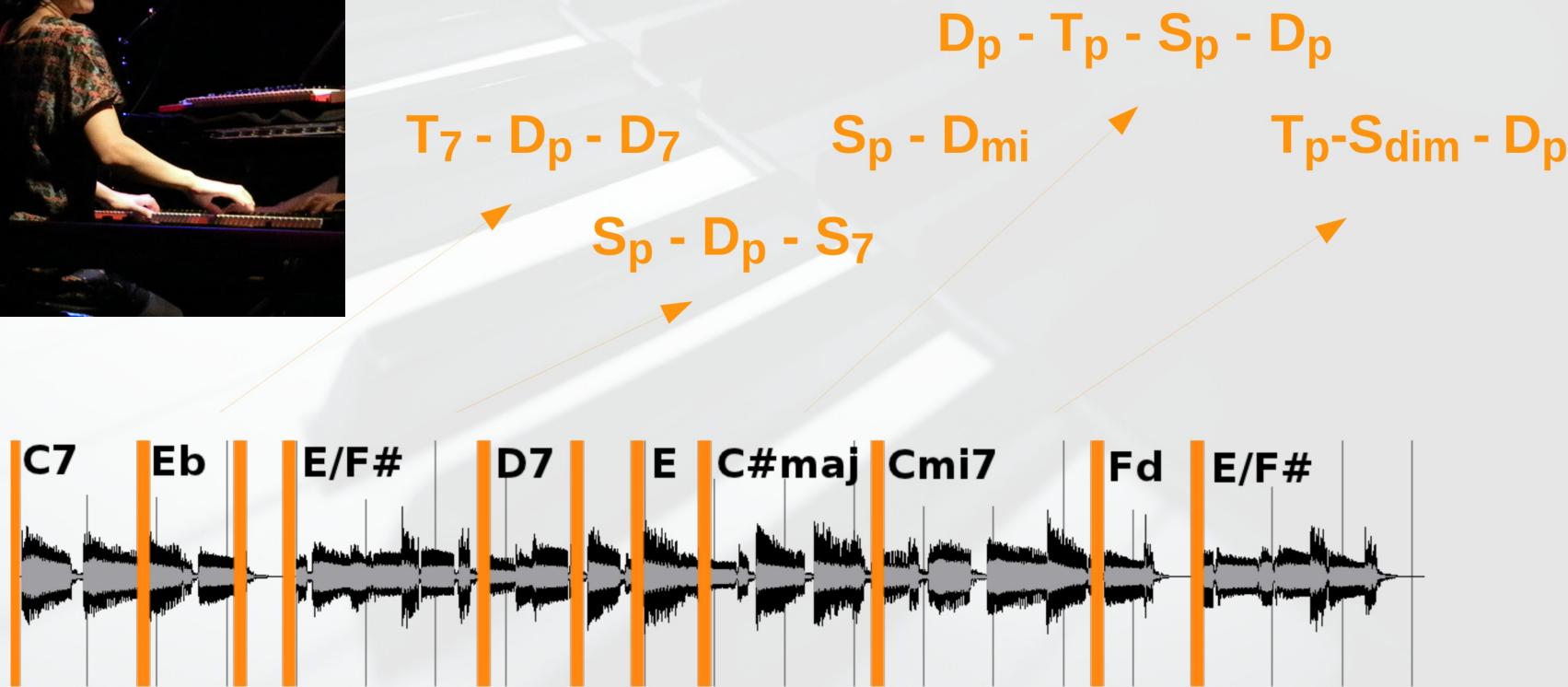


Demo 2: Complex Harmony Movement

Hiromi: 010101 (Binary System)



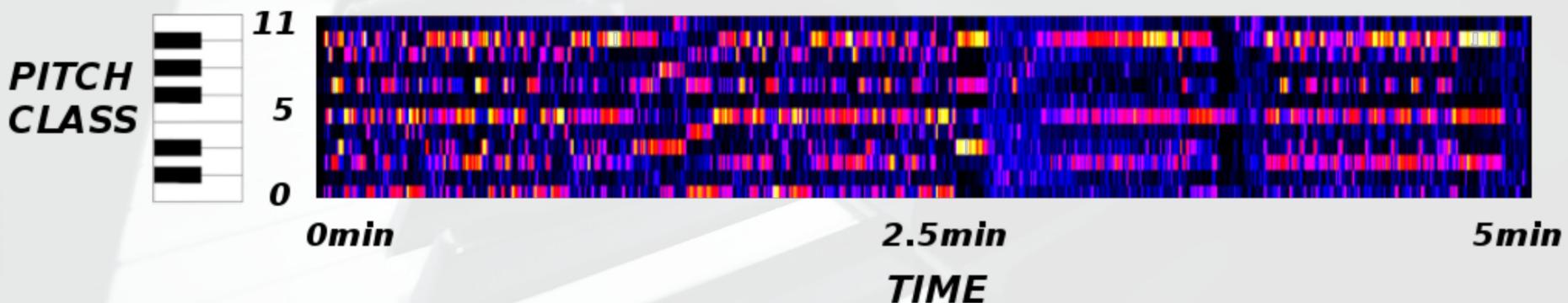
Modifications of basic harmonic functions



Harmony feature 1: Chroma Vectors „raw feature“

Raw Chroma Vectors

George Michael: Somebody To Love (Cover version)

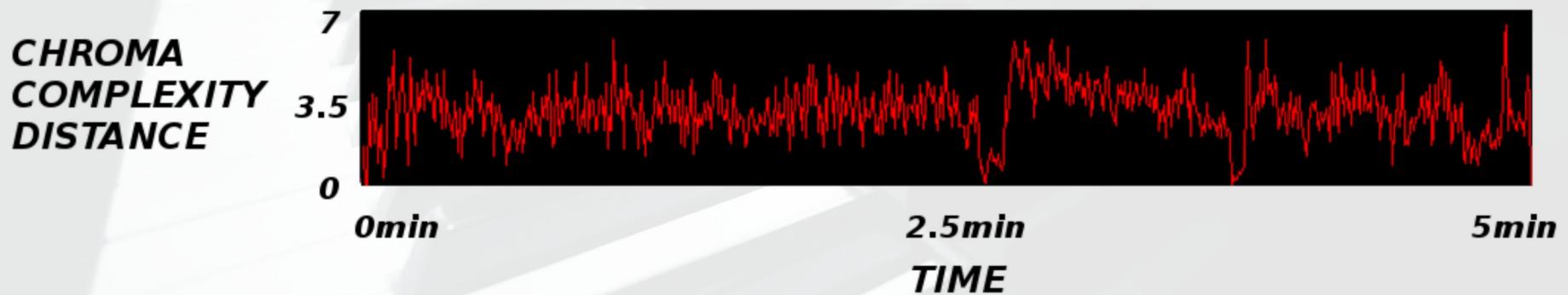


- Frequently used in MIR / Cover Song Identification
- **12-dimensional feature**, fixed sample rate (22Hz)
- Advantage: Proved to contain enough harmony information
- Drawback: Shifting needed (Cover song can be played in another key)

Harmony feature 2: Chroma Distances „fingerprint“

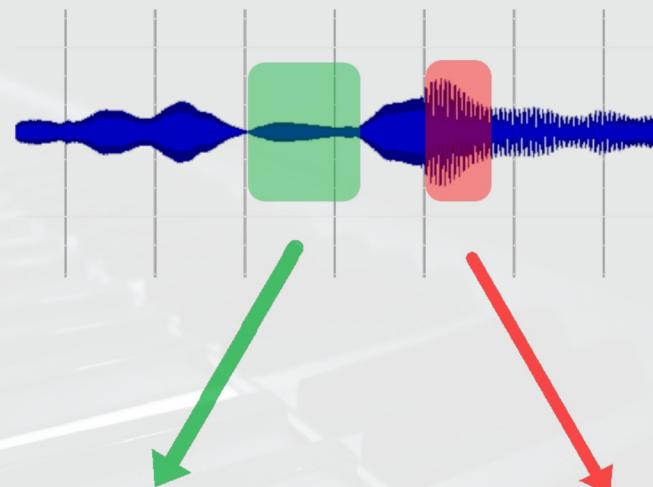
Chroma Complexity Distance (ChromaCD)

George Michael: Somebody To Love (Cover version)



- Idea: A distance derived from a pair of subsequent Chroma vectors
- **1 dimension (scalar)**, fixed sample rate (22Hz)
- Advantage: Reduced size, abstraction from the key, still can detect specific song parts (instrument solos, rhythmic patterns)
- Drawback: Loss of information

In depth: Feature 2 (Chroma Complexity Distance)

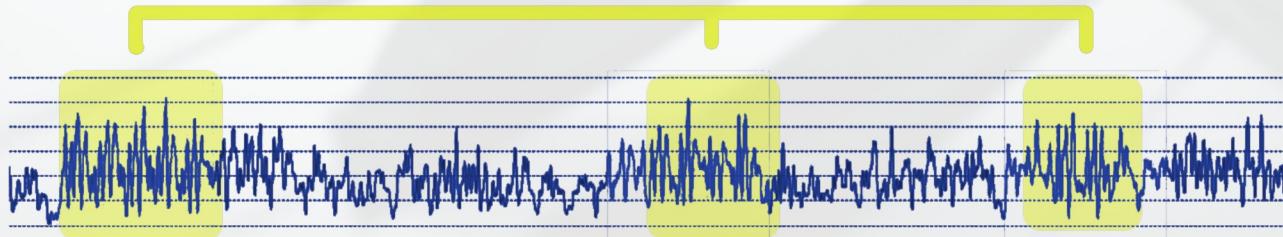


There are no tonal changes

The tonality changes rapidly

Michael Jackson: The Smooth Criminal

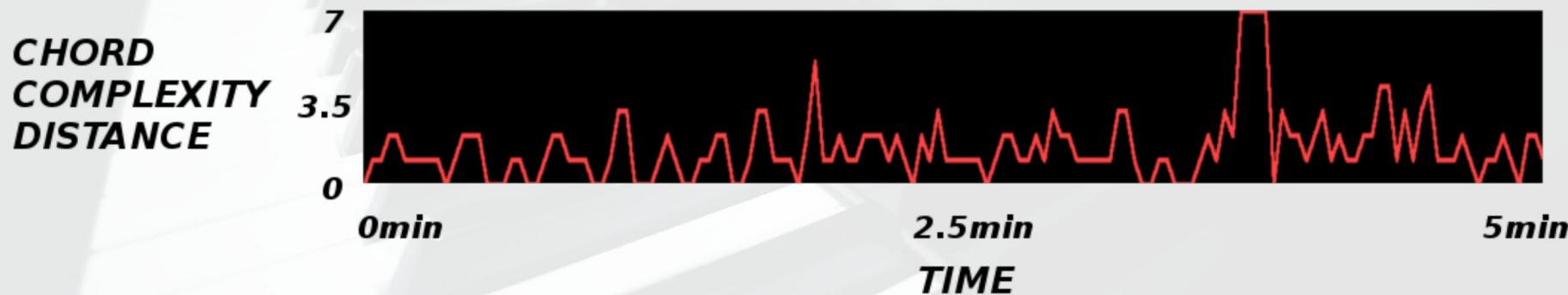
The same guitar riffs



Harmony feature 3: Chord Distances „fingerprint“

Chord Complexity Distance (ChordCD)

George Michael: Somebody To Love (Cover version)



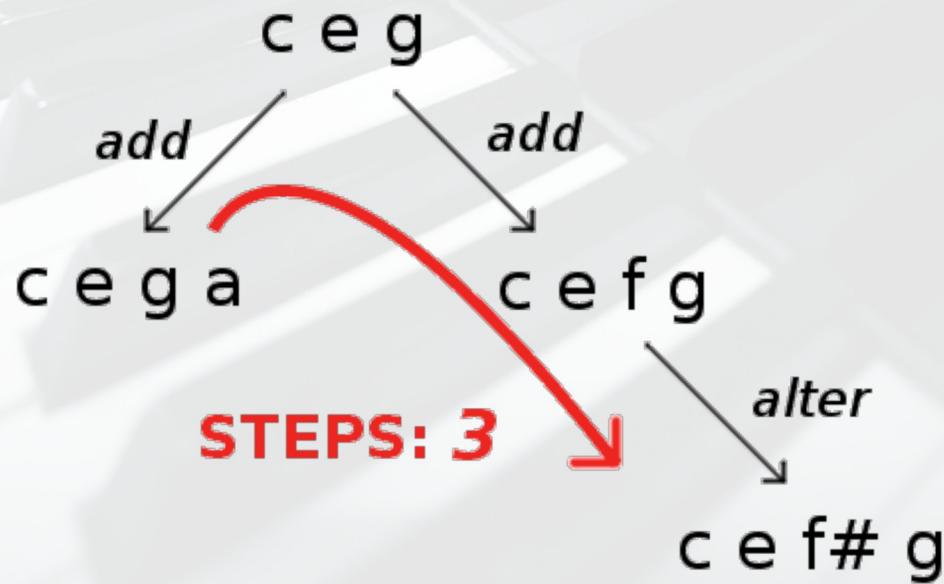
- Idea: A distance derived from a pair of subsequent Chords
- **1 dimension (scalar), chord segmentation detects the new block**
- Advantage: Greatly reduced size, abstraction from the key, abstraction from the specific song parts (only considers harmony)
- Drawback: Great loss of information

In depth: Feature 3 (Chord Complexity Distance)

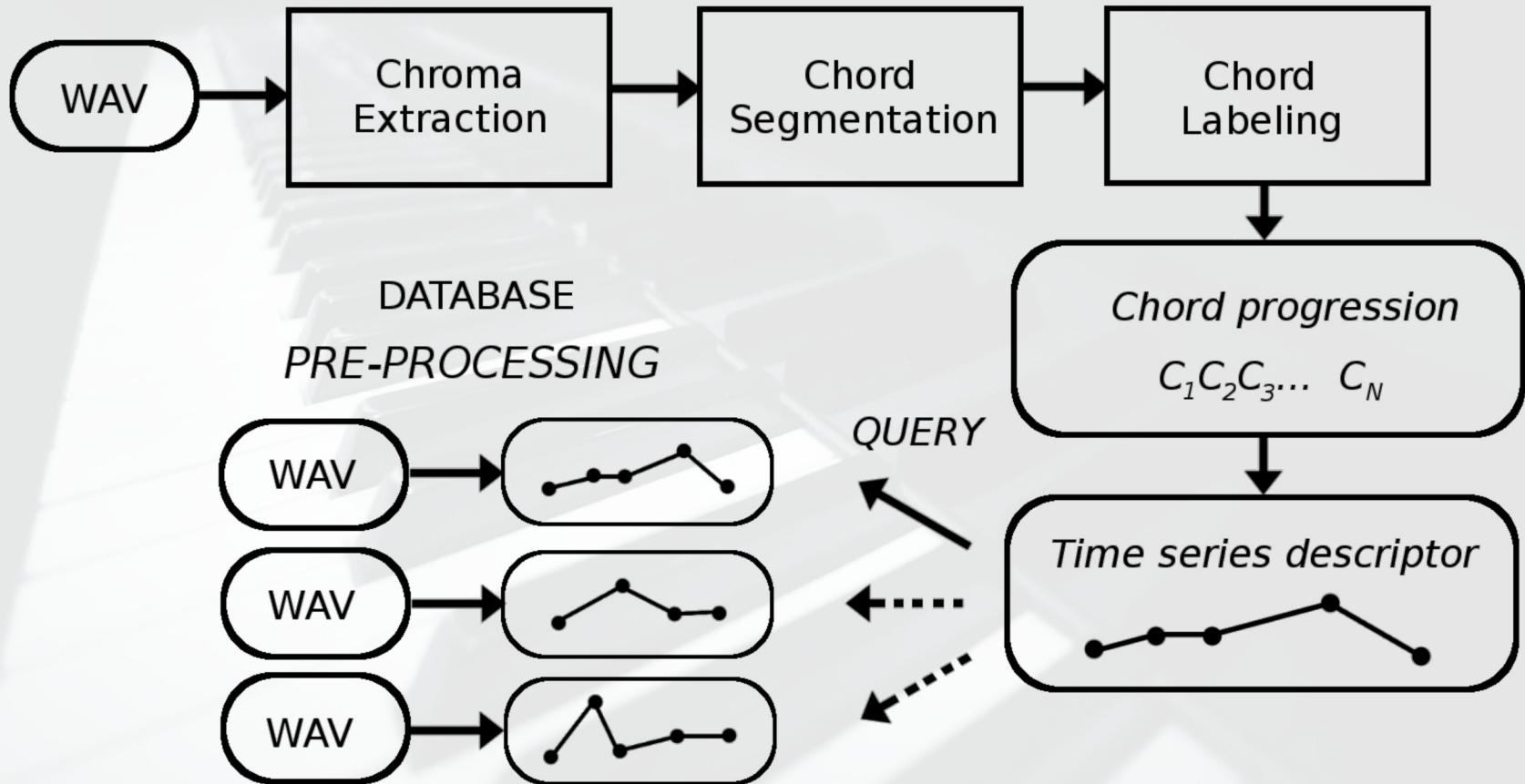
Chord Complexity Distance (ChordCD)

(a variation of Edit Distance)

key: C major

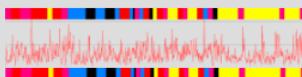


Experiments



All computation were performed on ANSELM supercomputer in **IT4Innovations
national
supercomputing
center**

harmony-analyser



***harmony-analyser is a set of visual tools for music harmony analysis of
WAV/MIDI input, powered by JHarmonyAnalyser library***

The difference we bring is the approach based on music theory, chord and chroma distances. JHarmonyAnalyser uses recent music theory models to extract musical meaning and distances between chords and chroma vectors. We aim to develop open-source music player, which is musician / musicologist-friendly and aid recent music information retrieval tasks.

harmony-analyser tools and JHarmonyAnalyser library are licenced under the [GNU GPL License](#).

Tools are compatible with GPL Licensed [Vamp plugins](#) which can be used for additional analysis.

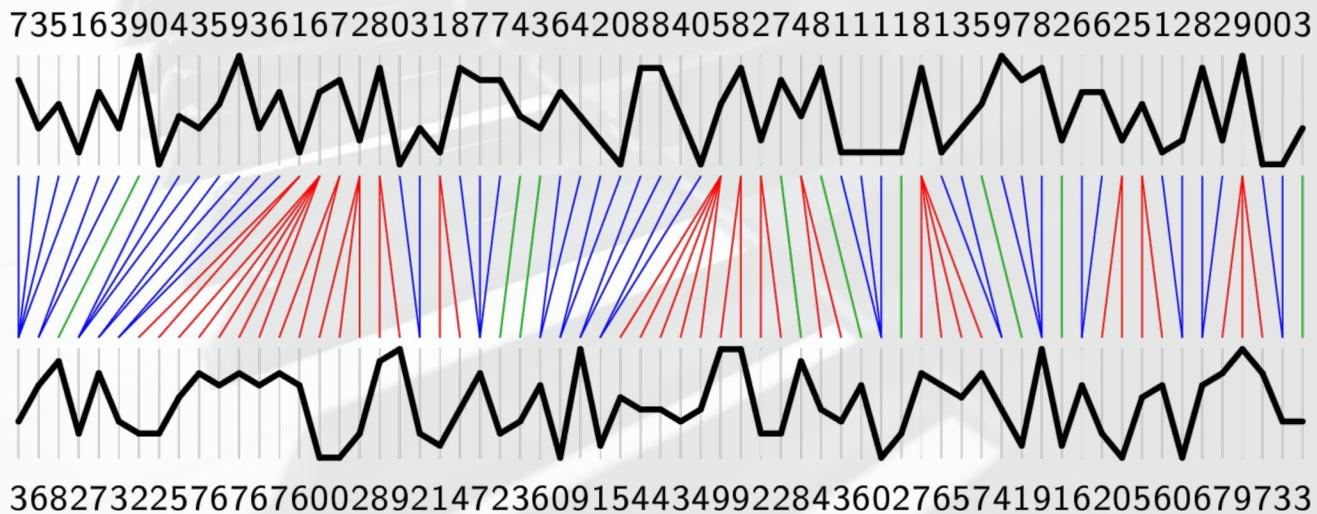
To contribute, please follow our guideline in [GitHub repository](#).

Releases

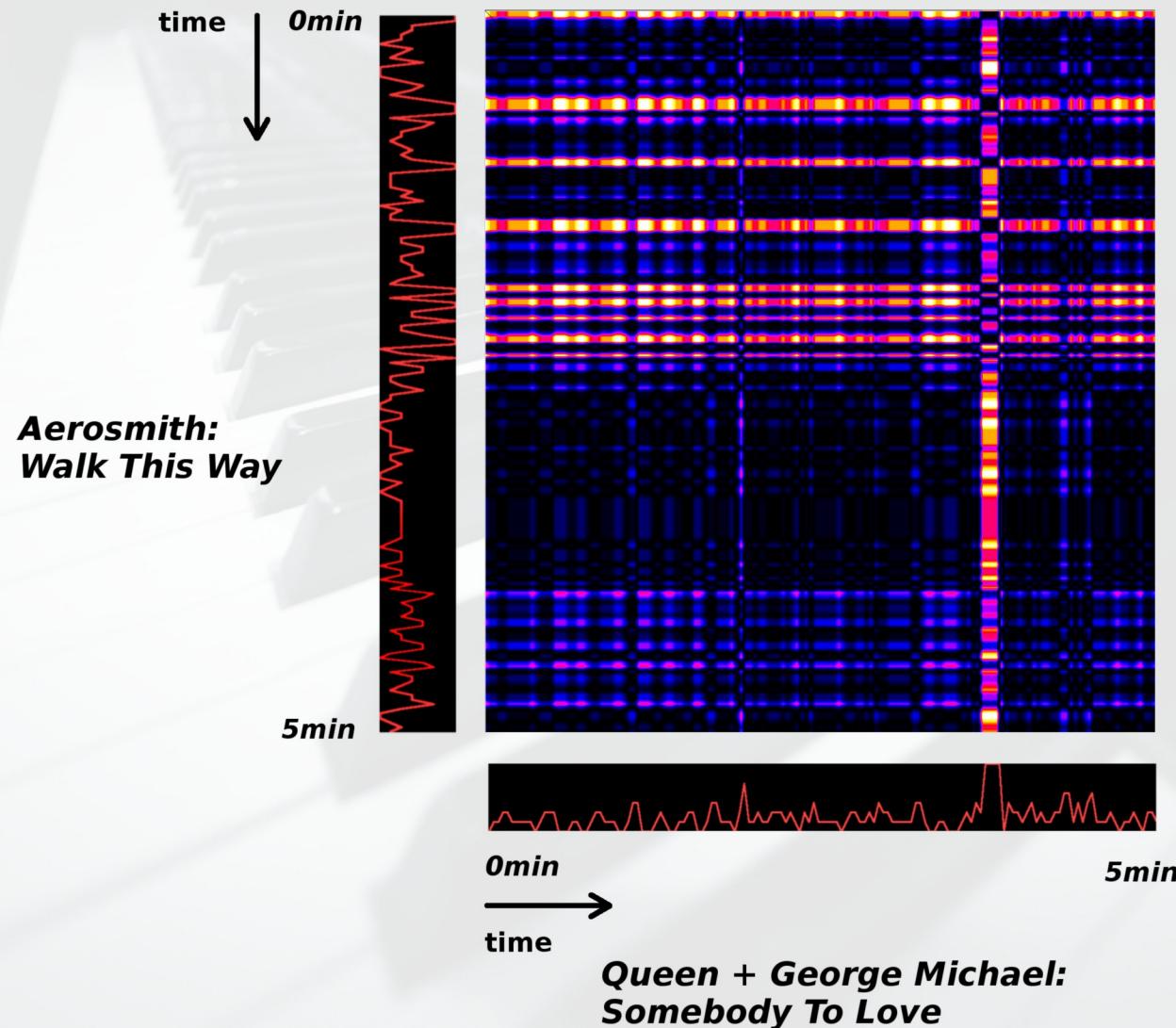
Please choose from the releases below:

Dynamic Time Warping – Idea

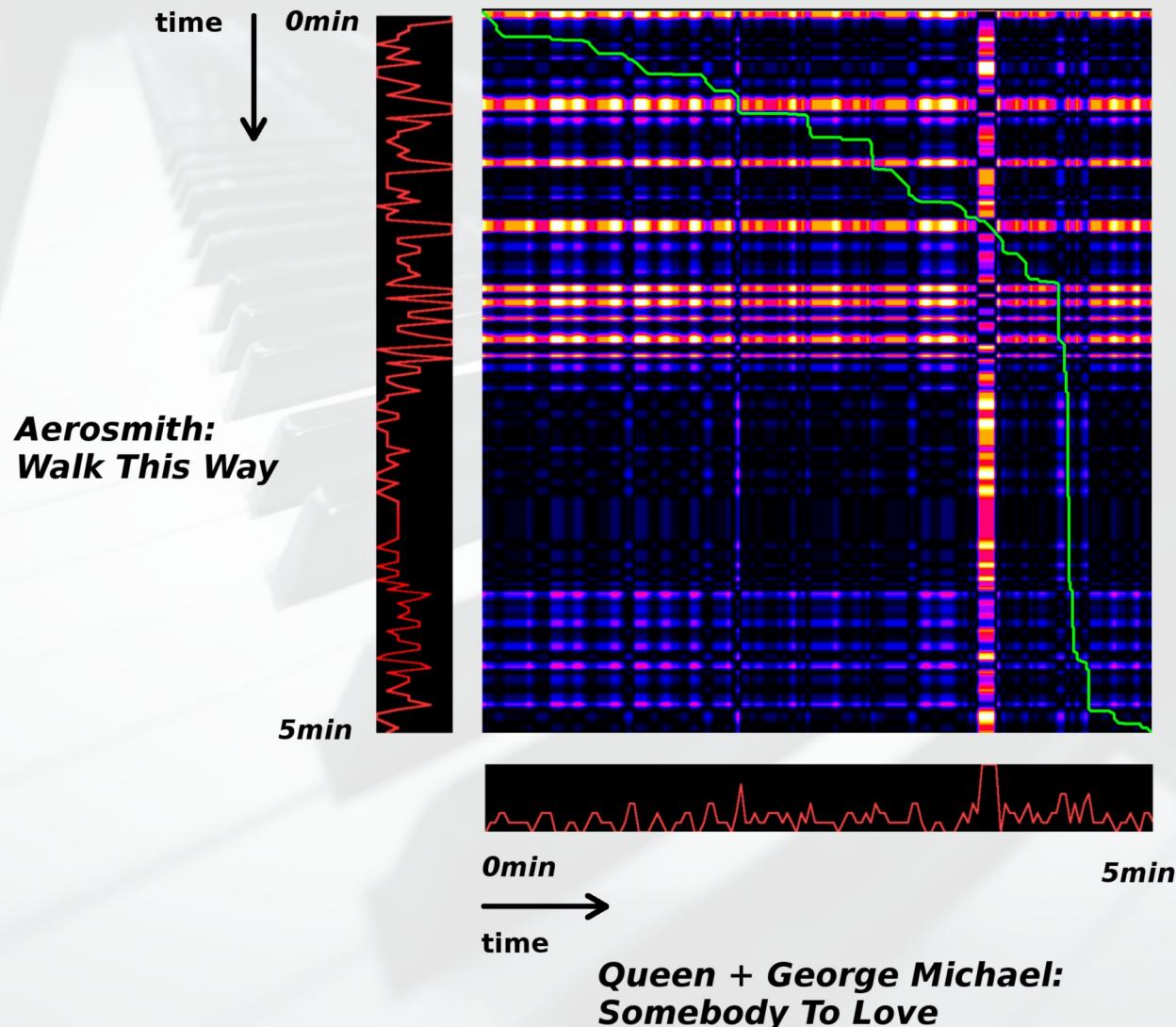
Dynamic Time Warping as alignment of 2 time series



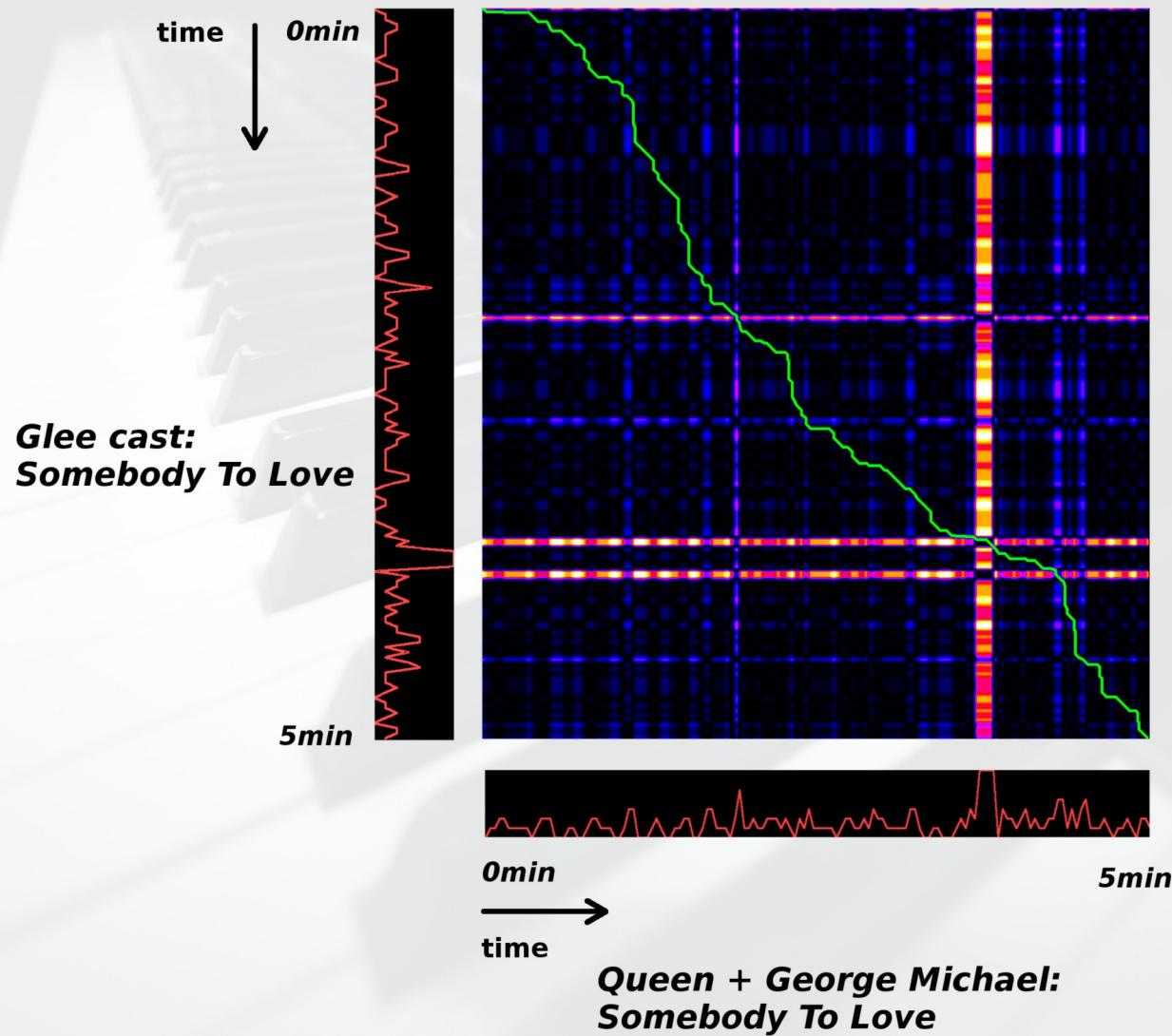
Dynamic Time Warping – Step 1: Distance matrix



Dynamic Time Warping – Step 2: Warping path



Dynamic Time Warping – Step 2: Warping path



Dynamic Time Warping Scores

Score 1 (accumulated distance)

$$DTW(S, T) = \min \left\{ \sqrt{\sum_{l=1}^L w_l} \right.$$

Score 2 (accumulated distance / length)

$$DTW(S, T) = \min \left\{ \frac{1}{L} \sqrt{\sum_{l=1}^L w_l} \right.$$

Score 3 (horizontal/vertical movements)

$$DTW_{sim}(S, T) = \frac{|S| + |T|}{|S| + l + |T| + u}$$

Score 4 (ratio to the maximal DTW)

$$DTW_{sim}(S, T) = 1 - \frac{\sqrt{DTW(S, T)}}{\sqrt{DTW_{max}(S, T)}}$$

Score 5 (ratio to the maximal DTW x differences)

$$DTW_{sim}(S, T) = \left(1 - \frac{\sqrt{DTW(S, T)}}{\sqrt{DTW_{max}(S, T)}} \right) \cdot \frac{\min(|S|, |T|)}{\max(|S|, |T|)}$$

Results of the comparison

	Raw chroma vectors	ChromaCD	ChordCD
AVERAGE RANK	321 / 1000	341 / 1000	403 / 1000
COMPUTATION TIME	550s	411s	100s

Comparison of mean average rank score on SecondHandSongs dataset.

	Raw chroma vectors	ChromaCD	ChordCD
AVERAGE RANK	13.963 / 80	21.538 / 80	26.688 / 80
COMPUTATION TIME	56s	51s	25ms

Comparison of mean average rank score on covers80 dataset.

Results of the comparison

Score	Raw chroma vectors
score (4.1)	0.482
score (4.2)	0.103
score (4.3)	0.417
score (4.4)	0.454
score (4.5)	0.082

MAP results for each DTW score and feature for covers80 dataset.

Score 1 (accumulated distance)

$$DTW(S, T) = \min \left\{ \sqrt{\sum_{l=1}^L w_l} \right\}$$

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$$DTW(S, T) = \min \left\{ \frac{1}{L} \sqrt{\sum_{l=1}^L w_l} \right\}$$

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Conclusion and Future work

1. We have compared harmony features for CSI
 - For future work: „raw“ chroma features show better accuracy
2. We have compared 5 DTW scores for CSI
 - 3 scores stand strong for CSI, 2 lack the accuracy
3. We have proposed harmony fingerprints
 - Can save time for the big data set pre-processing
4. New Data sets for CSI
 - Future work (ISMIR) – karaoke dataset

Thank you for your attention

In depth: Feature 2 (Chroma Complexity Distance)

Chroma Complexity Distance (ChromaCD)

$$\delta(x, y) = \sum_{i=1}^{12} |w(x)_i x_i - w(y)_i y_i|$$

- x, y : 12-dimensional chroma vectors
- $w(x), w(y)$: 12-dimensional weight vectors (the context of x to y transition)
- The weight vectors are dependent on the local chords and key
- Represents how complex is the chroma vector distance in the tonal context

In depth: Feature 2 (Chroma Complexity Distance)

$$\delta(x, y) = \sum_{i=1}^{12} |w(x)_i x_i - w(y)_i y_i|$$

WEIGHT VECTORS

key: C major

PITCH CLASS		chord: C major	chord: C major	chord: G major
		w(x)	w(y)	w(z)
11	B	1	1	0
10	A#	2	2	2
9	A	1	1	1
8	G#	2	2	2
7	G	0	0	0
6	F#	2	2	2
5	F	1	1	1
4	E	0	0	1
3	D#	2	2	2
2	D	1	1	0
1	C#	2	2	2
0	C	0	0	1

Model of Chord Complexity Distance

- Similar to formal grammars
- 2 successive chords in a musical piece

=

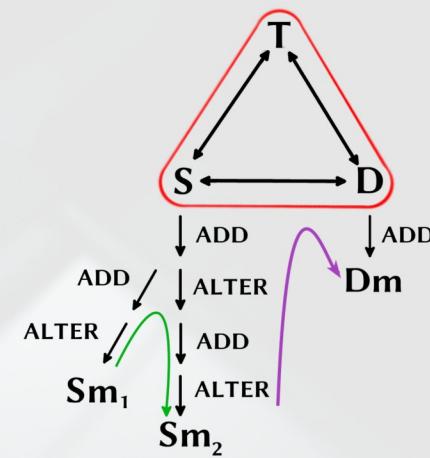
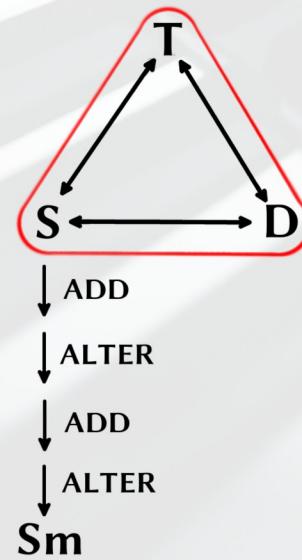
start strign & end string

- Example:

CEG $\xrightarrow{\text{ADD}}$ CEFG $\xrightarrow{\text{ALTER}}$ CEF#G $\xrightarrow{\text{ALTER}}$ CEF#G#

- 2 rules applicable on sentential form:

- ADD – adds a new tone
- ALTER – alters the tone



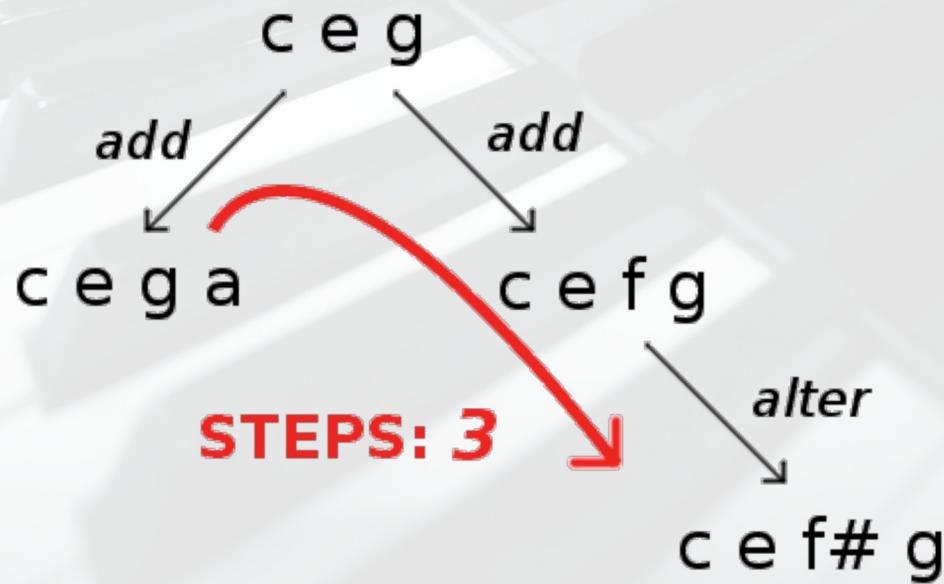
$$tc(Sm_1, Sm_2) = 5 \quad tc(Sm_2, Dm) = 5$$

In depth: Feature 3 (Chord Complexity Distance)

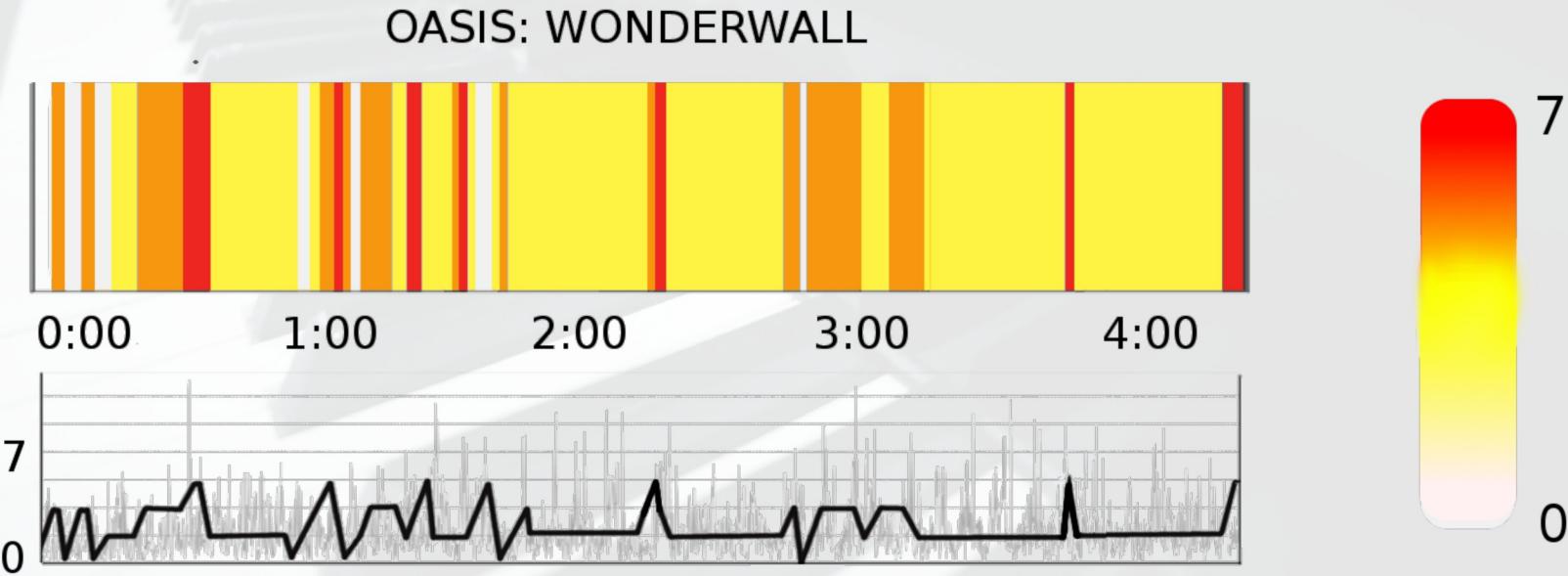
Chord Complexity Distance (ChordCD)

(a variation of Edit Distance)

key: C major



In depth: Feature 3 (Chord Complexity Distance)



In depth: Feature 3 (Chord Complexity Distance)

There are many different Chord Distances, ChordCD is only one possibility

The most common one is: **Tonal Pitch Space** (Fred Lerdahl)

(a)	0							(0)					
(b)	0				7			(0)					
(c)	0		4		7			(0)					
(d)	0	2	4	5	7	9	11	(0)					
(e)	0	1	2	3	4	5	6	7	8	9	10	11	(0)

TPS of C major chord in a C major key

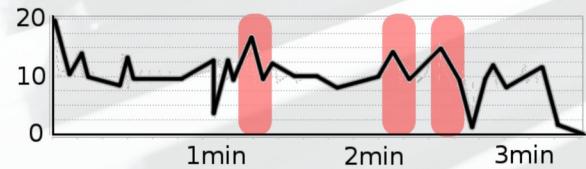
In depth: Feature 3 (Chord Complexity Distance)

ChordCD vs Tonal Pitch Space

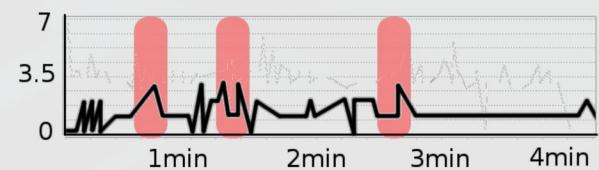
ChordCD



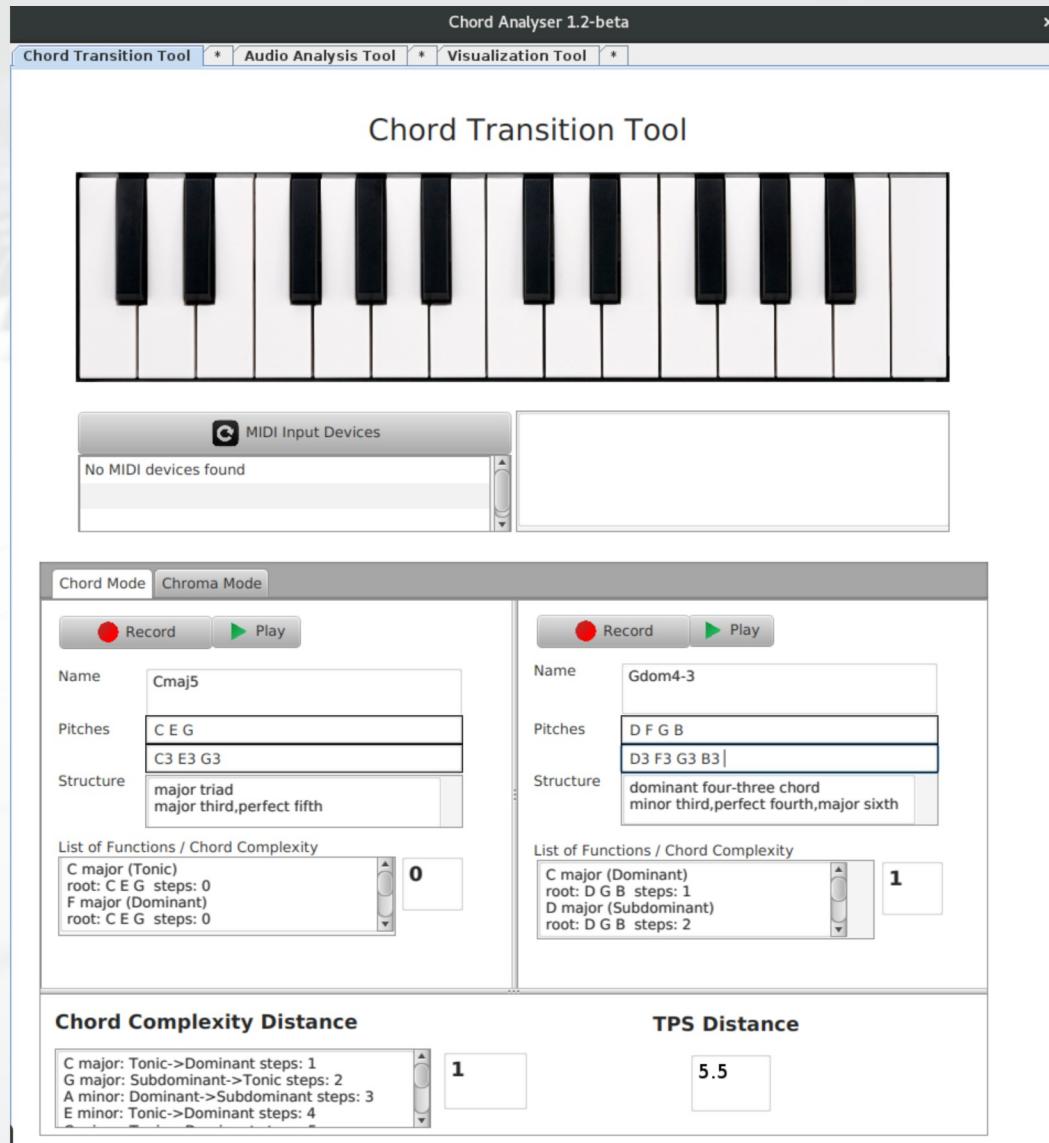
Tonal Pitch Space



Bastian Baker: Hallelujah



Oasis: Wonderwall



Chord Mode Chroma Mode

Record Play

Name	Cmaj5
Pitches	C E G C3 E3 G3
Structure	major triad major third,perfect fifth

List of Functions / Chord Complexity

C major (Tonic) root: C E G steps: 0	0
F major (Dominant) root: C E G steps: 0	▼

Record Play

Name	Gdom4-3
Pitches	D F G B D3 F3 G3 B3
Structure	dominant four-three chord minor third,perfect fourth,major sixth

List of Functions / Chord Complexity

C major (Dominant) root: D G B steps: 1	1
D major (Subdominant) root: D G B steps: 2	▼

Chord Complexity Distance

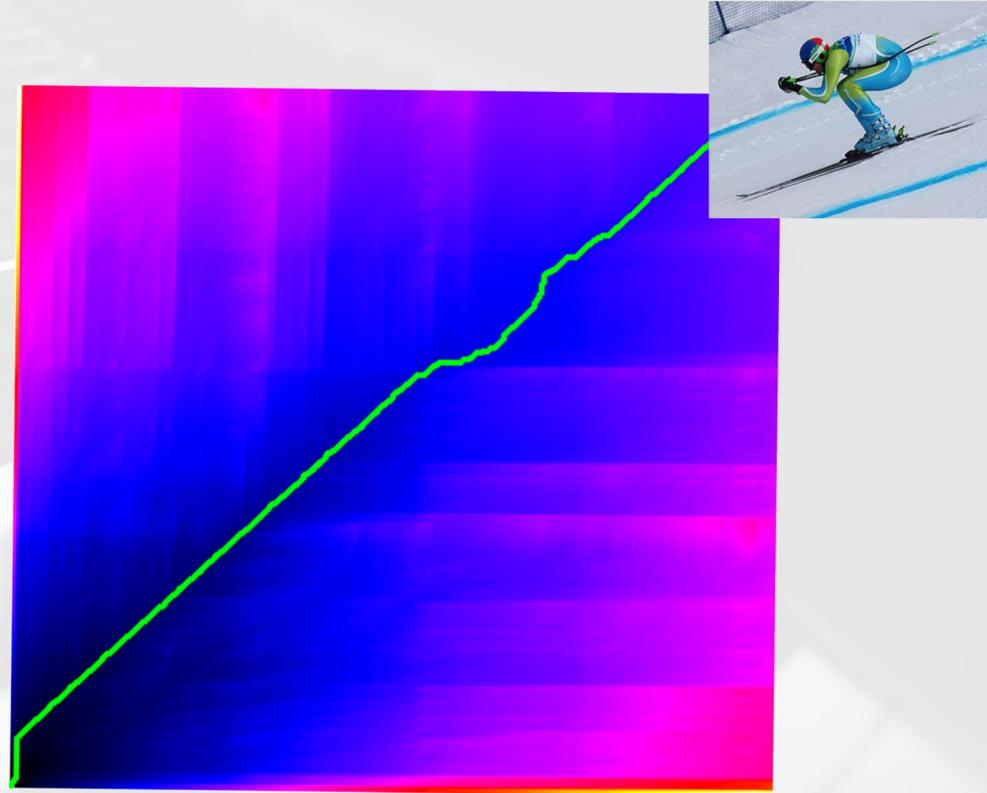
C major: Tonic->Dominant steps: 1 G major: Subdominant->Tonic steps: 2 A minor: Dominant->Subdominant steps: 3 E minor: Tonic->Dominant steps: 4	1
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TPS Distance

5.5

Dynamic Time Warping – Step 2: Warping path

Analogy – „Downhill skiing“

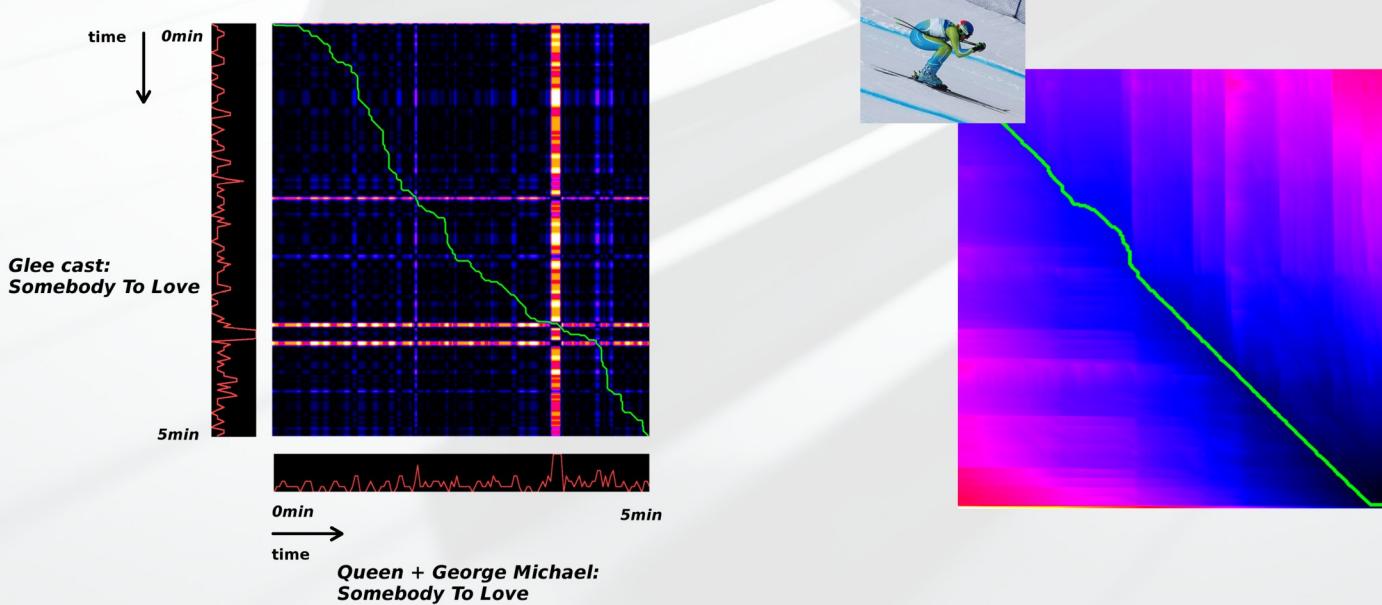


Dynamic Time Warping Scores

Score 1 (accumulated distance)

$$DTW(S, T) = \min \left\{ \sqrt{\sum_{l=1}^L w_l} \right\}$$

L = length of the path; w_l = distance of the step l

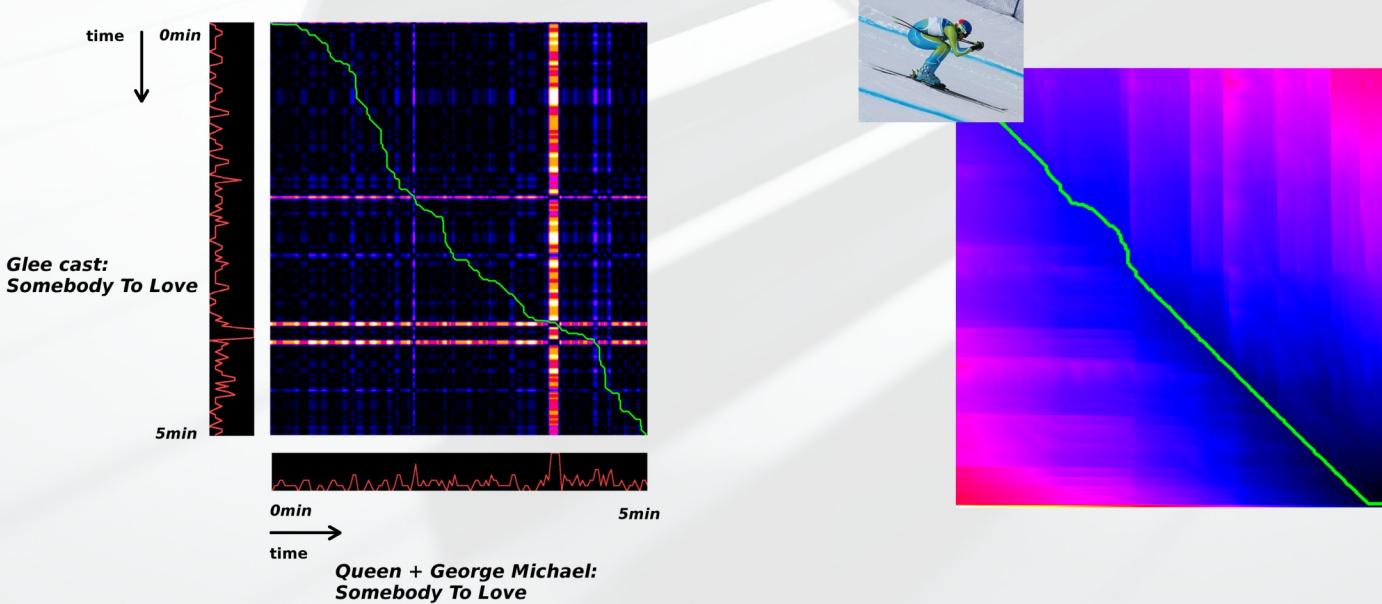


Dynamic Time Warping Scores

Score 2 (accumulated distance / path length)

$$DTW(S, T) = \min \left\{ \frac{1}{L} \sqrt{\sum_{l=1}^L w_l} \right\}$$

L = length of the path; w_l = distance of the step l

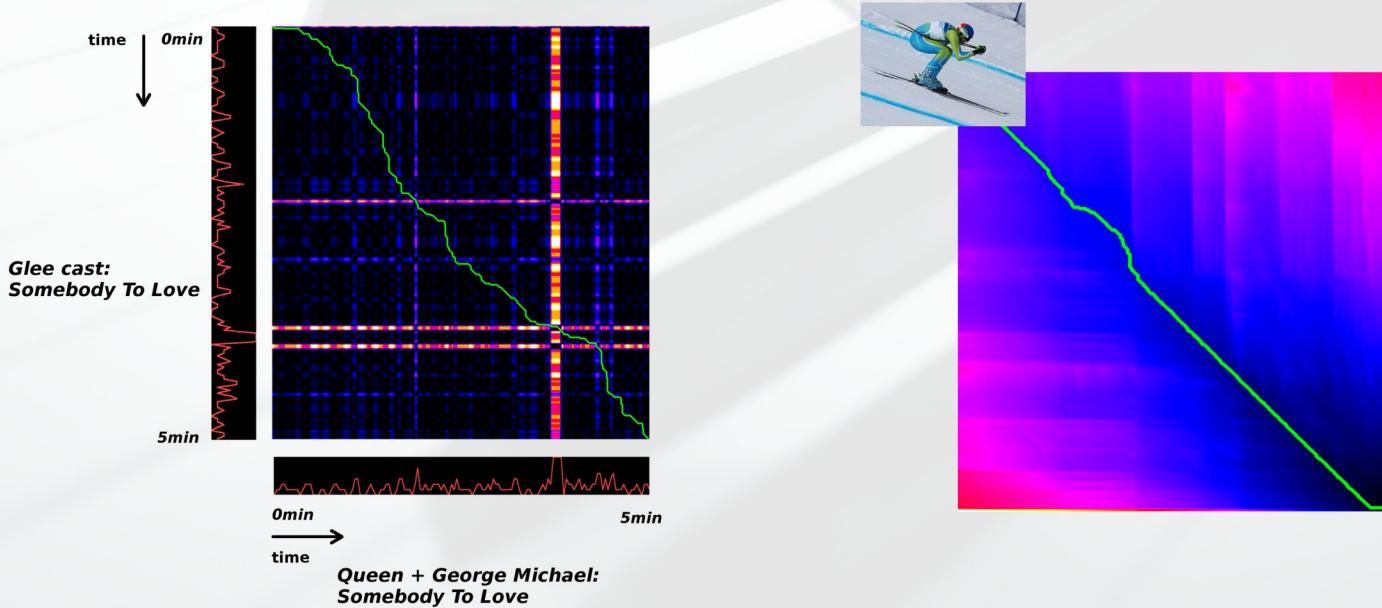


Dynamic Time Warping Scores

Score 3 (horizontal / vertical movements)

$$DTW_{sim}(S, T) = \frac{|S| + |T|}{|S| + l + |T| + u}$$

$|S|, |T|$ = length of series; l, u = number of horizontal / vertical movements

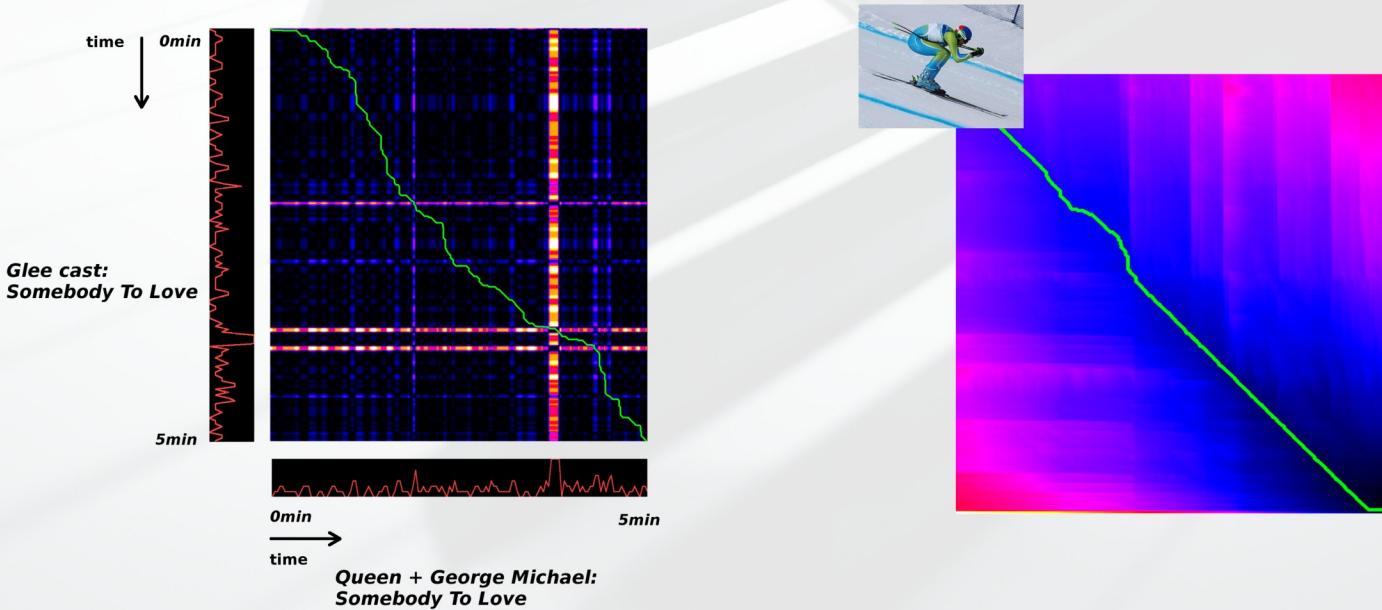


Dynamic Time Warping Scores

Score 4 (ratio to the maximal DTW)

$$DTW_{sim}(S, T) = 1 - \frac{\sqrt{DTW(S, T)}}{\sqrt{DTW_{max}(S, T)}}$$

$DTW_{max}(S, T)$ = maximal theoreticly possible DTW, when S and T are different in all points



Dynamic Time Warping Scores

Score 5 (ratio to the maximal DTW x differences ratio)

$$DTW_{sim}(S, T) = \left(1 - \frac{\sqrt{DTW(S, T)}}{\sqrt{DTW_{max}(S, T)}} \right) \cdot \frac{\min(|S|, |T|)}{\max(|S|, |T|)}$$

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