Deep Learning for Music Analysis and Generation

Rhythm

Beat tracking, downbeat tracking & tempo estimation (audio \rightarrow score)



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FMP Notebook

https://www.audiolabs-erlangen.de/resources/MIR/FMP/C6/C6.html

Part	Title	Notions, Techniques & Algorithms	HTML	IPYNB
B impyler	<u>Basics</u>	Basic information on Python, Jupyter notebooks, Anaconda package management system, Python environments, visualizations, and other topics	[html]	[ipynb]
O (21) (11) (11)	Overview	Overview of the notebooks (https://www.audiolabs-erlangen.de/FMP)	[html]	[ipynb]
1	Music Representations	Music notation, MIDI, audio signal, waveform, pitch, loudness, timbre	[html]	[ipynb]
2	Fourier Analysis of Signals	Discrete/analog signal, sinusoid, exponential, Fourier transform, Fourier representation, DFT, FFT, STFT	[html]	[ipynb]
3	Music Synchronization	Chroma feature, dynamic programming, dynamic time warping (DTW), alignment, user interface	[html]	[ipynb]

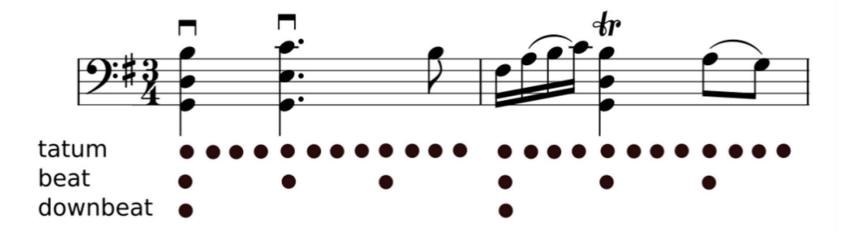
Part	Title	Notions, Techniques & Algorithms	HTML	IPYNB
4	Music Structure Analysis	Similarity matrix, repetition, thumbnail, homogeneity, novelty, evaluation, precision, recall, F- measure, visualization, scape plot	[html]	[ipynb]
5	Chord Recognition	Harmony, music theory, chords, scales, templates, hidden Markov model (HMM), evaluation	[html]	[ipynb]
6	Tempo and Beat Tracking	Onset, novelty, tempo, tempogram, beat, periodicity, Fourier analysis, autocorrelation	[html]	[ipynb]
7	Content-Based Audio Retrieval	Identification, fingerprint, indexing, inverted list, matching, version, cover song	[html]	[ipynb]
8	Musically Informed Audio Decomposition	Harmonic/percussive separation, signal reconstruction, instantaneous frequency, fundamental frequency (F0), trajectory, nonnegative matrix factorization (NMF)	[html]	[ipynb]

ISMIR 2021 Tutorial

https://tempobeatdownbeat.github.io/tutorial/intro.html

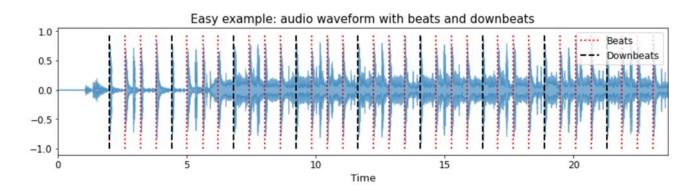
Tempo, Beat, and Downbeat Estimation

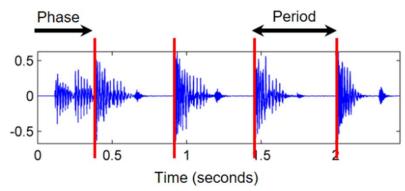
By Matthew E. P. Davies, Sebastian Böck, Magdalena Fuentes



What is Beat/Downbeat Tracking?

https://tempobeatdownbeat.github.io/tutorial/ch2_basics/definition.html





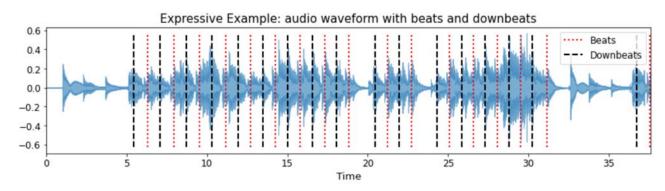


Figure 6.1 from [Müller, FMP, Springer 2015]

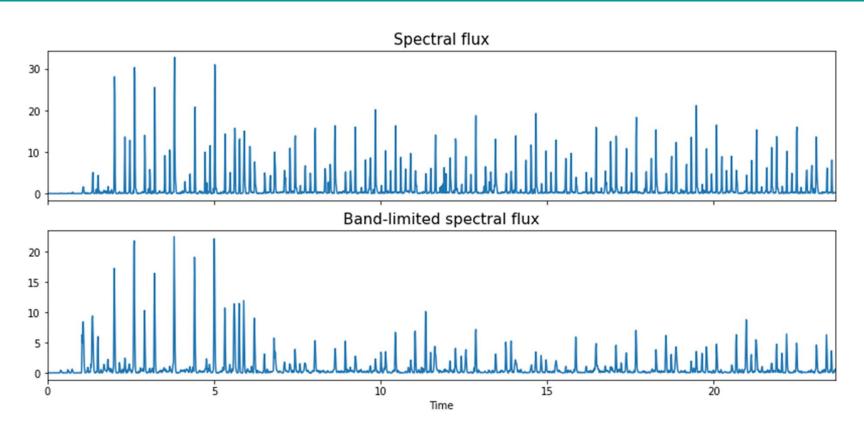
Beat/Downbeat Annotation

https://tempobeatdownbeat.github.io/tutorial/ch2_basics/annotation.html

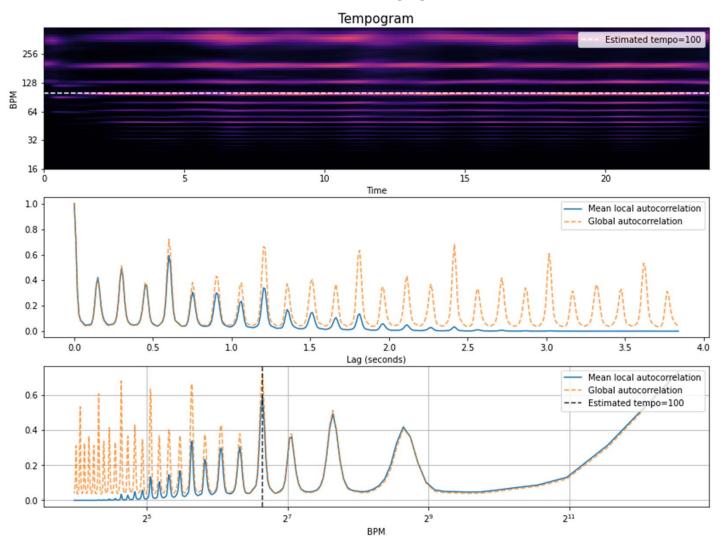
- Annotation is hard!
- It takes a long time, and the more challenging the material to annotate the greater the likelihood of this being helpful for learning.
- On the plus side, annotation is a fantastic way to learn about the task of beat and downbeat estimation so it's a really great excercise.
- We always need more data, so do consider doing some annotating!
- As hard as we try, annotation "mistakes" are made, so they made need correcting.
- This makes comparative evaluation more challenging, so it's always worthwile to ensure you are using the most up to date version of any annotations.

Baseline Approach

https://tempobeatdownbeat.github.io/tutorial/ch2_basics/baseline.html



Baseline Approach



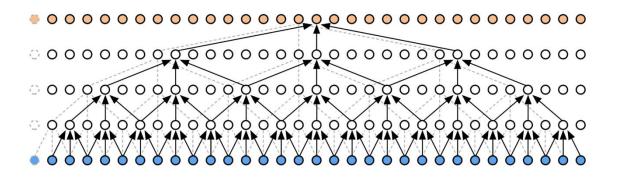
Deep Learning Approaches

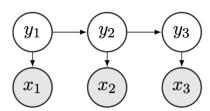
https://tempobeatdownbeat.github.io/tutorial/ch3_going_deep/overview.html

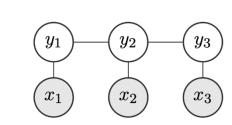


https://tempobeatdownbeat.github.io/tutorial/ch3_going_deep/dnns.html

https://tempobeatdownbeat.github.io/tut orial/ch3_going_deep/postprocessing.html







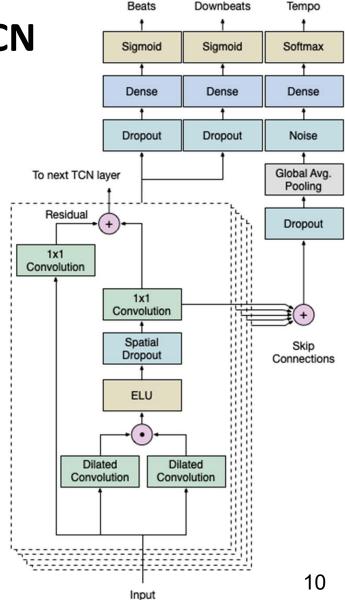
Hands on!

https://colab.research.google.com/drive/1tuOqNyO9gdMmYJsj33fP_QOfpRsm2tmt?usp=sharing

Ref: Böck & Davies, "Deconstruct, analyse, reconstruct: How to improve tempo, beat, and downbeat estimation," ISMIR 2020

Main Architecture - TCN

```
def residual block(x, i, activation, num filters, kernel size, padding, dropout rate=0, name=''):
       # name of the layer
       name = name + '_dilation_%d' % i
       # 1x1 conv. of input (so it can be added as residual)
       res_x = Conv1D(num_filters, 1, padding='same', name=name + '_1x1_conv_residual')(x)
       # two dilated convolutions, with dilation rates of i and 2i
       conv 1 = Conv1D(
              filters=num filters.
              kernel_size=kernel_size,
              dilation rate=i.
              padding=padding,
             name=name + '_dilated_conv_1',
       ) (x)
       conv 2 = Conv1D(
              filters=num filters.
              kernel_size=kernel_size,
              dilation rate=i * 2,
              padding=padding,
              name=name + '_dilated_conv_2',
       ) (x)
       # concatenate the output of the two dilations
       concat = keras.layers.concatenate([conv_1, conv_2], name=name + '_concat')
       # apply activation function
       x = Activation(activation, name=name + '_activation')(concat)
       # apply spatial dropout
       x = SpatialDropout1D(dropout_rate, name=name + '_spatial_dropout_%f' % dropout_rate)(x)
       # 1x1 conv. to obtain a representation with the same size as the residual
       x = Conv1D(num_filters, 1, padding='same', name=name + '_1x1_conv')(x)
```



PreNet

```
def create model (input shape, num filters=20, num dilations=11, kernel size=5, activation='elu', dropout rate=0.15):
       # input laver
       input_layer = Input(shape=input shape)
       # stack of 3 conv layers, each conv, activation, max. pooling & dropout
       conv_1 = Conv2D(num_filters, (3, 3), padding='valid', name='conv_1_conv')(input_layer)
       conv_1 = Activation(activation, name='conv_1_activation')(conv_1)
       conv 1 = MaxPooling2D((1, 3), name='conv 1 max pooling')(conv 1)
       conv 1 = Dropout(dropout rate, name='conv 1 dropout')(conv 1)
       conv_2 = Conv2D(num_filters, (1, 10), padding='valid', name='conv_2_conv')(conv_1)
       conv 2 = Activation(activation, name='conv 2 activation')(conv 2)
       conv 2 = MaxPooling2D((1, 3), name='conv 2 max pooling')(conv 2)
       conv_2 = Dropout(dropout_rate, name='conv_2_dropout')(conv_2)
       conv_3 = Conv2D(num_filters, (3, 3), padding='valid', name='conv_3 conv')(conv_2)
                                                                                              input
       conv 3 = Activation(activation, name='conv 3 activation')(conv 3)
       conv_3 = MaxPooling2D((1, 3), name='conv_3_max_pooling')(conv_3)
       conv 3 = Dropout(dropout rate, name='conv 3 dropout')(conv 3)
       # reshape layer to reduce dimensions
                                                                                                                                                        1x3
       x = Reshape((-1, num filters), name='tcn input reshape')(conv 3)
                                                                                                    3x3
       # TCN lavers
                                                                                                    conv
       dilations = [2 ** i for i in range(num_dilations)]
                                                                                                                                                            to TCN
       ten, skip = TCN(
              num_filters=[num_filters] * len(dilations).
              kernel size=kernel size.
                                                                                              5.81.1
                                                                                                         3.79.20
                                                                                                                  3.26.20
                                                                                                                              3.15.20
                                                                                                                                         3.5.20
                                                                                                                                                    1.3.20
                                                                                                                                                             1.1.20
              dilations=dilations.
```

Data Sequence Handling & Target Widening

```
# infer (global) tempo from beats
def infer tempo(beats, hist smooth=15, fps=FPS, no tempo=MASK VALUE):
       ibis = np. diff(beats) * fps
       bins = np. bincount(np. round(ibis). astype(int))
       # if no beats are present, there is no tempo
       if not bins. any():
             return NO TEMPO
       intervals = np. arange (1en (bins))
       # smooth histogram bins
       if hist_smooth > 0:
              bins = madmom. audio. signal. smooth (bins, hist smooth)
       # create interpolation function
       interpolation fn = interpld(intervals, bins, 'quadratic')
       # generate new intervals with 1000x the resolution
       intervals = np. arange (intervals[0], intervals[-1], 0.001)
       tempi = 60.0 * fps / intervals
       # apply quadratic interpolation
       bins = interpolation fn(intervals)
       peaks = argrelmax(bins, mode='wrap')[0]
       if len(peaks) == 0:
              # no peaks, no tempo
             return no tempo
       else:
              # report only the strongest tempo
              sorted peaks = peaks[np.argsort(bins[peaks])[::-1]]
              return tempi[sorted peaks][0]
```

```
# pad features
def cnn_pad(data, pad_frames):
    """Pad the data by repeating the first and last frame N times."""
    pad_start = np.repeat(data[:1], pad_frames, axis=0)
    pad_stop = np.repeat(data[-1:], pad_frames, axis=0)
    return np.concatenate((pad_start, data, pad_stop))
```

```
def widen beat targets(self, size=3, value=0.5):
       for y in self. beats. values():
              # skip masked beat targets
              if np.allclose(y, MASK_VALUE):
                     continue
              np. maximum(y, maximum filterld(y, size=size) * value, out=y)
def widen_downbeat_targets(self, size=3, value=0.5):
       for y in self. downbeats. values():
              # skip masked downbeat targets
              if np. allclose(y, MASK_VALUE):
                     continue
              np. maximum(y, maximum filterld(y, size=size) * value, out=y)
def widen tempo targets(self, size=3, value=0.5):
       for y in self. tempo. values():
              # skip masked tempo targets
              if np. allclose(y, MASK_VALUE):
              np.maximum(y, maximum filterld(y, size=size) * value, out=y)
```

Data Augmentation

"We use a simple approach of adding the same training examples with changed hop-size when computing the STFT. This results in the same song being represented with a different numbers of frames. This way the beat positions are "streched" or "squeezed" and the tempo changes accordingly."

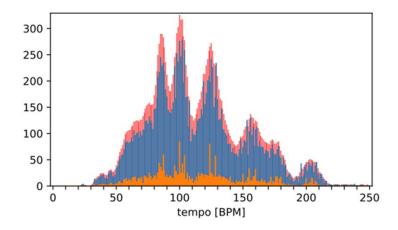


Figure 2: Tempo distribution of original tempo annotations (orange, foreground), after data augmentation (blue) and target widening (red, background).

Post-processing

```
def detect tempo (bins, hist smooth=11, min bpm=10):
       min bpm = int(np.floor(min bpm))
       tempi = np. arange (min_bpm, len(bins))
       bins = bins[min_bpm:]
       # smooth histogram bins
       if hist_smooth > 0:
              bins = madmom. audio. signal. smooth (bins, hist_smooth)
       # create interpolation function
       interpolation fn = interpld(tempi, bins, 'quadratic')
       # generate new intervals with 1000x the resolution
       tempi = np.arange(tempi[0], tempi[-1], 0.001)
       # apply quadratic interpolation
      bins = interpolation_fn(tempi)
       peaks = argrelmax(bins, mode='wrap')[0]
       if len(peaks) == 0:
              # no peaks, no tempo
              tempi = np.array([], ndmin=2)
       elif len(peaks) == 1:
              # report only the strongest tempo
              ret = np.array([tempi[peaks[0]], 1.0])
              tempi = np.array([tempi[peaks[0]], 1.0])
       else:
              # sort the peaks in descending order of bin heights
              sorted_peaks = peaks[np.argsort(bins[peaks])[::-1]]
              # normalize their strengths
              strengths = bins[sorted_peaks]
              strengths /= np. sum (strengths)
```

Library: Madmom

https://github.com/CPJKU/madmom