Intelligent Systems

Excersice 4- Representation

Simon Reichhuber
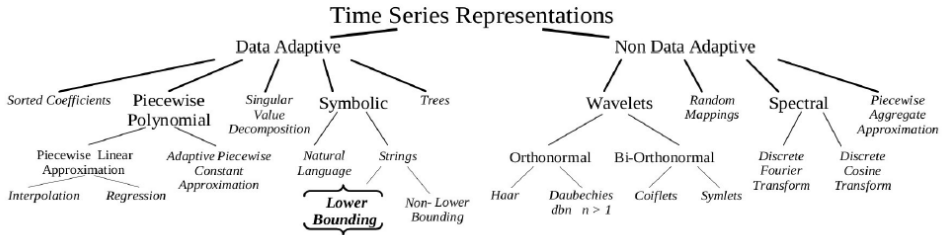
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Data representation

The FT represents the time series in the frequency domain. The signal is constructed as a sequence of sine and cosine terms.
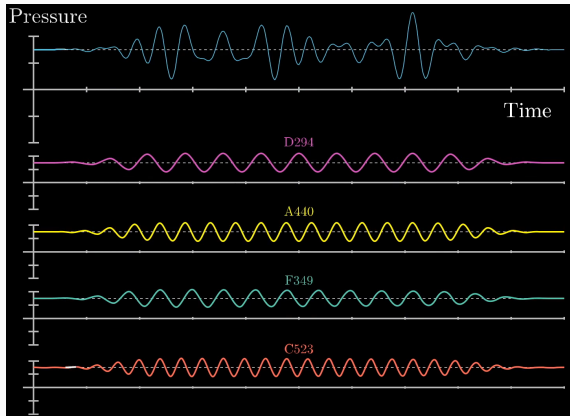

Figure 1: Fourier Transform

The FT represents the time series in the frequency domain. The signal is constructed as a sequence of sine and cosine terms.
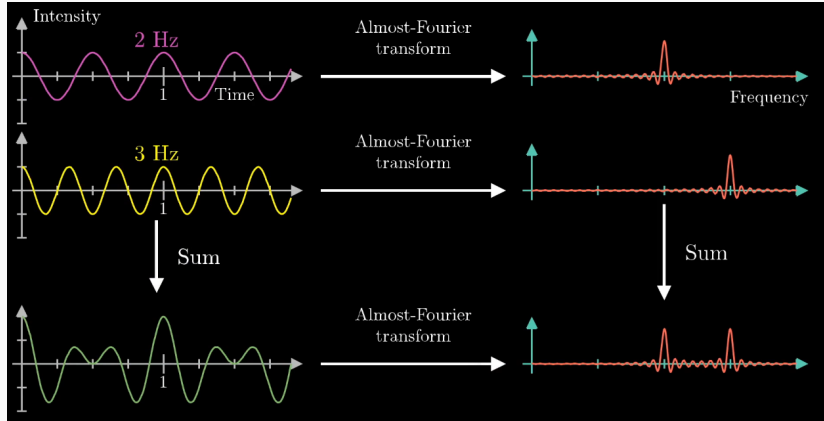

Figure 2: Fourier Transform

A. Explain the idea of the *Shape Definition Language* and its application?

B. Approximate the time series with the following approximations:
- *Piecewise Aggregate Approximation* (*PAA*) with 4 segemtns.
- *Clipping* to binary values ($\rightarrow$ search the procedure on the internet).
- *Picewise Linear Approximation* with 4 segments.
- *Run-Length Encoding* (*RLE*).

C. Aggregate the timeseries to the following statistical measures:
- *Mean*
- *Standard deviation*
- *Mode*

D. What are the advantages and disatvantages of the *clipping* procedure?

E. What is the main difference between the *Adaptive Picewise Aggregate Approximation* (*APAA*) and the *PAA*?

Given the case that the most important information of a time series can be extracted from the rough shape, limited terms of the *Shape Definition Language* (*SDL*) are enough to model it, i.e. *Up, up, stable, zero, down, Down*. These represent the different slopes of the time series. Advantageously, the representation can easily be processed by algorithms for symbolic sequences, like *Longest Common Subsequence* (*LCSS*).
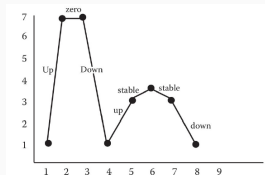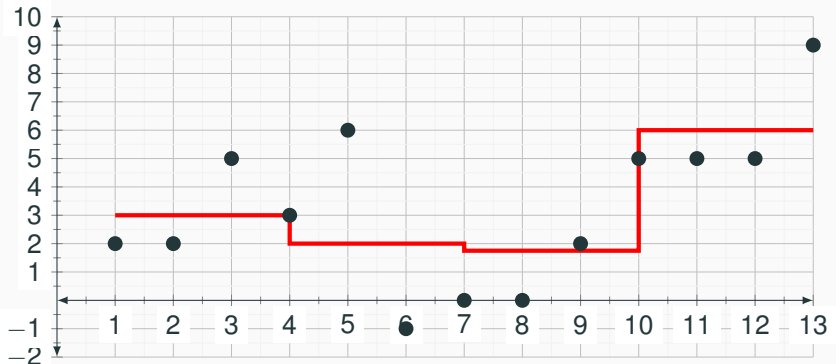

Figure 3: Shape Definition Language[1]

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[1]Mitsa, Theophano. Temporal data mining. CRC Press, 2010

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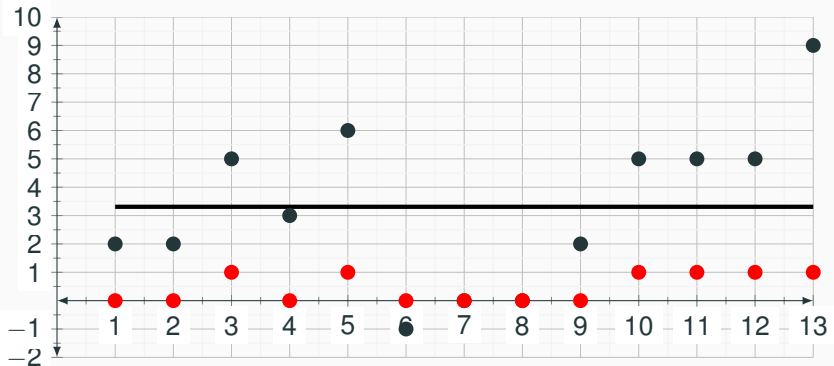
- PAA Segment Length $= \frac{t_{end} - t_{start}}{\#Segments}$
- 4 segments $\rightarrow$ PAA Segment Length $= \frac{13-1}{4} = 3$
- Segment 1: $\frac{2+2+5+3}{4} = 3$
- Segment 2: $\frac{3+6+(-1)+0}{4} = 2$
- ...

- Calculate time series' mean: $\mu = \frac{1}{N} \sum_{i=1}^{N} x_i$

- The clipped values are given by: $y_i^* = \begin{cases} 1 \text{ , for } x_i \geq \mu \\ 0 \text{ , otherwise} \end{cases}$

- $\mu = \frac{2+2+5+3+6+(-1)+0+0+2+5+5+5+9}{13} \approx 3.31$

- Clipping: $0, 0, 1, 0, 1, 0, 0, 0, 0, 1, 1, 1, 1$
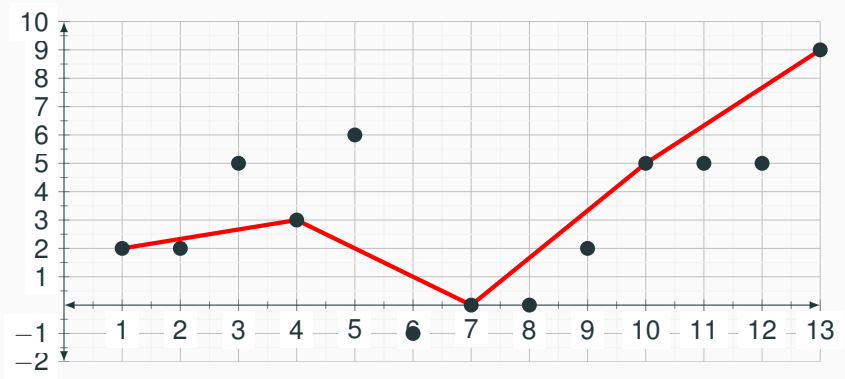
- Counter $\{n\}$ represents the number of repititions of the following symbol
- $2, 2, 5, 3, 6, -1, 0, 0, 2, 5, 5, 5, 9 \rightarrow$
  $\{2\}2, 5, 3, 6, -1, \{2\}0, 2, \{3\}5, 9$

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Given N D-dimensional points $x_i \in \mathbb{R}^d, i = 1, \ldots, N$, the following statistical measures can be defined:

- *Mean:* $\mu = \frac{1}{N} \sum_{i=1}^{N} x_i \quad \rightarrow 3.31$
- *Standard deviation*: $\sigma = \sqrt{\frac{1}{N-1} \sum_{i=1}^{N} (x_i - \mu)^2} \rightarrow 2.84$
- *Mode*: $m = argmax_x |\{y | y \in \{x_1, \ldots, x_N\}, x = y\}| \rightarrow 5$

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What are the advantages and disatvantages of the *clipping* procedure?

Advantages:

- Extreme compression rate (*float* $\rightarrow$ *bool*)
- Simple representation (only binary values)
- Rough patterns can be found easily

Disatvantage:

- Very unprecise representation of the signal

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What is the main difference between the *Adaptive Picewise Aggregate Approximation* (*APAA*) and the *PAA*?
**Main difference:**

- Variable length of the (temporal) sections
- Adaptive according to local details of a time series
  - Sections of frequent movement will be parted in smaller intervals
  - A Section without significant information will be represented as a large interval

What is the main difference between the *Adaptive Picewise Aggregate Approximation* (*APAA*) and the *PAA*?

Advantage:

- The error (e.g. Least-Squares) between the raw data and the representation is reduced.[2]
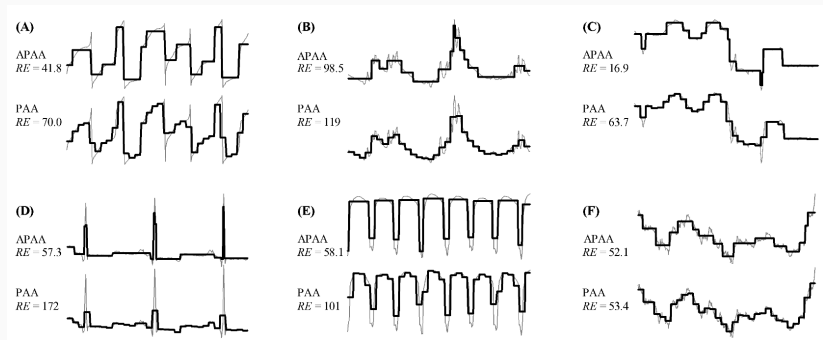
[2]Koegh et. al. 2001

Principal component analysis

A. What is the goal of the *Principal Component Analysis* (*PCA*) and what is its basic assumption.

B. What is the benefit of the *PCA*?

C. Describe the following items:

- Zero-mean feature
- Variance
- Standard deviation
- Covariance matrix
- Arithmetic mean
- Eigenvector
- Eigenvalue
- Projection onto new feature space

D. How can we get a dimensionality reduction with the means of Eigenvalues?

What is the goal of the *Principal Component Analysis* (*PCA*) and what is its basic assumption. **Basic assumption:**

- The larger the variance, the higher is the level of information.

Goal:

- Dimensionality reduction:
    - Remove dimensions with poor information gain
    - Generate new dimensions that better fit to the structure of the data, i.e. at which the data has the largest variance.

Benefits:

- **Reduce computing time**: Through the usage of data mining algorithms applied on dimensionality-reduced data sets.
- **Feature selection**: Very easy; just select the PCA meta features.
- **Comprehensability**: Easy detection of structures, e.g. by projecting the data onto the two or three most important meta features.

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D. How can we get a dimensionality reduction with the means of Eigenvalues?

Given: Data set with *N* samples and *D* dimensions (features).

1. Standardisation:

- Calculate the arithmetic mean for each feature j:
  $\mu_j = \frac{1}{N} \sum_{i=1}^{N} x_{j,i}$

- Zero-mean sample:
  $x'_{j,i} = x_{j,i} - \mu_j$

- Calculate the standard deviation for each feature j:
  $\sigma_j = \sigma_j = \sqrt{\frac{1}{N-1} \sum_{i=1}^{N} x'^2_{j,i}}$

- Standardisation for every feature j:
  $x_{j,i} = \frac{x'_{j,i}}{\sigma_j}$

2. Calculate covariance matrix:

- $s_{j,j'} = \frac{1}{N-1} \sum_{i=1}^{N} x'_{j,i} \cdot x'_{j',i}$

- $S = \begin{pmatrix} s_{1,1} & \cdots & s_{1,D} \\ \vdots & \ddots & \vdots \\ s_{D,1} & \cdots & s_{D,D} \end{pmatrix}$

3. Calculate eigenvectors and eigenvalues:

- Eigenvalues represent the ratio of the variance along their corresponding eigenvectors
- The eigenvector with the largest eigenvalue represents the direction, in which the data has the largest variance
- The eigenvector with the second largest eigenvalue represents a orthogonal direction w.r.t. the first eigenvector, in which the data has the second largest variance, etc.
- Eigenvectors are the *principal components* (*PC*), which are more suitable to model the structure of the data than the original features (assuming that high information gain corresonds to high variance).

4. Dimensionality reduction: Choose the most important eigenvalues according to their eigenvalues:

- Method 1: The sum of the remaining eigenvalues shall be larger than a predefined ratio (e.g. 0.75) of the sum of all eigenvalues.
- Method 2: Dimensions shall be removed if the eigenvalue of the corresponding eigenvector is lower than the mean of all eigenvalues.

4. Projection onto the new feature space:

- Spanned by the new selected eigenvectors (*PC*s).

Python PCA

A. Download the file *04_Representation.ipyn* from *OpenOlat*.

B. In order to solve the tasks, you can use the library *numpy*.

C. Compare your results afterwards with the help of *sklearn*.