

Indexing Time Series

Based on original slides by Prof. Dimitrios Gunopulos and Prof. Christos Faloutsos with some slides from tutorials by Prof. Eamonn Keogh and Dr. Michalis Vlachos. Excellent tutorials (and not only) about Time Series can be found there:

http://www.cs.ucr.edu/~eamonn/tutorials.html

A nice tutorial on Matlab and Time series is also there:

http://www.cs.ucr.edu/~mvlachos/ICDM06/

Time Series Databases

- A time series is a sequence of real numbers, representing the measurements of a real variable at equal time intervals
 - Stock prices
 - Volume of sales over time
 - Daily temperature readings
 - ECG data
- A time series database is a large collection of time series

Time Series Data

25.1750 25.1750 25.2250 25.2500

25.2500 25.2750 25.3250

25.3500 25.3500 25.4000

25.4000 25.3250 25.2250

25.2000 25.1750

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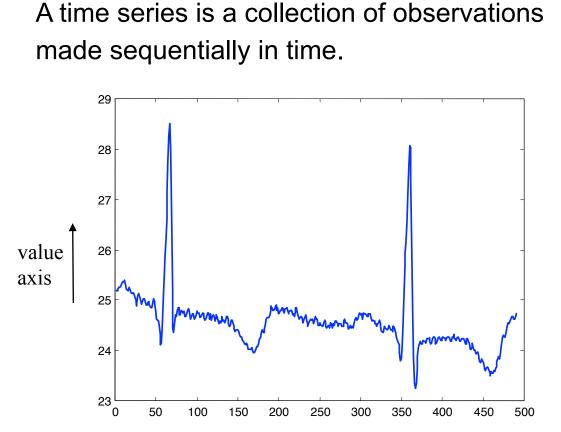
24.6250

24.6750 24.6750 24.6250

24.6250

24.6250 24.6750

24.7500



time axis

Time Series Problems



(from a database perspective)

The Similarity Problem

$$X = x_1, x_2, ..., x_n \text{ and } Y = y_1, y_2, ..., y_n$$

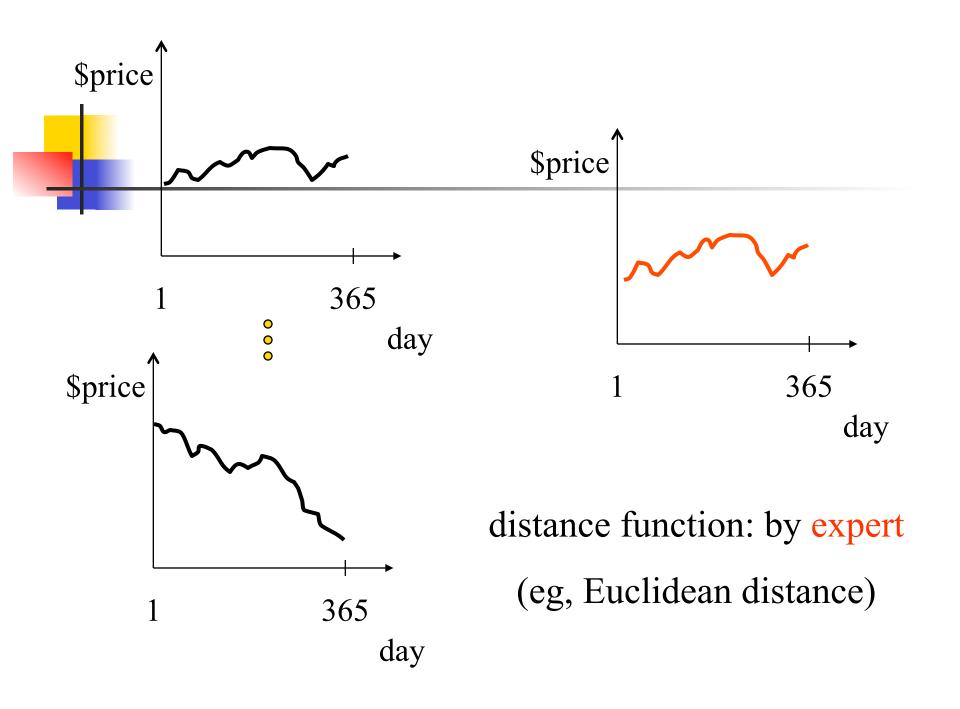
- Define and compute Sim(X, Y)
 - E.g. do stocks X and Y have similar movements?
- Retrieve efficiently similar time series (Indexing for Similarity Queries)

Types of queries

- whole match vs sub-pattern match
- range query vs nearest neighbors
- all-pairs query



- Find companies with similar stock prices over a time interval
- Find products with similar sell cycles
- Cluster users with similar credit card utilization
- Find similar subsequences in DNA sequences
- Find scenes in video streams



Problems

- - Define the similarity (or distance) function
 - Find an efficient algorithm to retrieve similar time series from a database
 - (Faster than sequential scan)

The Similarity function depends on the Application

Metric Distances

- What properties should a similarity distance have to allow (easy) indexing?
- D(A,B) = D(B,A) Symmetry
- D(A,A) = 0 Constancy of Self-Similarity
- D(A,B) >= 0 Positivity
- $D(A,B) \le D(A,C) + D(B,C)$ Triangular Inequality
- Some times the distance function that best fits an application is not a metric... then indexing becomes interesting....

Euclidean Similarity Measure

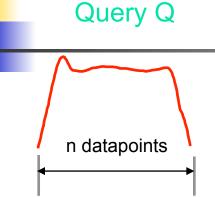
- View each sequence as a point in n-dimensional Euclidean space (n = length of each sequence)
- Define (dis-)similarity between sequences X and Y as

$$L_p = \left(\sum_{i=1}^n |x_i - y_i|^p\right)^{1/p}$$

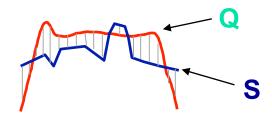
p=1 Manhattan distance

p=2 Euclidean distance

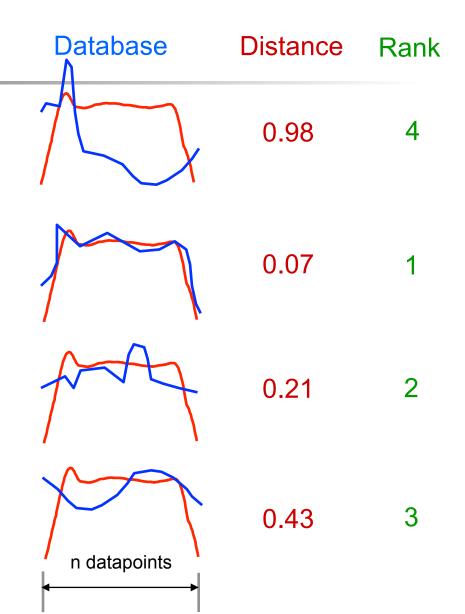
Euclidean model



Euclidean Distance between two time series Q = $\{q_1, q_2, ..., q_n\}$ and S = $\{s_1, s_2, ..., s_n\}$



$$D(Q,S) = \sqrt{\sum_{i=1}^{n} (q_i - s_i)^2}$$



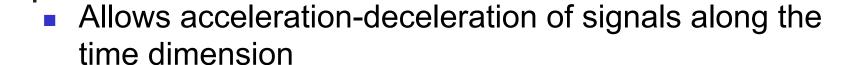


Advantages

- Easy to compute: O(n)
- Allows scalable solutions to other problems, such as
 - indexing
 - clustering
 - etc...

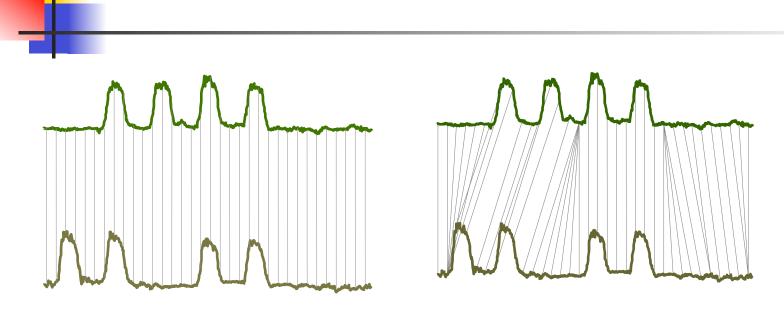
Dynamic Time Warping

[Berndt, Clifford, 1994]



- Basic idea
 - Consider $X = x_1, x_2, ..., x_n$, and $Y = y_1, y_2, ..., y_n$
 - We are allowed to extend each sequence by repeating elements
 - Euclidean distance now calculated between the extended sequences X' and Y'
 - Matrix M, where m_{ij} = d(x_i, y_j)

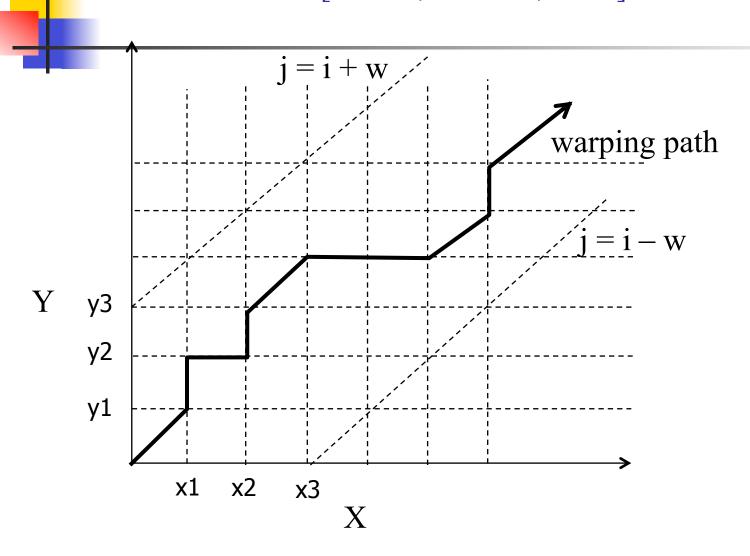
Example



Euclidean distance vs DTW

Dynamic Time Warping

[Berndt, Clifford, 1994]



Restrictions on Warping Paths

- Monotonicity
 - Path should not go down or to the left
- Continuity
 - No elements may be skipped in a sequence
- Warping Window

$$|i-j| \le w$$

Example

	s1	s2	s3	s4	s5	s6	s7	s8	s9
q1	3.76	8.07	1.64	1.08	2.86	0.00	0.06	1.88	1.25
q2	2.02	5.38	0.58	2.43	4.88	0.31	0.59	3.57	2.69
q3	6.35	11.70	3.46	0.21	1.23	0.29	0.11	0.62	0.29
q4	16.8	25.10	11.90	1.28	0.23	4.54	3.69	0.64	1.10
q5	3.20	7.24	1.28	1.42	3.39	0.04	0.16	2.31	1.61
q6	3.39	7.51	1.39	1.30	3.20	0.02	0.12	2.16	1.49
q7	4.75	9.49	2.31	0.64	2.10	0.04	0.00	1.28	0.77
8p	0.96	3.53	0.10	4.00	7.02	1.00	1.46	5.43	4.33
q9	0.02	1.08	0.27	8.07	12.18	3.39	4.20	10.05	8.53

Matrix of the pair-wise distances for element si with qj

Example

<i>s</i> 1	s2	s3	<i>s</i> 4	<i>s</i> 5	<i>s</i> 6	<i>s</i> 7	<i>s</i> 8	s9
<i>q</i> 1 3.76	11.83	13.47	14.55	17.41	17.41	17.47	19.35	20.60
q25.78	9.14	9.72	12.15	17.03	17.34	17.93	21.04	22.04
<i>q</i> 3 12.13	17.48	12.60	9.93	11.16	11.45	11.56	12.18	12.47
q4 29.02	37.23	24.50	11.21	10.16	14.70	15.14	12.20	13.28
q5 32.22	36.26	25.78	12.63	13.55	10.20	10.36	12.67	13.81
<i>q</i> 6 35.61	39.73	27.17	13.93	15.83	10.22	10.32	12.48	13.97
q7 40.36	45.10	29.48	14.57	16.03	10.26	10.22	11.50	12.27
q8 41.32	43.89	29.58	18.57	21.59	11.26	11.68	15.65	15.83
<i>q</i> 9 41.34	42.40	29.85	26.64	30.75	14.65	15.46	21.73	24.18

Matrix computed with Dynamic Programming based on the: $dist(i,j) = dist(si, yj) + min \{dist(i-1,j-1), dist(i, j-1), dist(i-1,j)\}$

Formulation

 Let D(i, j) refer to the dynamic time warping distance between the subsequences

$$x_1, x_2, ..., x_i$$

 $y_1, y_2, ..., y_j$

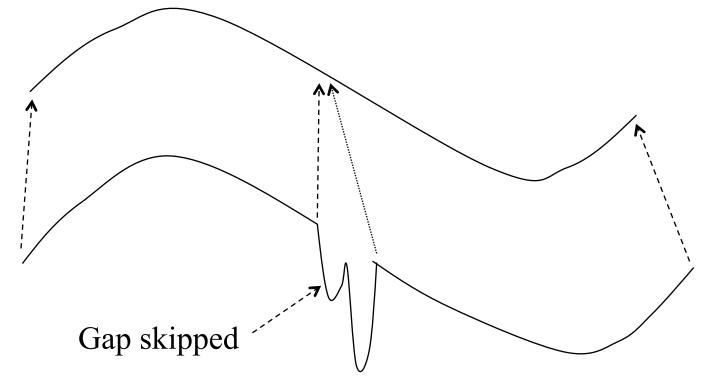
$$D(i, j) = |x_i - y_i| + min\{ D(i - 1, j), D(i - 1, j - 1), D(i, j - 1) \}$$

Solution by Dynamic Programming

- Basic implementation = O(n²) where n is the length of the sequences
 - will have to solve the problem for each (i, j) pair
- If warping window is specified, then O(nw)
 - Only solve for the (i, j) pairs where | i j | <= w</p>

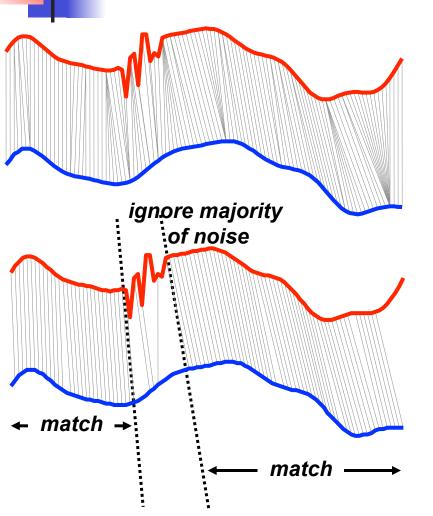
Longest Common Subsequence Measures

(Allowing for Gaps in Sequences)



Longest Common Subsequence (LCSS)

LC\$S is more resilient to noise than DTW.



Disadvantages of DTW:

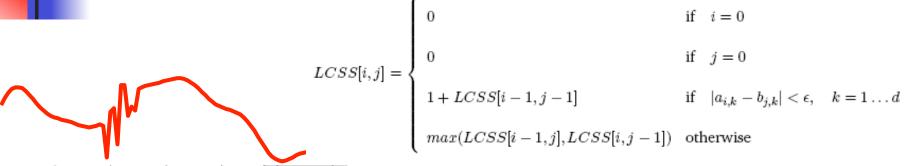
- A. All points are matched
- B. Outliers can distort distance
- C. One-to-many mapping

Advantages of LCSS:

- A. Outlying values not matched
- B. Distance/Similarity distorted less
- C. Constraints in time & space

Longest Common Subsequence

Similar dynamic programming solution as DTW, but now we measure similarity not distance.



Can also be expressed as distance

$$D_{LCSS}(A, B) = 1 - \frac{LCSS_{\delta,\epsilon}(A, B)}{\min(n, m) \text{ or } \max(n, m)}$$

Similarity Retrieval



Find all time series S where

$$D(Q,S) \leq \varepsilon$$

- Nearest Neighbor query
 - Find all the k most similar time series to Q
- A method to answer the above queries: Linear scan ... very slow

A better approach GEMINI

GEMINI



- extract m features (numbers, eg., avg., etc.)
- map into a point in m-d feature space
- organize points with off-the-shelf spatial access method ('SAM')
- retrieve the answer using a NN query
- discard false alarms

GEMINI Range Queries

Build an index for the database in a feature space using an R-tree

Algorithm RangeQuery(Q, ε)

- 1. Project the query Q into a point q in the feature space
- \mathbf{z} Find all candidate objects in the index within $\mathbf{\epsilon}$
- 3. Retrieve from disk the actual sequences
- 4. Compute the actual distances and discard false alarms

GEMINI NN Query



Algorithm K_NNQuery(Q, K)

- Project the query Q in the same feature space
- 2. Find the candidate K nearest neighbors in the index
- Retrieve from disk the actual sequences pointed to by the candidates
- 4. Compute the actual distances and record the maximum
- 5. Issue a RangeQuery(Q, εmax)
- 6. Compute the actual distances, return best K

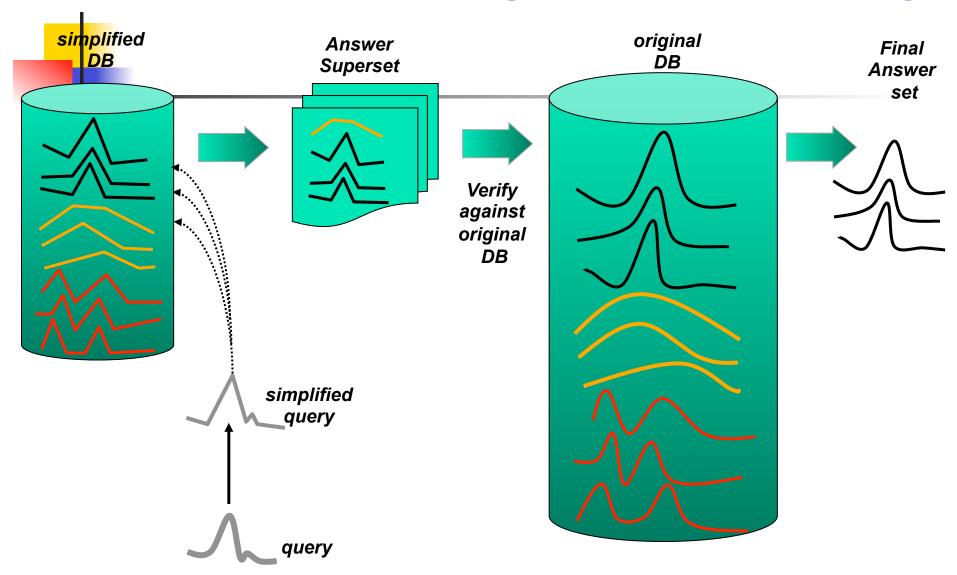
GEMINI



$$D_{feature}(F(x), F(y)) \leq D(x, y)$$

• Note that, the closer the feature distance to the actual one, the better.

Generic Search using Lower Bounding



Problem

How to extract the features? How to define the feature space?

- Fourier transform
- Wavelets transform
- Averages of segments (Histograms or APCA)
- Chebyshev polynomials
- your favorite curve approximation...

Fourier transform

- DFT (Discrete Fourier Transform)
- Transform the data from the time domain to the frequency domain
- highlights the periodicities
- SO?

DFT

A: several real sequences are periodic

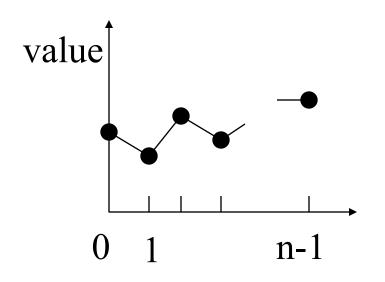
Q: Such as?

A:

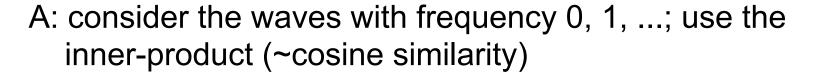
- sales patterns follow seasons;
- economy follows 50-year cycle (or 10?)
- temperature follows daily and yearly cycles

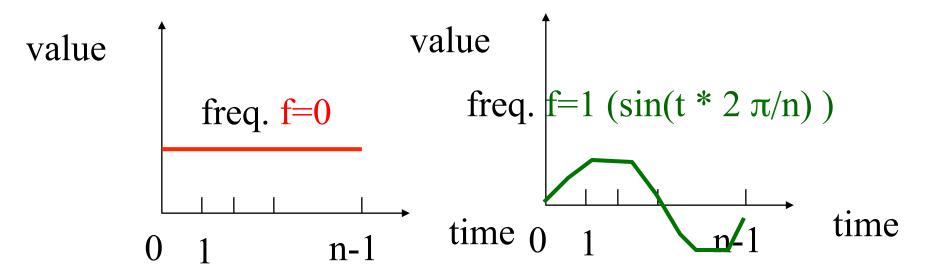
Many real signals follow (multiple) cycles

Decomposes signal to a sum of sine and cosine waves. Q:How to assess 'similarity' of **x** with a (discrete) wave?

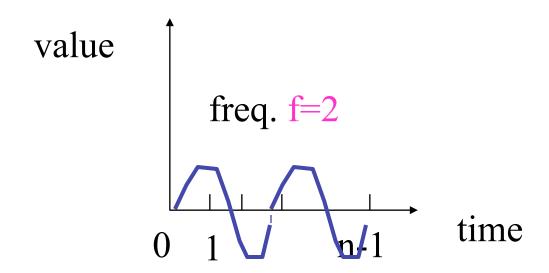


$$\mathbf{x} = \{x_0, x_1, \dots x_{n-1}\}$$
 $\mathbf{s} = \{s_0, s_1, \dots s_{n-1}\}$
time





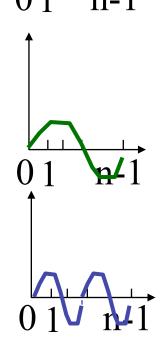
A: consider the waves with frequency 0, 1, ...; use the inner-product (~cosine similarity)

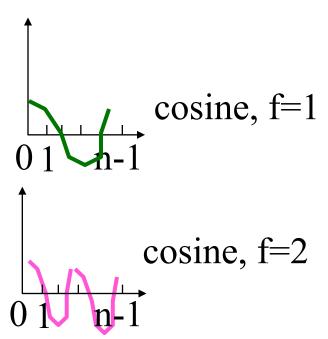




sine, freq =1

sine, freq = 2





How does it work?

- Basis functions are actually n-dim vectors,
 orthogonal to each other
- 'similarity' of x with each of them: inner product
- DFT: ~ all the similarities of x with the basis functions

How does it work?

Since $e^{jf} = cos(f) + j sin(f) (j=sqrt(-1))$, we finally have:

DFT: definition

Discrete Fourier Transform (n-point):

$$X_{f} = 1/\sqrt{n} \sum_{t=0}^{n-1} x_{t} * \exp(-j2\pi tf/n)$$

$$(j = \sqrt{-1}) \qquad \text{inverse DFT}$$

$$x_{t} = 1/\sqrt{n} \sum_{t=0}^{n-1} X_{f} * \exp(+j2\pi tf/n)$$

DFT: properties

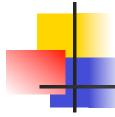
Observation - SYMMETRY property:

$$X_f = (X_{n-f})^*$$

("*": complex conjugate: $(a + b j)^* = a - b j$)

Thus we use only the first half numbers

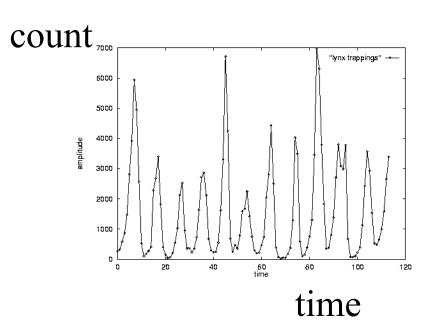
DFT: Amplitude spectrum

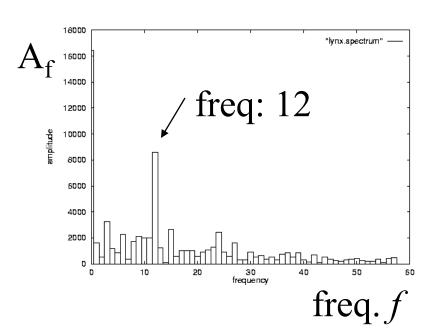


•Amplitude

$$A_f^2 = \operatorname{Re}^2(X_f) + \operatorname{Im}^2(X_f)$$

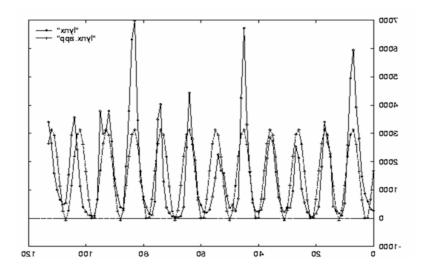
•Intuition: strength of frequency 'f'

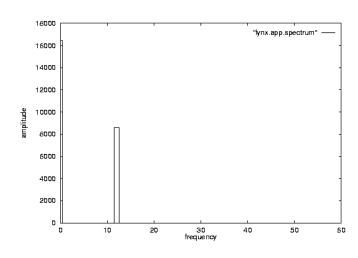


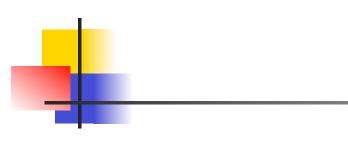


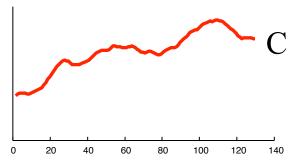
DFT: Amplitude spectrum

- excellent approximation, with only 2 frequencies!
- so what?









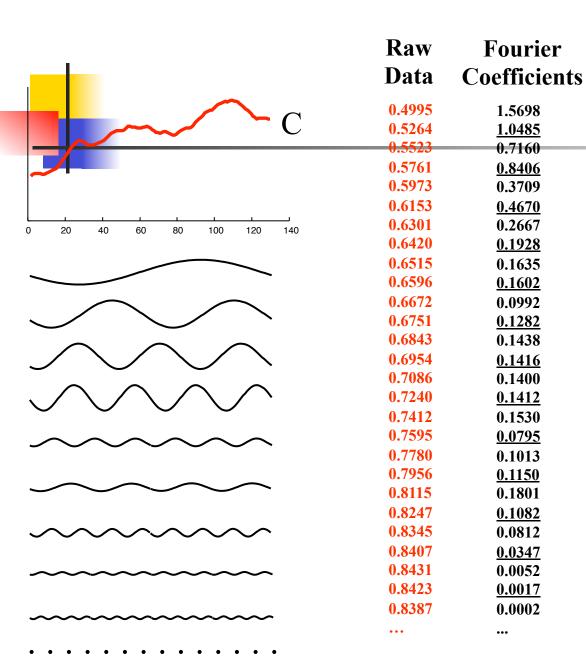
$$n = 128$$

Raw Data

0.4995 0.5264 0.5761 0.5973 0.6153 0.6301 0.6420 0.6515 0.6596 0.6672 0.6751 0.6843 0.6954 0.7086 0.7240 0.7412 0.7595 0.7780 0.7956 0.8115 0.8247 0.8345 0.8407 0.8431 0.8423 0.8387

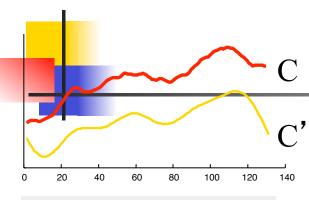
The graphic shows a time series with 128 points.

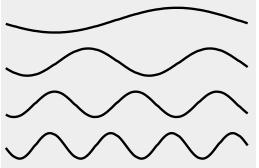
The raw data used to produce the graphic is also reproduced as a column of numbers (just the first 30 or so points are shown).



We can decompose the data into 64 pure sine waves using the Discrete Fourier Transform (just the first few sine waves are shown).

The Fourier Coefficients are reproduced as a column of numbers (just the first 30 or so coefficients are shown).





We have discarded $\frac{15}{16}$ of the data.

		Truncated
Raw	Fourier	Fourier
Data	Coefficients	Coefficients
0.4995	1 5698	1 5698

0.4995	1.5698
0.5264	1.0485
0.5523	0.7160
0.5761	0.8406
0.5973	0.3709
0.6153	<u>0.4670</u>
0.6301	0.2667
0.6420	<u>0.1928</u>
0.6515	0.1635
0.6596	<u>0.1602</u>
0.6672	0.0992
0.6751	0.1282
0.6843	0.1438
0.6954	<u>0.1416</u>
0.7086	0.1400
0.7240	<u>0.1412</u>
0.7412	0.1530
0.7595	<u>0.0795</u>
0.7780	0.1013
0.7956	<u>0.1150</u>
0.8115	0.1801
0.8247	<u>0.1082</u>
0.8345	0.0812
0.8407	<u>0.0347</u>
0.8431	0.0052
0.8423	<u>0.0017</u>
0.8387	0.0002

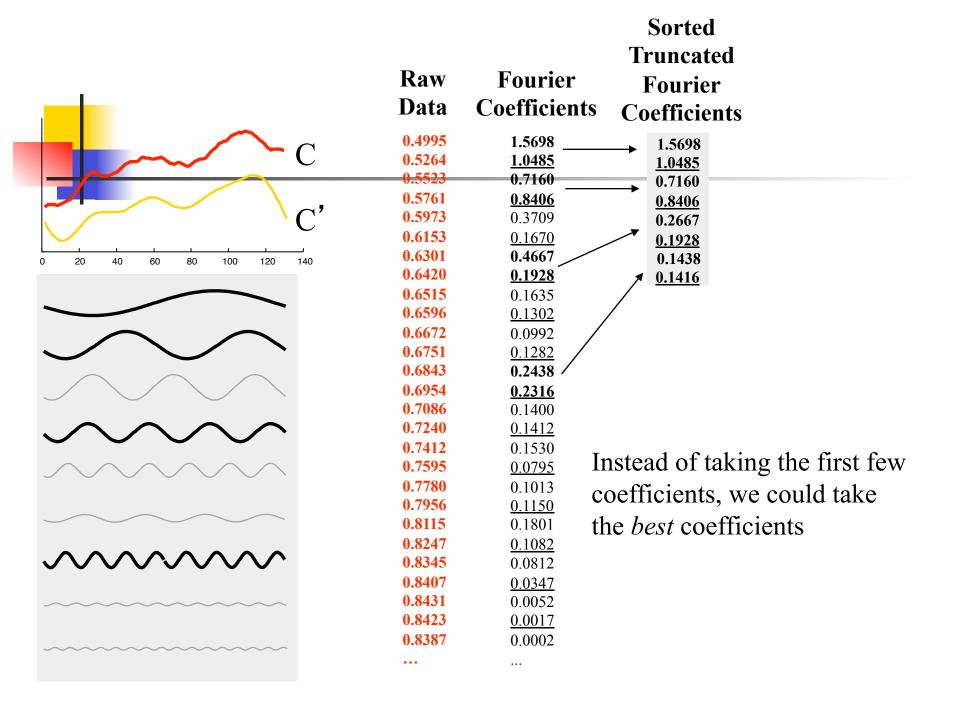
$$\begin{array}{c}
1.5698 \\
\underline{1.0485} \\
0.7160 \\
\underline{0.8406} \\
0.3709 \\
\underline{0.4670} \\
0.2667
\end{array}$$

$$n = 128$$

$$N = 8$$

$$C_{\text{ratio}} = 1/16$$

0.1928



DFT: Parseval's theorem

$$sum(x_t^2) = sum(|X_f|^2)$$

Ie., DFT preserves the 'energy' or, alternatively: it does an axis rotation:

$$\mathbf{x}$$
1
$$\mathbf{x} = \{\mathbf{x}0, \mathbf{x}1\}$$

$$\mathbf{x}$$

Lower Bounding lemma

Using Parseval's theorem we can prove the lower bounding property!

- So, apply DFT to each time series, keep first 3-10 coefficients as a vector and use an Rtree to index the vectors
- R-tree works with euclidean distance, OK.

Time series collections

Fourier and wavelets are the most prevalent and successful "descriptions" of time series.

- Next, we will consider collections of M time series, each of length N.
 - What is the series that is "most similar" to all series in the collection?
 - What is the second "most similar", and so on...

Time series collections

Some notation:

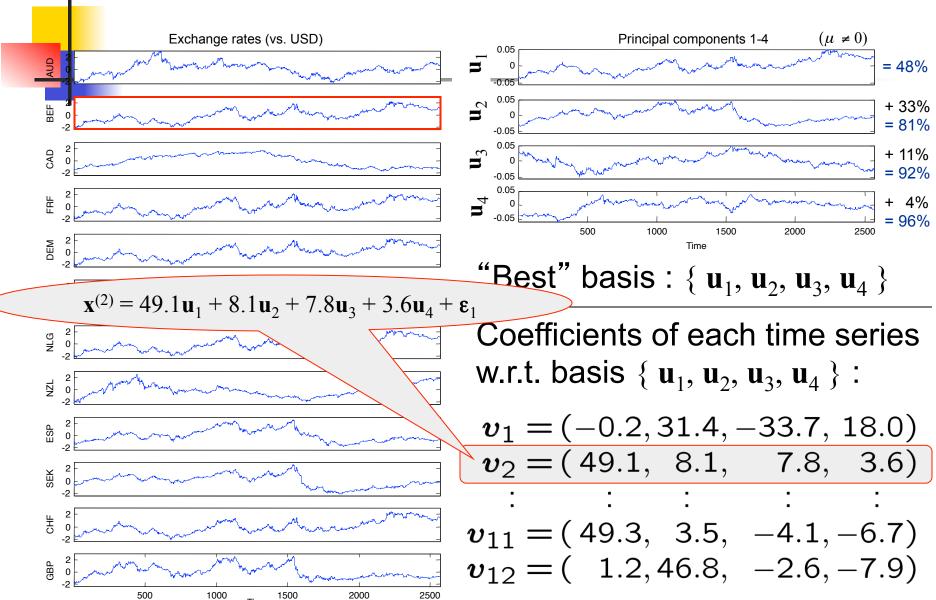
$$\mathbb{R}^N \ni \left(x_1^{(i)}, \dots x_N^{(i)}\right) \equiv \mathbf{x}^{(i)}$$
 : *i*-th time series vector

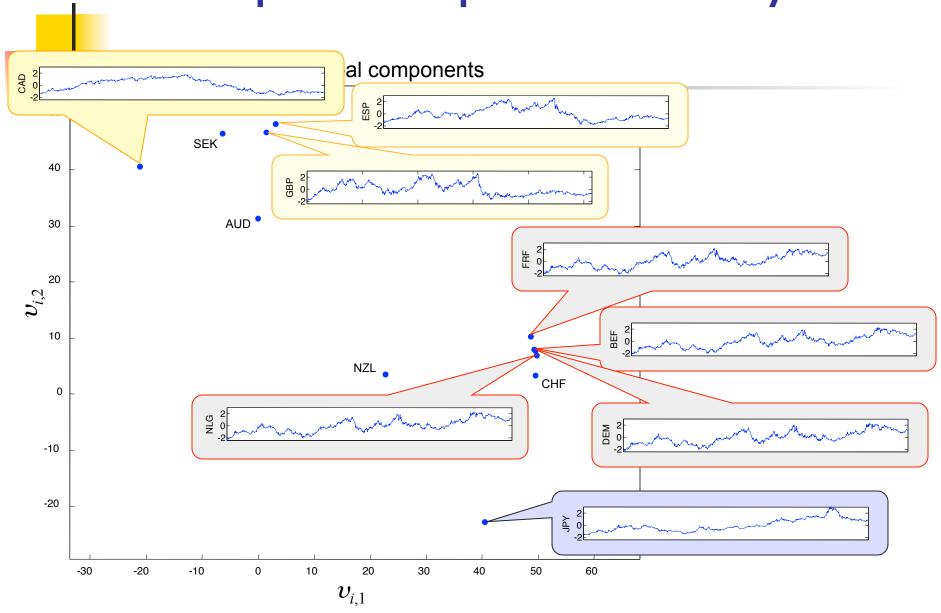
$$\mathbb{R}^M
ightarrow \left(x_t^{(1)}, \dots x_t^{(M)}\right) \equiv \mathbf{x}_t$$
 : vector of all series at time t

 $x_{t}^{(i)}$: value of the *i*-th series at time t

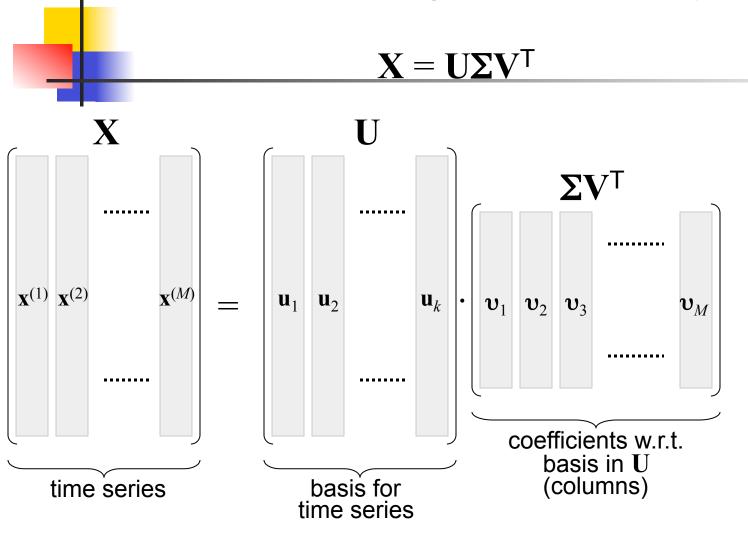
$$\mathbb{R}^{N \times M} \ni \mathbf{X} := \begin{pmatrix} x_1^{(1)} & x_1^{(2)} & \cdots & x_1^{(i)} & \cdots & x_1^{(M)} \\ x_1^{(1)} & x_1^{(2)} & \cdots & x_2^{(i)} & \cdots & x_2^{(M)} \\ \vdots & \vdots & \cdots & \vdots & \cdots & \vdots \\ x_t^{(1)} & x_t^{(2)} & \cdots & x_t^{(i)} & \cdots & x_t^{(M)} \\ \vdots & \vdots & \cdots & \vdots & \cdots & \vdots \\ x_N^{(1)} & x_N^{(2)} & \cdots & x_N^{(i)} & \cdots & x_N^{(M)} \end{pmatrix}$$
 values at time t , \mathbf{X}_t

Example

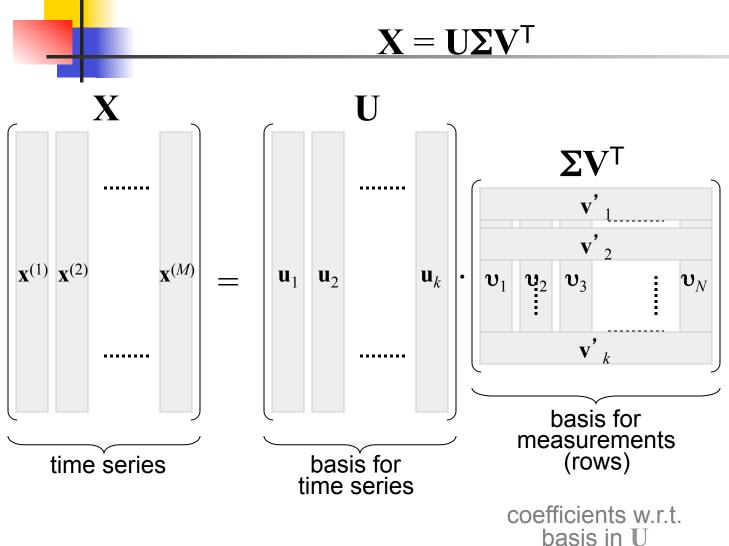




Matrix notation — Singular Value Decomposition (SVD)

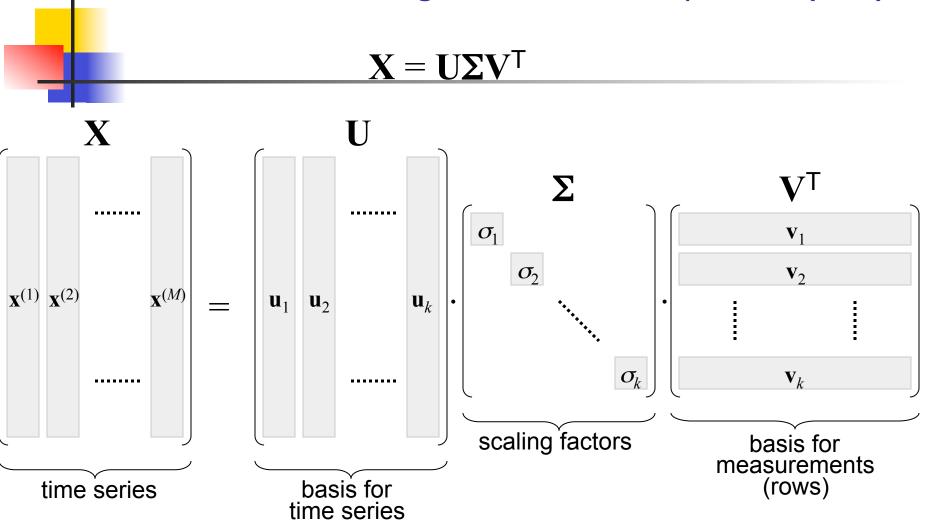


Matrix notation — Singular Value Decomposition (SVD)



basis in U (columns)

Matrix notation — Singular Value Decomposition (SVD)

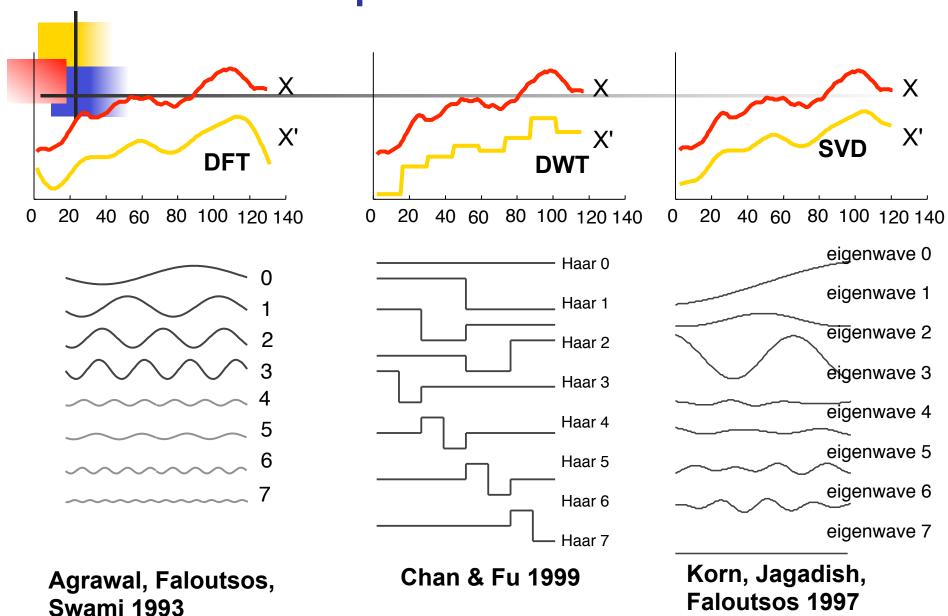




- PCA gives another lower dimensional transformation
- Easy to show that the lower bounding lemma holds

- but needs a collection of time series
- and expensive to compute it exactly

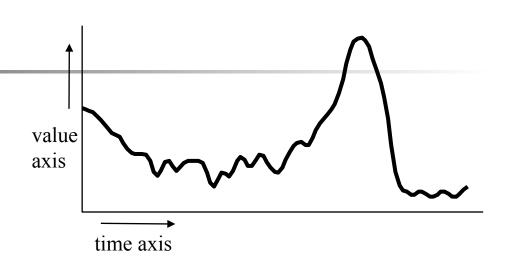
Feature Spaces



Piecewise Aggregate Approximation (PAA)

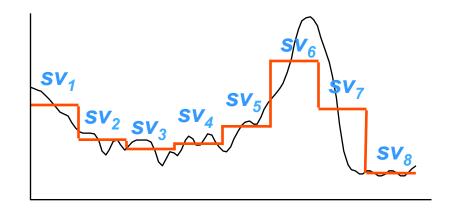


(n-dimensional vector) $S=\{s_1, s_2, ..., s_n\}$



n'-segment PAA representation (n'-d vector)

$$S = \{sv_1, sv_2, ..., sv_{n'}\}$$



PAA representation satisfies the lower bounding lemma (Keogh, Chakrabarti, Mehrotra and Pazzani, 2000; Yi and Faloutsos 2000)

Can we improve upon PAA?

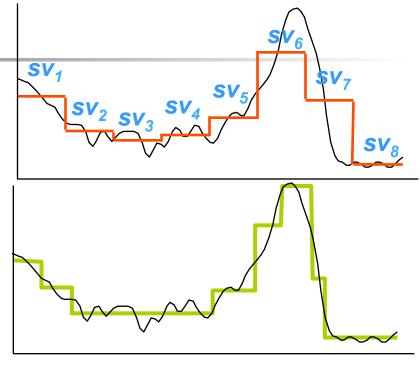
n'-segment PAA representation(n'-d vector)

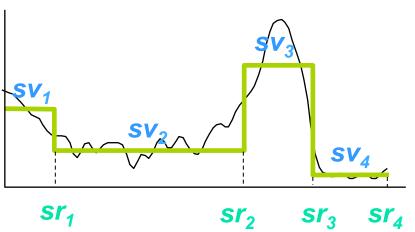
$$S = \{sv_1, sv_2, ..., sv_N\}$$

Adaptive Piecewise Constant Approximation (APCA)

n'/2-segment APCA representation (n'-d vector)

 $S = \{ sv_1, sr_1, sv_2, sr_2, ..., sv_M, sr_M \}$ (M is the number of segments = n'/2)

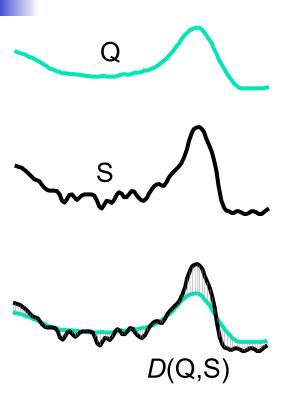




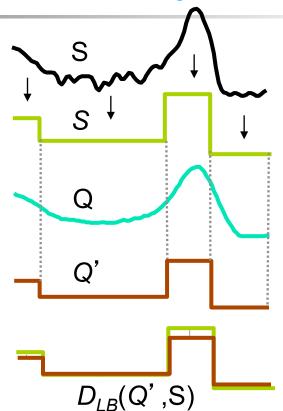
Distance Measure

ract (Euclidean) distance D(Q,S)

Lower bounding distance $D_{LB}(Q,S)$



$$D(Q,S) \equiv \sqrt{\sum_{i=1}^{n} (q_i - S_i)^2}$$



$$D_{LB}(Q',S)$$

$$\equiv \sqrt{\sum_{i=1}^{M} (sr_i - sr_{i-1})(qv_i - sv_i)^2}$$

Lower Bounding the Dynamic Time Warping

Recent approaches use the Minimum Bounding Envelope for bounding the **constrained** DTW

- Create a δ Envelope of the query Q (U, L)
- Calculate distance between MBE of Q and any sequence A
- One can show that: $D(MBE(Q)_{\delta},A) < DTW(Q,A)$
- δ is the constraint

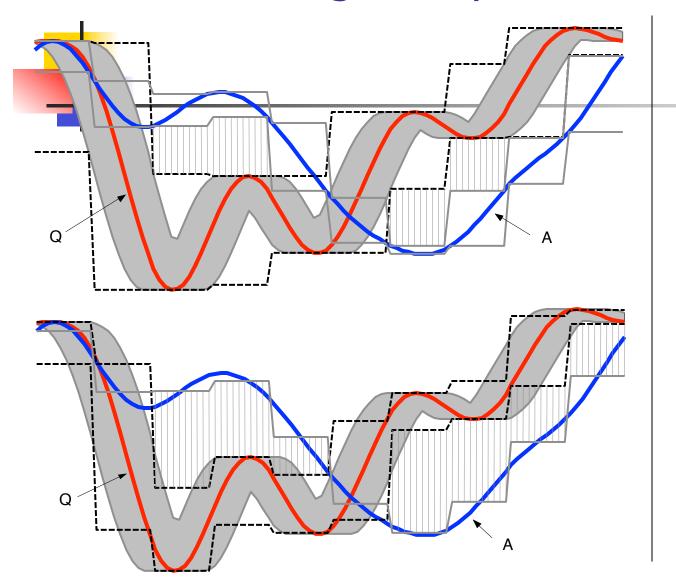
$$U[i] = \max_{-\delta \le r \le \delta} (Q[i+r])$$

$$L[i] = \min_{-\delta \le r \le \delta} (Q[i+r])$$

$$LB_{-}K(E(Q), A) = P$$

$$LB_{-}K(E(Q), A) = \int_{p}^{N} \begin{cases} |A[i] - U[i]|^{p} & if \quad A[i] > U[i] \\ |A[i] - L[i]|^{p} & if \quad A[i] < L[i] \\ 0 & otherwise \end{cases}$$

Lower Bounding the Dynamic Time Warping



LB by Keogh approximate MBE and sequence using MBRs LB = 13.84

LB by Zhu and Shasha approximate MBE and sequence using PAA

LB = 25.41

Computing the LB distance

- Use PAA to approximate each time series A in the sequence and U and L of the query envelop using k segments
- Then the LB_PAA can be computed as follows:

$$LB_PAA(\overline{E}(Q), \overline{A}) = \int_{i=1}^{k} \frac{N}{k} \begin{cases} \left| \overline{A}[i] - \overline{U}[i] \right|^{p} & if \quad \overline{A}[i] > \overline{U}[i] \\ \left| \overline{A}[i] - \overline{L}[i] \right|^{p} & if \quad \overline{A}[i] < \overline{L}[i] \end{cases}$$

$$0 \quad otherwise$$

where $\overline{A}[i]$ is the average of the i-th segment of the time series A, i.e.

$$\overline{A}[i] = \frac{k}{N} \sum_{j=\frac{N}{k}(i-1)+1}^{\frac{N}{k}i} A[j]$$

similarly we compute $\overline{U}[i]$ and $\overline{L}[i]$