Methods for Segmentation in Music Structure Analysis (MSA)

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General Goal

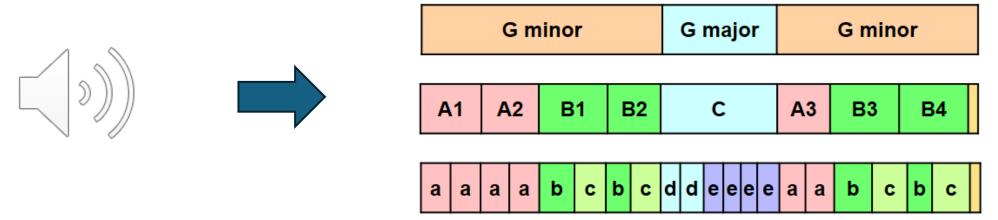


Figure 4.28 from [Müller, FMP, Springer 2015]

Hungarian Dance No. 5 by Johannes Brahms

Problem Definition

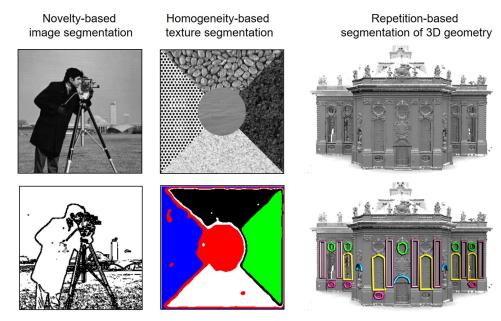
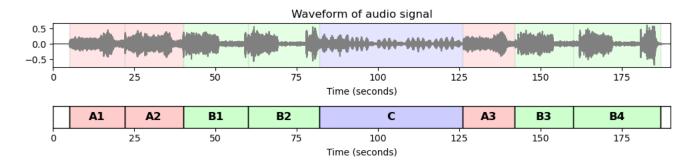


Figure 4.3 from [Müller, FMP, Springer 2015], 3D model by kind permission of [Sunkel et al., CGF, 2011]



Aspects to Consider





Different Approaches

- Homogeneity
- Repetition
- Novelty

Time scale	Dimension	Content
Short term	Timbre	Quality of the produced sound
	Orchestration	Sources of sound production
	Acoustics	Quality of the recorded sound
Middle term	Rhythm	Patterns of sound onsets
	Melody	Sequences of notes
	Harmony	Sequences of chords
Long term	Structure	Organization of the musical work

What we will see

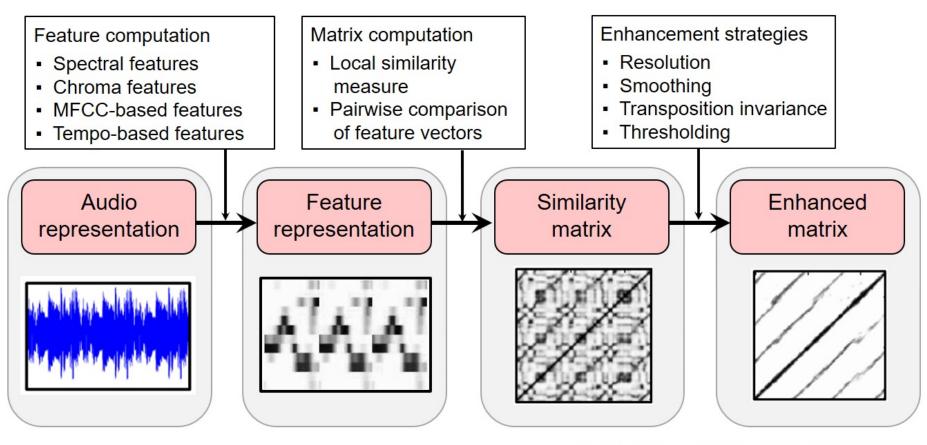


Figure 4.9 from [Müller, FMP, Springer 2015]

Libraries Needed

```
import matplotlib.pyplot as plt
import matplotlib.gridspec as gridspec

from IPython.display import Audio

import numpy as np

import librosa

from libfmp import *
import libfmp.b
import libfmp.c3
import libfmp.c4
import libfmp.c6
```



meinardmueller/ libfmp



libfmp - Python package for teaching and learning Fundamentals of Music Processing (FMP)

Al 5 G 69 1 184 Y 18 Centributors Used by Stars For



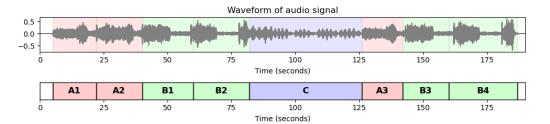
Importing the Audio File with Annotations

```
path = 'Hungarian Dance No. 5.mp3'
x, Fs = librosa.load(path)
x duration = x.shape[0] // Fs
print(f'x.shape: {x.shape}')
print(f'samplerate: {Fs}')
print(f'x.shape / sr: {x duration} seconds')
Audio(x, rate=Fs)
## Plot waveform with segmentation overlay
libfmp.b.plot signal(x, Fs, ax=ax[0], title='Waveform of audio
signal')
libfmp.b.plot segments overlay(ann, ax=ax[0],
time max=x duration,
print labels=False, label ticks=False, edgecolor='gray',
colors = color ann, fontsize=10, alpha=0.1)
## Plot segmentation
libfmp.b.plot segments(ann, ax=ax[1], time max=x duration,
colors=color ann, time label='Time (seconds)')
```

x.shape: (4189500,) samplerate: 22050

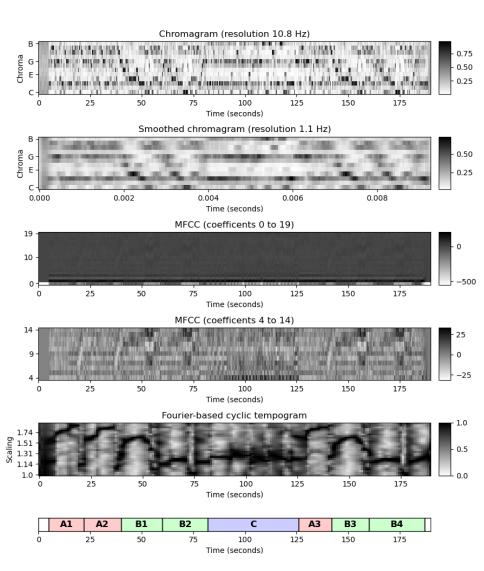
x.shape / sr: 190 seconds





Feature Representations

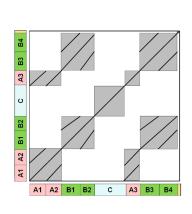
```
----- chromagram ------
chroma = librosa.feature.chroma stft(y=x, sr=Fs,
tuning=0, # A440 tuning
norm=2, # L2 normalization
hop length=H, # temporal resolution
n fft=N # size of FFT window
filt len = 41
down sampling = 10
filt_kernel = np.ones([1, filt_len])
chroma_smoothed = signal.convolve(chroma, filt_kernel, mode='same') / filt_len
chroma smoothed = chroma smoothed[:, ::down sampling]
# ----- mfcc -----
mfcc = librosa.feature.mfcc(y=x, sr=Fs, hop_length=H, n_fft=N)
coef = np.arange(4, 15)
mfcc upper = mfcc[coef, :]
 ----- tempogram ------
nov, sr_nov = libfmp.c6.compute_novelty_spectrum(x, Fs=Fs, N=2048, H=512,
gamma=100, M=10, norm=True)
nov, sr nov = libfmp.c6.resample signal(nov, Fs in=sr nov, Fs out=100)
X, T_coef, F_coef_BPM = libfmp.c6.compute_tempogram_fourier(nov, sr_nov,
N=1000, H=100, Theta=np.arange(30, 601))
```

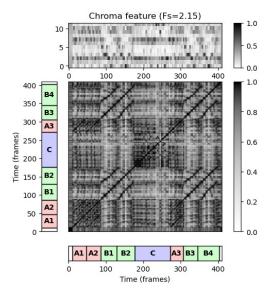


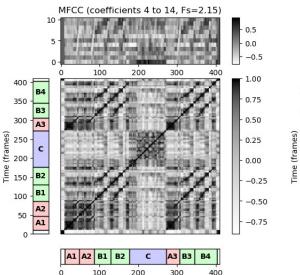
Segmentation Methods

Self-Similarity Matrix

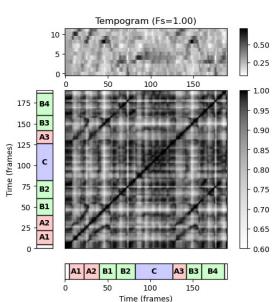
```
# Chroma Feature Sequence
N, H = 4096, 1024
chromagram = librosa.feature.chroma_stft(y=x, sr=Fs, tuning=0, norm=2, hop_length=H, n_fft=N)
X, Fs_X = libfmp.c3.smooth_downsample_feature_sequence(chromagram, Fs/H, filt_len=41, down_sampling=10)
# Annotation
ann_frames = libfmp.c4.convert_structure_annotation(ann, Fs=Fs_X)
# SSM
X = libfmp.c3.normalize_feature_sequence(X, norm='2', threshold=0.001)
S_chroma = compute_sm_dot(X,X)
fig, ax = plot_feature_ssm(X, 1, S_chroma, 1, ann_frames, x_duration*Fs_X, color_ann=color_ann, clim_X=[0,1], clim=[0,1], label='Time (frames)', title='Chroma feature (Fs=%0.2f)'%Fs X)
```







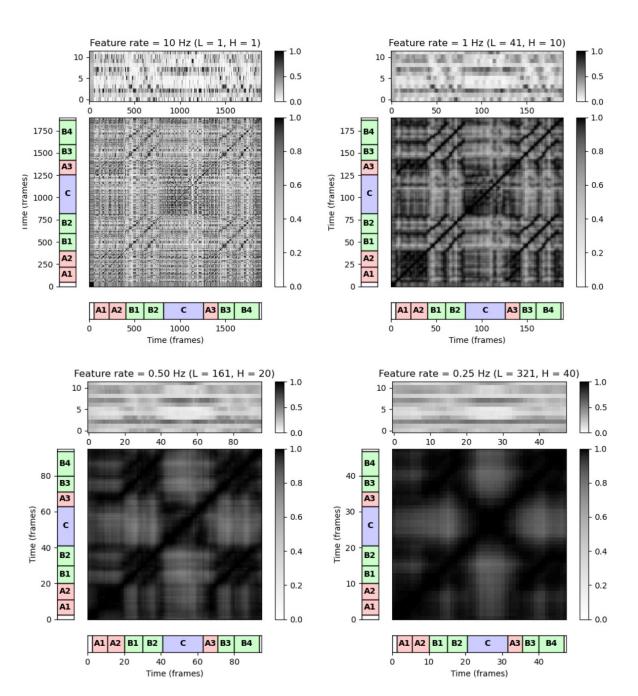
Time (frames)



Path Enhancements

Smoothing

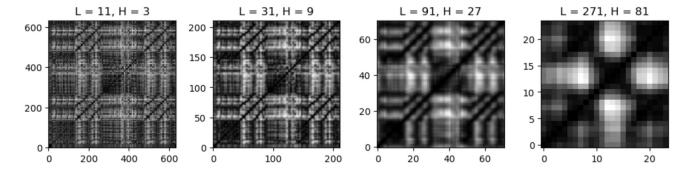
```
C = librosa.feature.chroma_stft(y=x, sr=Fs, tuning=0,
norm=2, hop_length=2205, n_fft=4410)
Fs C = Fs/2205
# Chroma Feature Sequence and SSM (10 Hz)
L, H = 1, 1
X, Fs_feature =
libfmp.c3.smooth downsample feature sequence(C, Fs C,
filt len=L, down sampling=H)
X = libfmp.c3.normalize_feature_sequence(X, norm='2',
threshold=0.001)
S = libfmp.c4.compute sm dot(X,X)
# Chroma Feature Sequence and SSM (1 Hz)
L, H = 41, 10
# Chroma Feature Sequence and SSM (0.5 Hz)
L, H = 161, 20
# Chroma Feature Sequence and SSM (0.25 Hz)
L, H = 321, 40
. . .
```



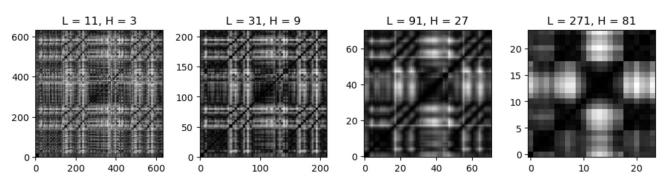
Median Filtering

```
# Chroma Feature Sequence and SSM (0.5 Hz)
L_{iter} = [11, 31, 91, 271]
H_{iter} = [3, 9, 27, 81]
num iter = len(L iter)
print('SSMs obtained using median filtering')
fig = plt.figure(figsize=(10,3))
for i in range(num_iter):
     L = L iter[i]
     H = H iter[i]
     X, Fs feature =
     libfmp.c3.median downsample feature sequence(C,
     Fs_C,
     filt len=L, down sampling=H)
     X = libfmp.c3.normalize_feature_sequence(X,
     norm='2', threshold=0.001)
     S = libfmp.c4.compute sm dot(X,X)
     . . .
```

SSMs obtained using average filtering

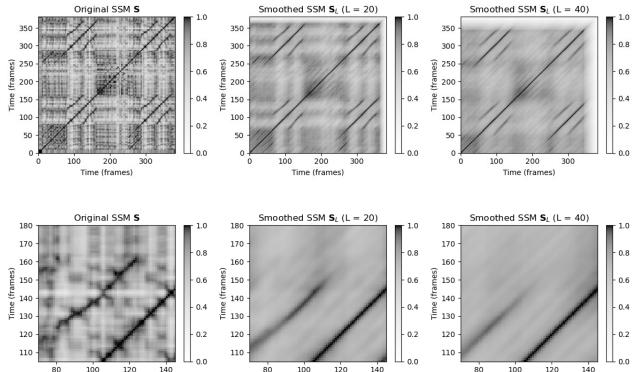


SSMs obtained using median filtering



Diagonal Smoothing

```
def filter diag sm(S, L):
     """Path smoothing of similarity matrix by forward filtering
     along main diagonal
     N = S.shape[0] # Number of rows
     M = S.shape[1] # Number of columns
     S L = np.zeros((N, M))
     S_{extend} L = np_{extend} ((N + L, M + L))
     S extend L[0:N, 0:M] = S # copy original matrix
     for pos in range(0, L):
            # add portion of matrix
           S_L = S_L + S_{extend} L[pos:(N + pos), pos:(M + pos)]
     S_L = S_L / L \# average
     return S L
. . .
L = 20
S_L = filter_diag_sm(S, L)
subplot_matrix_colorbar(S L, fig, ax[1], clim=[0,1], ylabel='Time
(frames)', xlabel='Time (frames)',
title=r'Smoothed SSM $\mathbf{S} {L}$ (L = %d)'%L)
. . .
```



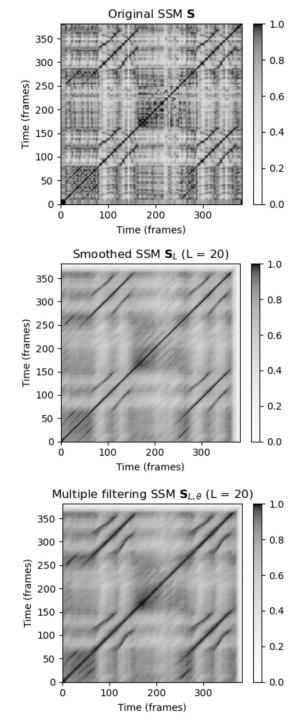
Time (frames)

Time (frames)

Time (frames)

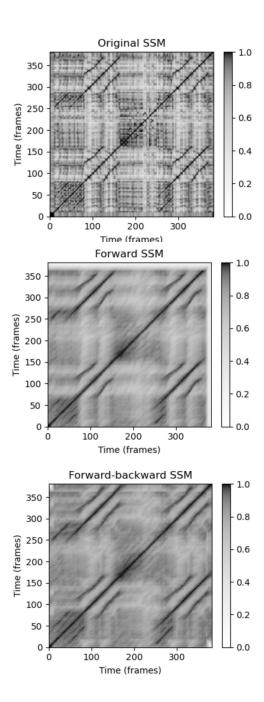
Multiple Filtering

```
def filter_diag_mult_sm(S, L=1, tempo_rel_set=np.asarray([1])):
     N = S_{\bullet} shape[0]
     M = S.shape[1]
     num = len(tempo rel set)
     S L final = np.zeros((N, M))
     for s in range(0, num):
          M_ceil = int(np.ceil(M / tempo_rel_set[s]))
           resample = np.multiply(np.divide(np.arange(1, M_ceil+1), M_ceil), M)
           np.around(resample, 0, resample)
           resample = resample - 1
           index_resample = np.maximum(resample, np.zeros(len(resample))).astype(np.int64)
           S resample = S[:, index resample]
           S_L = np.zeros((N, M_ceil))
           S extend L = np.zeros((N + L, M ceil + L))
           S_extend_L[0:N, 0:M_ceil] = S_resample
           for pos in range(0, L):
                      S L = S L + S extend L[pos:(N + pos), pos:(M ceil + pos)]
           SL = SL/L
           resample = np.multiply(np.divide(np.arange(1, M+1), M), M_ceil)
           np.around(resample, 0, resample)
           resample = resample - 1
           index_resample = np.maximum(resample, np.zeros(len(resample))).astype(np.int64)
           S_resample_inv = S_L[:, index resample]
           S_L_final = np.maximum(S_L_final, S_resample inv)
     return S_L_final
```



Forward-Backward Smoothing

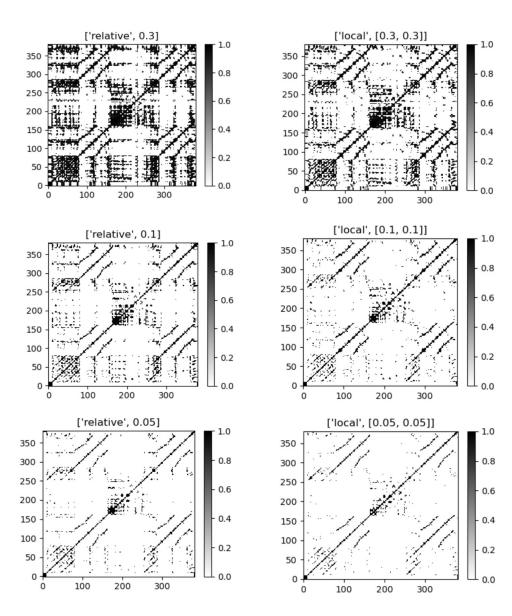
```
def filter diag mult sm(S, L=1, tempo rel set=np.asarray([1]), direction=0):
     N = S_s \text{shape}[0]
     M = S.shape[1]
     num = len(tempo rel set)
     S L final = np.zeros((N, M))
     for s in range(0, num):
           M_ceil = int(np.ceil(M / tempo_rel_set[s]))
           resample = np.multiply(np.divide(np.arange(1, M ceil+1), M ceil), M)
           np.around(resample, 0, resample)
           resample = resample - 1
           index resample = np.maximum(resample, np.zeros(len(resample))).astype(np.int64)
           S resample = S[:, index resample]
           S L = np.zeros((N, M ceil))
           S extend L = np.zeros((N + L, M ceil + L))
           # Forward direction
           if direction == 0:
                      S_extend_L[0:N, 0:M_ceil] = S_resample
                      for pos in range(0, L):
                                 S L = S L + S extend L[pos:(N + pos), pos:(M ceil + pos)]
           # Backward direction
           if direction == 1:
                      S_extend_L[L:(N+L), L:(M_ceil+L)] = S resample
                      for pos in range(0, L):
                                 S_L = S_L + S_{extend}L[(L-pos):(N + L - pos),
                                                         (L-pos):(M ceil + L - pos)]
```



. . .

Thresholding

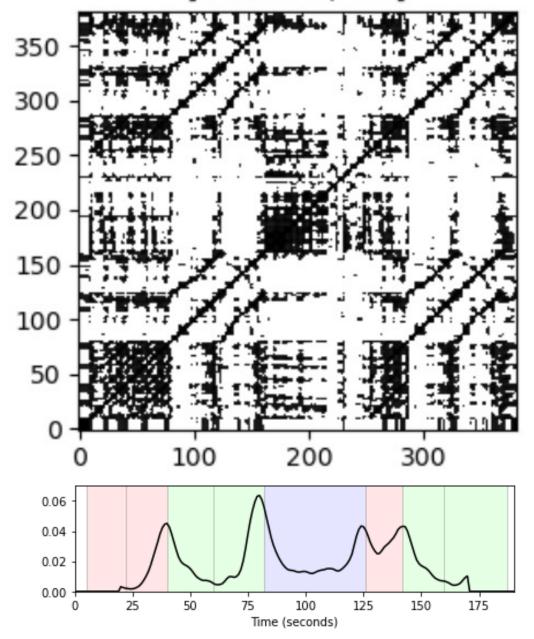
```
if strategy == 'absolute':
      thresh abs = thresh
      S thresh[S thresh < thresh] = 0
if strategy == 'relative':
      thresh rel = thresh
      num cells below thresh = int(np.round(S thresh.size*(1-thresh rel)))
      if num cells below thresh < num cells:</pre>
            values sorted = np.sort(S thresh.flatten('F'))
            thresh abs = values sorted[num cells below thresh]
            S_thresh[S_thresh < thresh abs] = 0
      else:
            S thresh = np.zeros([N, M])
if strategy == 'local':
      thresh rel row = thresh[0]
      thresh rel col = thresh[1]
      S binary row = np.zeros([N, M])
      num cells row below thresh = int(np.round(M * (1-thresh rel row)))
      for n in range(N):
            row = S[n, :]
            values sorted = np.sort(row)
            if num cells row below thresh < M:</pre>
                        thresh abs = values sorted[num cells row below thresh]
                        S_binary_row[n, :] = (row >= thresh_abs)
      S binary col = np.zeros([N, M])
      num_cells_col_below_thresh = int(np.round(N * (1-thresh_rel_col)))
      for m in range(M):
            col = S[:, m]
            values sorted = np.sort(col)
            if num cells col below thresh < N:</pre>
                  thresh abs = values sorted[num cells col below thresh]
                  S binary col[:, m] = (col >= thresh abs)
      S thresh = S * S binary row * S binary col
return S_thresh
```



['relative', 0.3]

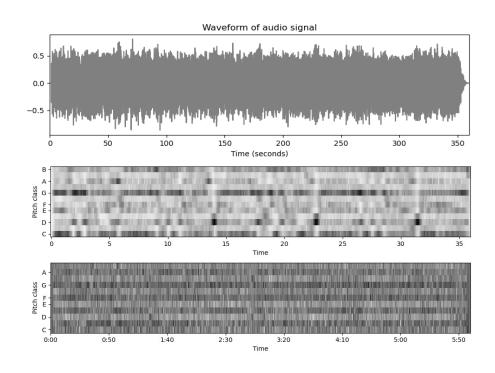
Novelty-Based approach

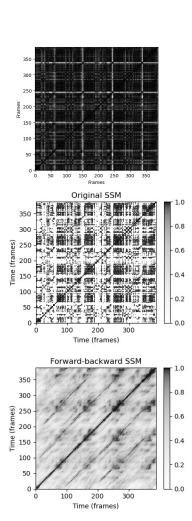
```
def compute_kernel_checkerboard_gaussian(L, var=1, normalize=True):
     taper = np.sqrt(1/2) / (L * var)
     axis = np.arange(-L, L+1)
     qaussian1D = np.exp(-taper**2 * (axis**2))
     gaussian2D = np.outer(gaussian1D, gaussian1D)
     kernel box = np.outer(np.sign(axis), np.sign(axis))
     kernel = kernel_box * gaussian2D
     if normalize:
           kernel = kernel / np.sum(np.abs(kernel))
     return kernel
def compute novelty ssm(S, kernel=None, L=10, var=0.5, exclude=False):
     if kernel is None:
           kernel = compute_kernel_checkerboard_gaussian(L=L, var=var)
     N = S.shape[0]
     M = 2*L + 1
     nov = np.zeros(N)
     S padded = np.pad(S, L, mode='constant')
     for n in range(N):
           nov[n] = np.sum(S padded[n:n+M, n:n+M] * kernel)
     if exclude:
           right = np.min([L, N])
           left = np.max([0, N-L])
           nov[0:right] = 0
           nov[left:N] = 0
     return nov
```

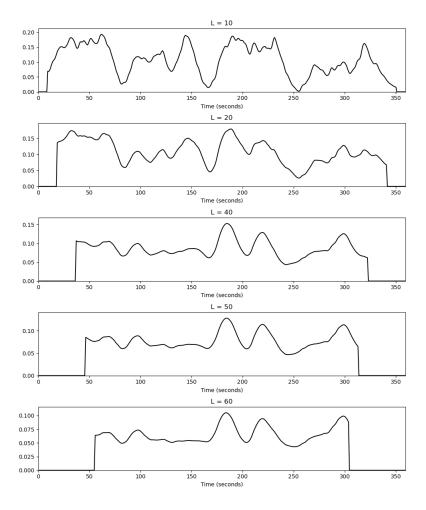


Summary

Like a Rolling Stone - Bob Dylan







Thank you for your attention.