

# Lecture Notes for Managing and Mining Multiplayer Online Games Summer term 2018

## Chapter 6: Temporal Analysis

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http://www.dbs.ifi.lmu.de/cms/VO\_Managing\_Massive\_Multiplayer\_Online\_Games

## **Chapter Overview**

- Behavior and Sequences
- Comparing Sequences
- Finding frequent subsequences
- Markov chains
- Hidden Markov-Chains
- Time series and feature-transformations
- Comparing time series
- Poisson-Processes

## player behavior

#### examples for player behavior

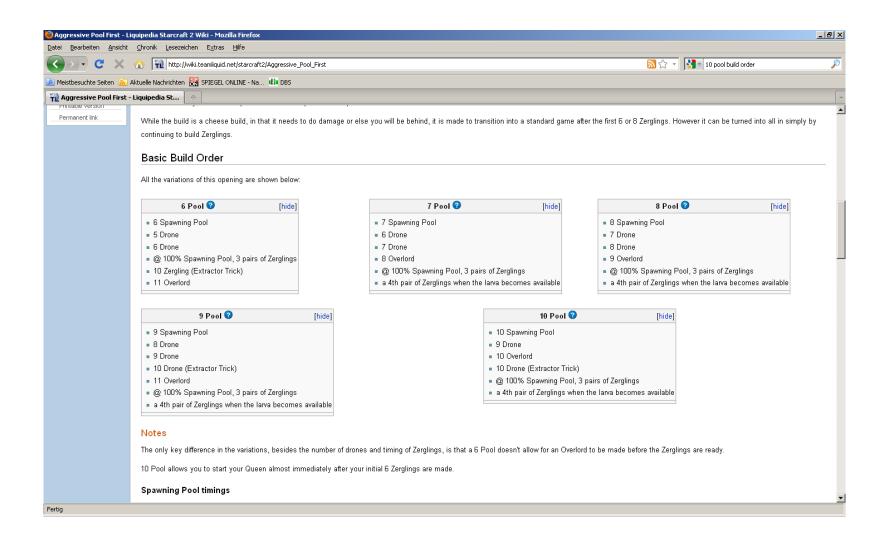
- sequence of moves in chess
- sequence of movement, action and interaction in a MMORPG
- sequence of orders to units in RTS Games
- behavior consists of a sequence of possible actions
- Simplest models for behavior are strings or sequences.

**Definition**: Let  $A=\{A_1, ..., A_n\}$  be a finite alphabet of n possible player actions, then the l-Tuple  $(a_1, ..., a_l) \in Ax...xA$  is a sequence of l length over A.

#### Remark:

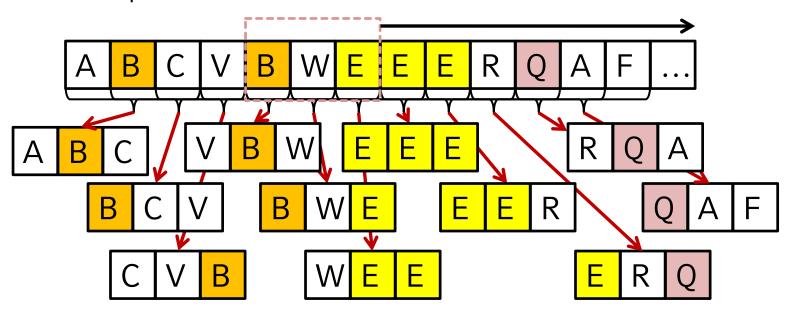
- Model describes only observations and does not differentiate between possible and impossible sequences.
- Model neglects the time between actions.

## Example: SC II Zerg Rushes



## Subsequences and Partitioning

- Which player is observed at a given time and for how long?
- The longer a player is observed, the less likely it becomes that another player behaves similarly
- To find typical behavioral patterns a sequence is usually divided into subsequences.
- Windowing (partitions a sequence)
   Slide a window of length k over the sequence and consider all subsequences. ( here k = 3)



## Subsequences and Partitioning

**problem**: A sequence of length I has I - (k-1) k-elemental subsequences and many of those are irrelevant.

**idea**: Only sequences appearing with a certain frequency are of interest.

#### **Frequent Subsequence Mining**

Find all subsequences in a sequence database appearing more frequently than *minsup*. (cf. Frequent Itemset Mining)

- $\Rightarrow$  length of the sequence is arbitrary.
- ⇒ search space is larger than the search space of itemset mining. (several occurrences of elements and orders)

## Frequent Subsequence Mining

- frequency fr(S,G) of S in sequence G:
   count occurrence of S in G
- relative frequency of S:

$$\varphi(S,G) = \frac{fr(S,G)}{|G| - |S| - 1}$$

sequence description of G:

$$\mathcal{S}(G) = \{ (S, \varphi(S, G)) \mid S \in G \}$$

mining sequential patterns is well explored
 => many approaches and algorithms

Properties of a Suffix Tree ST for the alphabet A with sequence G where IGI = n:

- to rule out ambivalence, words are padded with a terminal symbol (A), commonly \$.
- *ST* has exactly *n*+1 leaf nodes numbered from 0 to *n*, on the way from the root to the leaf *i* the suffix of length *n*-*i* is filed.
- Edges represent elements of A(\$) (uncompressed form), nonempty partial-sequences of A(\$) respectively
- Edges, emanating from the same starting node, must begin with different elements of *A*.

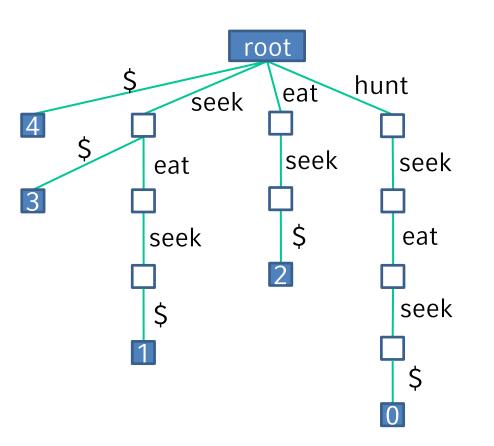
Creation in O([input string]), Search in O([query string])

- example: alphabet A ={eat, hunt, seek, flee, defend}
- insert:

```
S_1 = (seek, hunt, eat, seek)
S_2 = (seek, flee,hunt)
```

- example: Alphabet A ={eat, hunt, seek, flee, defend}
- insert:

```
S_1 = (hunt, seek, eat, seek) (hunt, seek, eat, seek, $) S_2 = (seek, flee, hunt) (seek, flee, hunt, $)
```



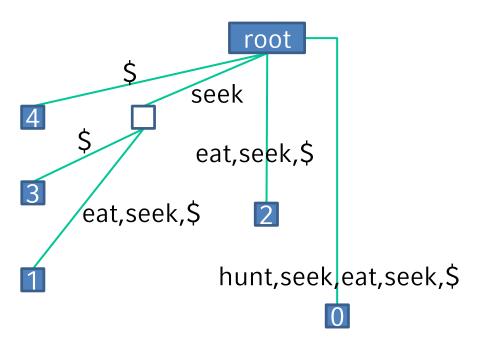
#### uncompressed variant:

every edge is labeled with an element of A{\$}

compressed variant: combine sub-paths without branches into one edge

- example: Alphabet A ={eat, hunt, seek, flee, defend}
- insert:

```
S_1 = (hunt, seek, eat, seek) (hunt, seek, eat, seek, $) S_2 = (seek, flee, hunt) (seek, flee, hunt, $)
```

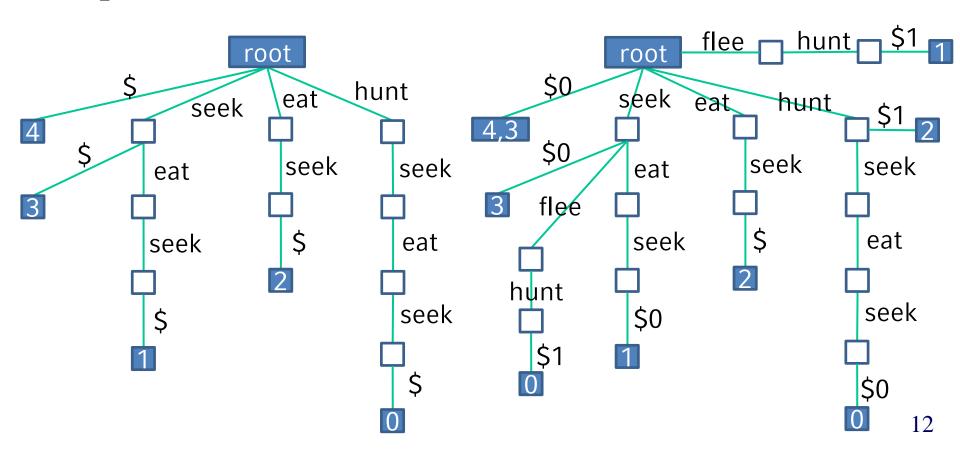


uncompressed variant: Every edge is labeled with an element of A{\$}

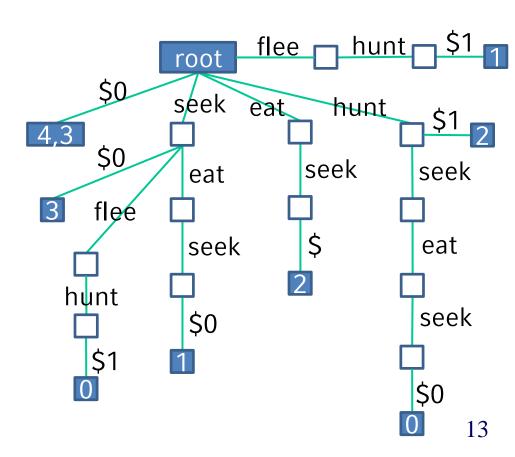
compressed variant: combine sub-paths without branches into one edge

- example: Alphabet A ={eat, hunt, seek, flee, defend}
- insert:

 $S_1$  = (hunt, seek, eat, seek) (hunt, seek, eat, seek, \$)  $S_2$  = (seek, flee, hunt) (seek, flee, hunt, \$)



- example: Alphabet A ={eat, hunt, seek, flee, defend}
- sample queries:
- Is q a Suffix?
- Is q a Substring?
- How often occurs *q*?



- Example: Alphabet A ={eat, hunt, seek, flee, defend}
- Sample request:
- Is q a Suffix?
- $\Rightarrow$  follow path (q\$) starting at root,

If reaching a leaf, then it is a Suffix

- Is q a Substring?
- => follow path (q) starting at root,

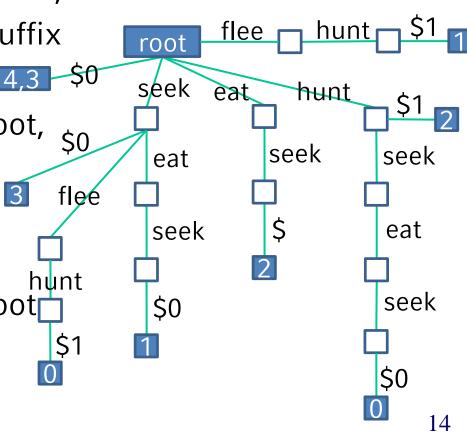
If processing possible,

then Substring

- How often occurs q?
- => follow path (q) starting at root

#leaves below terminal nodes

= #Occurences



## Comparing two Sequences

given: Alphabet A and a sequence database

DB  $\{(x_1, ..., x_k) | k \in IN \land x_i \in A \text{ for } 1 \le i \le k\}.$ 

task: compute the similarity of S1, S2  $\in$  DB.

**Hamming Distance**: number of different entries over all positions.

For 2 sequences with |S1|=|S2|=k:

$$Dist_{Ham}(S1, S2) = \sum_{i=0}^{k} \begin{cases} 0 & if \quad s_{1,i} = s_{2,i} \\ 1 & else \end{cases}$$

**Remark**: For sequences of different length, the shorter sequence is filled with the gap symbol "-".

example: S1 = (A,B,B,A,B) und S2 = (A,A,A,A,A)

$$(A,B,B,A,B)$$
  
 $(A,A,A,A,A)$  Dist<sub>Ham</sub>  $(S1,S2)=3$ 

#### Levenshtein Distance

- Hamming Distance: Computing the minimum cost to transform S1 into S2. Only substitutions of single elements are allowed in doing so. (Turn B into A.)
- Hamming Similarity: Counts the number of similar elements.
- idea: Extend the allowed transformations to include deletion and insertion of symbols.
- Levenshtein Distance: Minimum expense to transform *S1* into *S2* using 3 operations *Delete, Insert* and *Substitute*.

$$(A,B,B,A,B)$$
  $(A,B,B,A,B)$   $Sim_{Lev}(S1,S2)=3$ 

## Calculating Levenshtein Distance

given: Two sequences S1, S2 over the alphabet A with |S1|=n and |S2|=m.

task:  $Dist_{Lev}(S1,S2)$ 

Calculating Levenshtein Distance with dynamic programming:

Let D be a  $n \times m$ -Matrix over IN with:

$$\begin{split} D_{0,0} &= 0 \\ D_{0,i} &= i, \quad 0 \leq i \leq n \\ D_{j,0} &= j, \quad 0 \leq j \leq m \\ D_{i-1,j-1} &+ 0, falls \quad s_{1i} = s_{2,j} \\ D_{i-1,j-1} &+ 1, (Substitution) \\ D_{i,j-1} &+ 1, (Insertion) \\ D_{i-1,j} &+ 1, (Deletion) \end{split}$$
 für  $1 \leq i \leq n, \quad 1 \leq j \leq m$ 

After construction of matrix D,  $D_{n,m}$  contains the Levenshtein-distance between both input sequences.

## Example Levenshtein Distance

S1 = auto, S2 = ute

	ı	а	u	t	0
ı	0	1	2	3	4
u	1				
t	2				
е	3				



	-	а	u	t	0
-	0	1	2	3	4
u	1	1-:	1 <sup>V</sup>		
t					
е					



_		ı	а	u	t	0
	-	0	1	2	3	4
	u	1	1	1	2	3
	t	2	2	2	1	2
	е	3	3	3	2	2

	-	а	u	t	0
-	0	1	2	3	4
u	1	1	1	2	3
t	2	2	2	1	2
е	3	3	3	2	2

#### **Edit Distances**

- generalization of Levenshtein-Distances:
  - different cost matrix: substitution costs 4, deletion 1, insertion 2...
  - more operations:
    - transposing order

• duplicating, ...

$$(A,B,B,B,B)$$
 3 duplicates of B

costs may differ for different values:

$$Subst.(A,B) \neq Subst.(A,Z)$$

• works for sequences based on real-valued alphabets, for example: For A = IR: Subst(5,1) = I5-1I

## Markov Chains and Sequences

- sequences of actions are subject to certain rules
- modeling with finite automatons (testing sequence for validity)
- Markov chains are probabilistic automatas:
  - allowed state transitions
  - probability distributions for state transitions.
- 1<sup>st</sup> order Markov assumption: The state at time *t+1* depends solely on the state at time *t*.
- the order of a Markov chain is the number of predecessor states on which the choice of the next state might depend.

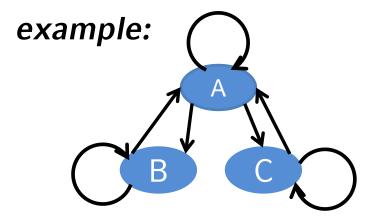
#### First Order Markov-Chains

**definition**: A Markov chain M is defined for a state set A and a stochastic transition-matrix  $|A| \times |A| = D$ .

#### explanations:

- A may contain a start- and a absorption-state (Modeling Start and End)
- stochastic Matrix: rows add up to 1.

(row *i* contains the distribution of successors for state *i*)



	-	Α	В	С
-	0.0	0.3	0.3	0.4
А	0.1	0.25	0.5	0.15
В	0.1	0.5	0.4	0.0
С	0.1	0.1	0.7	0.1

$$p(ACBB) = P(A \mid -) \cdot P(C \mid A) \cdot P(B \mid C) \cdot P(B \mid B) \cdot P(-\mid B)$$
  
= 0.3 \cdot 0.15 \cdot 0.4 \cdot 0.7 \cdot 0.4 \cdot 0.1

#### Hidden Markov Models

#### training a Markov chain:

 break the training sequence down into 2-grams and determe the relative frequency.
 (How often is A followed by B?)

$$P(B \mid A) = \frac{fr(AB)}{fr(A)}$$

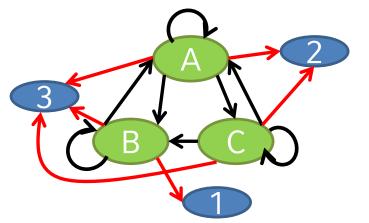
#### problem:

- observations often do not match the observed behavior:
  - action log is available, but game-play has to be analyzed
  - incorrect execution obfuscates actual intentions
  - analysis of an AI state changes
     (observed actions may be employed in different states)

#### Hidden Markov Models

**Definition**: A Hidden Markov Model M is defined by a state set A, a stochastic transition matrix  $|A| \times |A| = D$ , an observation set B and a stochastic output-matrix  $|A| \times |B| = F$ .

**Example**:  $A = \{A, B, C\}, B = \{1, 2, 3\}$ 



D	-	Α	В	С
-	0.0	0.3	0.3	0.4
Α	0.1	0.25	0.5	0.15
В	0.1	0.5	0.4	0.0
С	0.1	0.1	0.7	0.1

I	2	3
0.0	0.2	0.8
0.5	0.0	0.5
0.0	0.5	0.5
	0.5	0.0     0.2       0.5     0.0

*P(122)*: define all possible state triples, generated by 122 : BAA, BAC

$$P(122) = P(BAA) \cdot P(122 \mid BAA) + P(BAC) \cdot P(122 \mid BAC)$$

#### Use of HMM

- **Evaluation**: How likely is an observation  $O=(o_1,...,o_k)$  with  $o_i \in B$  for the HMM (A,B,D,F)? (Forward Estimation)
- **Recognition**: Given the observation  $O=(o_1, ..., o_k)$  and the HMM (A,B,D,F) which sequence  $(s_1, ..., s_k)$  with  $s_i \in A$  gives the best explanation for O? (Viterbi-Algorithm)
- Training: Given the observation O=(o<sub>1</sub>, ..., o<sub>k</sub>), how can we modify D and F to maximize P(OI(A,B,D,F))?
   (Baum-Welch Estimation)

#### **Evaluation: Forward Variables**

**given**:  $O = (o_1, ..., o_k)$  and (A, B, D, F)

**task:** P(O|(A,B,D,F))

**naive solution:** calculate P(O|S) for all k-elemental sequences S on A. (number grows exponentially with k)

improved solution: utilize Markov assumption

define forward-variable  $\alpha_{i}(t)$  as

$$\alpha_{j}(t) = P(o_{1}, o_{2}, ..., o_{t}, s_{t} = a_{j} | (ABDF))$$

calculation by induction:

$$\alpha_{j}(1) = d_{-,j} \cdot f_{j,o_{1}} \quad ,1 \leq j \leq |A|$$

$$\alpha_{j}(t+1) = \left(\sum_{i=1}^{|A|} \alpha_{i}(t) \cdot d_{i,j}\right) \cdot f_{j,ot+1} \quad ,1 \leq t \leq k-1$$

calculating with |A|<sup>2</sup>·k operations:

$$P(O \mid (A, B, D, F)) = \sum_{i=1}^{|A|} P(O, s_i = a_i \mid (A, B, D, F)) = \sum_{i=1}^{|A|} \alpha_i(k)$$

## Recognition: Viterbi Algorithm

given:  $O=(o_1, ..., o_k)$ , and Model (A, B,D,F).

task:  $S=(s_1, ..., s_k)$ , which maximizes P(O|S, (A, B, D, F)).

• define  $\delta(t)$  as the highest probability of a sequence on A of length t for the observation O.

$$\delta_{j}(t) = \max_{s_{1},...,s_{t-1}} P(s_{1},...,s_{t-1},O \mid (A,B,D,F))$$

calculation by induction

$$\begin{split} \mathcal{S}_{j}(1) &= d_{-,j} \cdot f_{j,o_{1}} &, 1 \leq j \leq |A| \\ \mathcal{S}_{j}(t+1) &= \left( \max_{1 \leq i \leq |A|} \left( \mathcal{S}_{i}(t) d_{i,j} \right) \right) \cdot f_{j,o_{t+1}} &, 1 \leq j \leq k-1 \\ \psi_{j}(1) &= 0 &, 1 \leq j \leq |A| \\ \psi_{j}(t+1) &= \arg\max_{1 \leq i \leq |A|} \left( \mathcal{S}_{i}(t) d_{i,j} \right) &, 1 \leq j \leq k-1 \end{split}$$

 similar to forward algorithm, but more efficient since only the best solution is pursued.

#### **Backward Variables**

analogously to Forward-Variable a Backward-Variable can be defined, used in training the HMM.

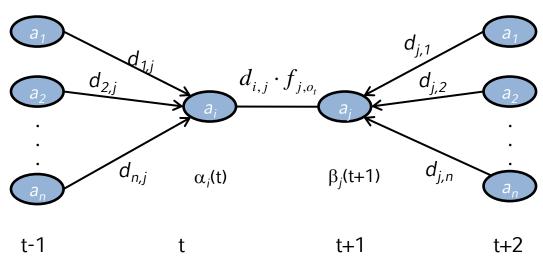
define Backward-Variable  $\beta_i(t)$  as

$$\beta_{j}(t) = P(o_{t+1}, ..., o_{k} \mid s_{t} = a_{j}, (ABDF))$$

Calculation by Induction:

$$\beta_i(k) = 1$$
 ,  $1 \le i \le |A|$ 

$$\beta_i(t-1) = \sum_{j=1}^{|A|} d_{i,j} \cdot f_{j,o_t} \cdot \beta_j(t)$$
 ,  $2 \le t \le k$ 



## Training: Baum-Welch Estimation

**given**:  $O = (o_1, ..., o_k)$ , A and B.

task: D, F, maximizing P(OI(A,B,D,F)).

Locally optimize solution with Expectation Maximization (EM)

Define  $\xi_{i,j}$  (t) as the likelihood of being in state  $a_i$  at the point in time t and being in state  $a_i$  at the point in time t+1:

$$\begin{split} \xi_{i,j}(t) &= & P(s_t = a_i, s_{t+1} = a_j \mid O, (A, B, D, F)) \\ &= \frac{\alpha_i(t) \cdot d_{i,j} \cdot f_{j,o_{t+1}} \beta_j(t+1)}{P(O \mid (A, B, D, F))} \\ &= \frac{\alpha_i(t) \cdot d_{i,j} \cdot f_{j,o_{t+1}} \beta_j(t+1)}{\sum_{k=1}^{|A|} \sum_{l=1}^{|A|} \alpha_k(t) \cdot d_{k,l} \cdot f_{l,o_{t+1}} \beta_j(t+1)} \end{split}$$

• Define  $\gamma_i$  (t) as the probability of being in state  $a_i$  at the point in time t:

$$\gamma_i(t) = \sum_{j=0}^{|A|} \xi_{i,j}(t)$$

## Training: Baum-Welch Estimation

- $\sum_{t=1}^{k-1} \xi_{i,j}(t)$  equals the expected number of state transitions from  $a_i$  to  $a_i$ .
- $\sum_{t=1}^{k-1} \gamma_i(t)$  equals the expected number of state transitions from  $a_i$  to other states.
- parameter are being recomputed as follows:

$$d_{-,a_i} = \gamma_i(1) \quad , d_{i,j} = \frac{\sum_{t=1}^{k-1} \xi_{i,j}(t)}{\sum_{t=1}^{k-1} \gamma_i(t)} \quad , f_{j,b_l} = \frac{\sum_{t \in \{t \mid o_t = b_l\}} \gamma_i(t)}{\sum_{t=1}^{k-1} \gamma_i(t)}$$

- training happens in alternating steps
  - calculate of  $\gamma_i$  (t),  $\xi_{i,j}$  (t) and P(OI(A,B,D,F))
  - updates of D and F (updates see above)
- algorithm terminates when
   P(OI(A,B,D,F)) grows less than

## Real-Value Sequences

- so far: Alphabet is a discrete domain
- Sequences can also be created based on real-value domains, for example  $IR^d$ .
- Frequent Pattern Mining on real-value domains is usually impossible.
- Comparing 2 real-value sequences on domain D with a distance function dist: D D IR<sub>0</sub>+.
  - Analogous to Hamming Distance one can determine the sum of distances for every position of the sequence.

$$dist_{sequ}(S_1, S_2) = \sum_{i=1}^{|S_1|} dist(s_{1,i}, s_{2,i}) + (|S_2| - |S_1|) \cdot \varphi, \quad für \quad |S_2| \ge |S_1|, \varphi \in IR^+$$

- Extension of edit distance is als possible: Substitution cost for *v* and *u* correlates to *dist(v,u)*.
- (More details follow later for Dynamic Time Warping)

#### Time series

 so far: sequences model the order of actions, but not the points in time.

but: in real time games timing is essential.

- ⇒ RTS games: build order are only effective if they can be realized in minimal time.
- ⇒ in MMORPGs the damage caused depends on the number of actions per time unit.
- ⇒ chess with chess clock: a move is also measured by the time needed to think.
- **time series**: Let *T* be a domain to model time and let *F* be an object presentation, then:

 $Z=((x_1,t_1),...,(x_l,t_l)) \in (F\times T)\times...\times (F\times T)$  is a time series of length l on F.

## **Examples for Time Series**

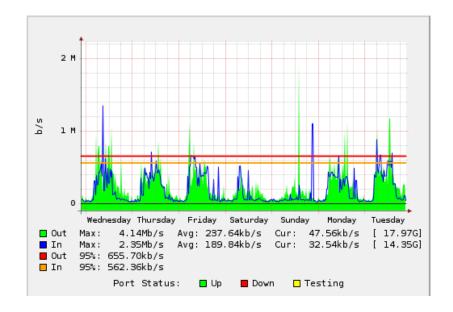
SC2-Logs: time series on discrete actions

```
0:00 TSLHyuN
                  Select Hatchery (10230)
0:00 TSLHyuN
                  Select Larva x3 (1027c,10280,10284), Deselect all
0:00 TSLHyuN
                  Train Drone
0:01 TSLHyuN
                  Train Drone
0:01 TSLHyuN
                  Select Drone x6 (10234,10238,1023c,10240,10244,10248),
Deselect all
0:01 TSLHyuN
                  Right click; target: Mineral Field (10114)
0:01 TSLHyuN
                  Deselect 6 units
0:02 TSLHyuN
                  Right click; target: Mineral Field (10170)
```

...

#### Network-Traffic:

- used in bot detection
- estimating game intensity

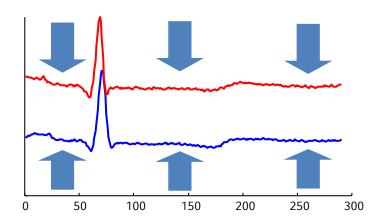


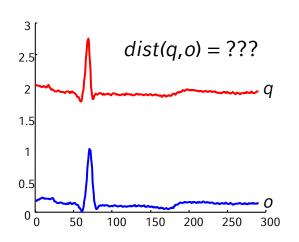
## Preprocessing Time series (1)

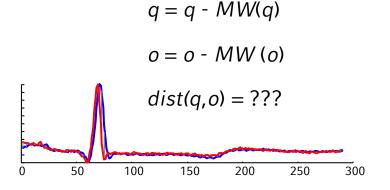
#### offset translation

- similar time series with different offsets
- shifting all time series around the
- mean *MW*:

1 
$$i |o|: o_i = o_i - MW(o)$$





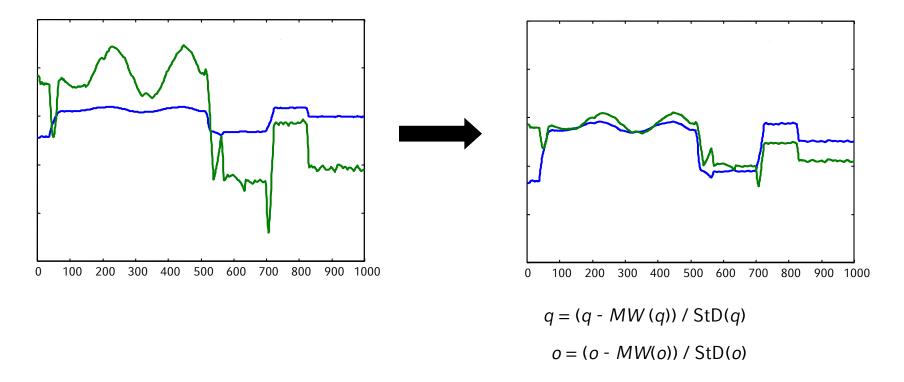


## preprocessing time series (2)

#### scaling amplitudes

- time series with similar progression but different amplitudes
- shifting the time series around the mean (MW) and normalizing the amplitude by standard deviation (StD):

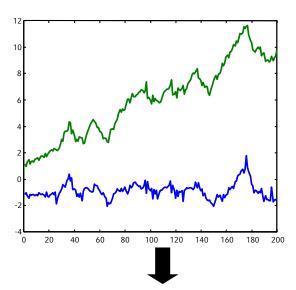
1  $i |o|: o_i = (o_i - MW(o)) / StD(o)$ 



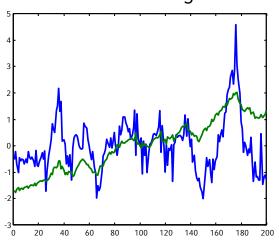
## preprocessing time series (3)

#### linear trends

- similiar time series with different trends
- Intuition:
  - determine regression line
  - move time series by means of this line

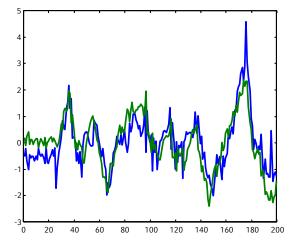


offset translation + amplitudes scaling



offset translation + Amplitudes scaling

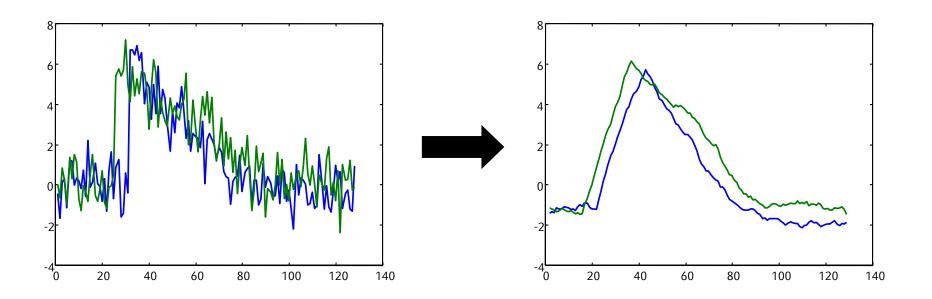




## Preprocessing time series (4)

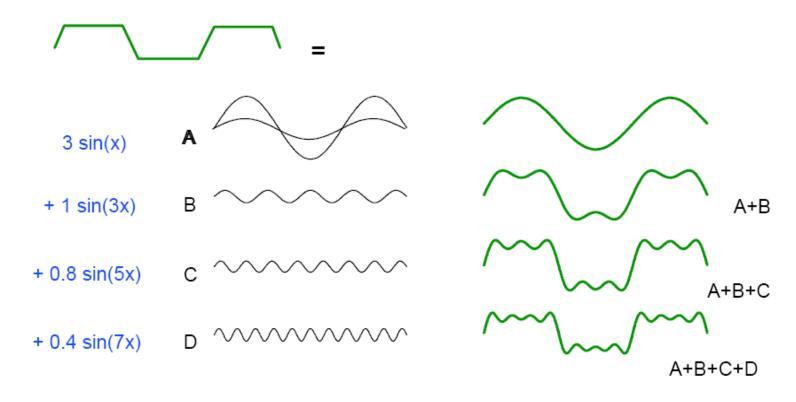
#### rectifying noise

- similar time series with a large amount of noise
- smoothing: determine for every value  $o_i$  the mean over all values  $[o_{i-k}, ..., o_i, ..., o_{i+k}]$  for a given k.



#### idea:

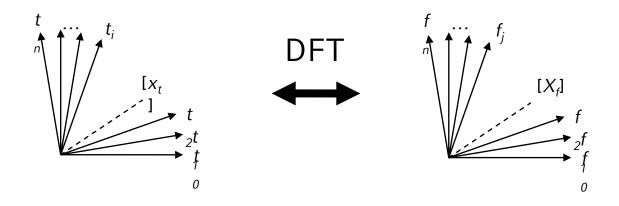
- describe arbitrary periodic functions as weighted sum of periodic base functions with different frequencies. A time series turns into a vector of constant length.
- base functions: sin and cos



**Fourier's theorem:** A periodic function (which is reasonable continuous) may be expressed as the sum of a series of sine and cosine terms with a specific amplitude.

#### properties:

- transformation does not change a function, only the presentation
- transformation is reversible => inverse DFT
- analogy: change of base in vector calculation



#### formal:

- given a time series of length n:  $x = [x_t]$ , t = 0, ..., n 1
- the DFT of x is a sequence  $X = [X_f]$  of n complex numbers for the frequencies f = 0, ..., n-1 with

$$X_{f} = \frac{1}{\sqrt{n}} \sum_{t=0}^{n-1} x_{t} \cdot e^{\frac{-i \cdot 2\pi \cdot f \cdot t}{n}} = \frac{1}{\sqrt{n}} \sum_{t=0}^{n-1} x_{t} \cos\left(\frac{2 \cdot \pi \cdot f \cdot t}{n}\right) - i \cdot \frac{1}{\sqrt{n}} \sum_{t=0}^{n-1} x_{t} \sin\left(\frac{2 \cdot \pi \cdot f \cdot t}{n}\right)$$
Real part Imaginary part

where *i* identifies the imaginary unit viz.  $i^2 = -1$ .

 the real part indicates the share of the cosine functions, whereas the imaginary part indicates the share of sine functions of the frequency f.

the inverse DFT restores the original signal:

$$x_{t} = \frac{1}{\sqrt{n}} \sum_{f=0}^{n-1} X_{f} \cdot e^{\frac{i \cdot 2 \cdot \pi \cdot f \cdot t}{n}}$$

$$t = 0, ..., n-1 \quad (t: \text{ points in time})$$

$$[x_{t}] \leftrightarrow [X_{f}] \text{ describes a Fourier-Paar,}$$

$$\text{viz. DFT}([x_{t}]) = [X_{f}] \text{ and DFT}^{-1}([X_{f}]) = [x_{t}].$$

- the DFT is a **linear map**, viz. from  $[x_t] \leftrightarrow [X_f]$  and  $[y_t] \leftrightarrow [Y_f]$  follows:
  - $[x_t + y_t] \leftrightarrow [X_t + Y_t]$  and
  - $[ax_t] \leftrightarrow [aX_f]$  for a Scalar *a* IR
- energy of a sequence
  - energy E(c) of c is the square of the amplitude:  $E(c) = |c|^2$ .
  - energy E(x) of a sequence x is the sum of all energies of the sequence:

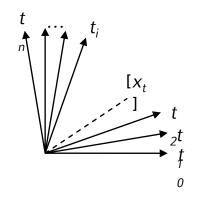
$$E(x) = ||x||^2 = \sum_{t=0}^{n-1} |x_t|^2$$

**Parseval's theorem**: Energy of a signal in a time range equals the energy in the frequency range.

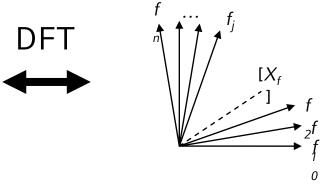
**Formal**: Let *X* the DFT of *x*, then follows:

$$\sum_{t=0}^{n-1} |x_t|^2 = \sum_{t=0}^{n-1} |X_f|^2$$

• consequence from Parseval's theorem and the DFT's linearity: The euclidean distance of two signals x and y correspond in time and frequency range:  $||x-y||^2 = ||X-Y||^2$ 







"Frequency range (-space)"

### **Basic Idea of query processing:**

The euclidean distance is used as a sequence's similarity function:

$$dist(x, y) = ||x - y|| = \sqrt{\sum_{t=0}^{n-1} |x_t - y_t|^2}$$

- parseval's theorem allows for distances to be calculated in the frequency range instead of the time range: dist(x,y) = dist(X,Y)
- in practice the lowest frequencies are the most important.
- the first frequency coefficients contain the most important information.
- for indexing the transformed sequences are shortened, for  $[X_f]$ , f = 0, 1, ..., n-1 coefficients only the first c coefficients  $[X_f < c]$ , c < n are indexed.

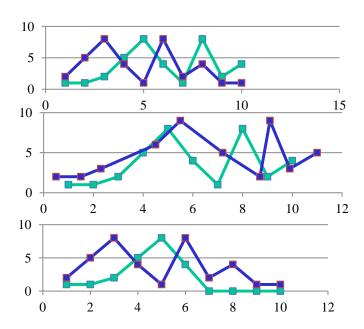
$$dist_c(x, y) = \sqrt{\sum_{f=0}^{c-1} |x_f - y_f|^2} \le \sqrt{\sum_{f=0}^{n-1} |x_f - y_f|^2} = dist(x, y)$$

- for the index a lower bound of the true distance can be calculated:
   filter-refinement:
  - filter step is based on shortened time series (index assisted)
  - refinement step determines true hits on complete time series

## **Distances of Time Series**

problems: Which points in time are to be compared?

- offset at the beginning:
   S2 is shifted in time to S1.
- clocking of reading: points in time of measuring differ.
- length of time series: measuring periods differ.

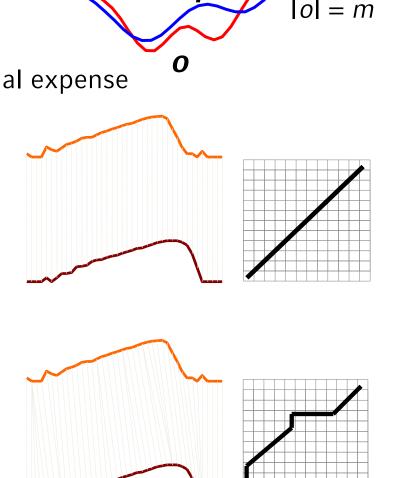


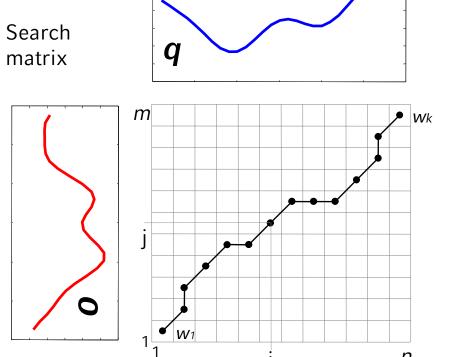
- time series with the same clocking and length can be compared as vectors. (dimension = point in time)  $Dist_{timeseries}(S1, S2) = \sum_{t=1}^{T} dist_{obj}(s_{1t}, s_{2t})$
- for variable length, clocking and offsets: adaption of edit-distance for sequences => Dynamic Time Warping

# Dynamic Time Warping Distanz

### calculation:

- given: time series q and o of different length
- find mapping of all  $q_i$  to o with minimal expense





|q| = n

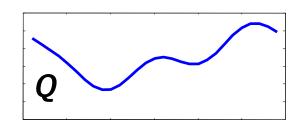
## Dynamic Time Warping Distance

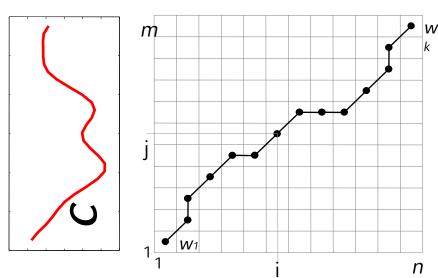
#### **Search Matrix**

- All possible mappings q to o can be interpreted as a "warping" path within the search matrix
- Of all these mappings, we search for the path with the lowest cost

$$DTW(q, o) = \min \left\{ \sqrt{\sum_{k=1}^{K} w_k} / K \right\}$$

Dynamic Programming
 => Run-time (n · m)
 (see Edit Distances)

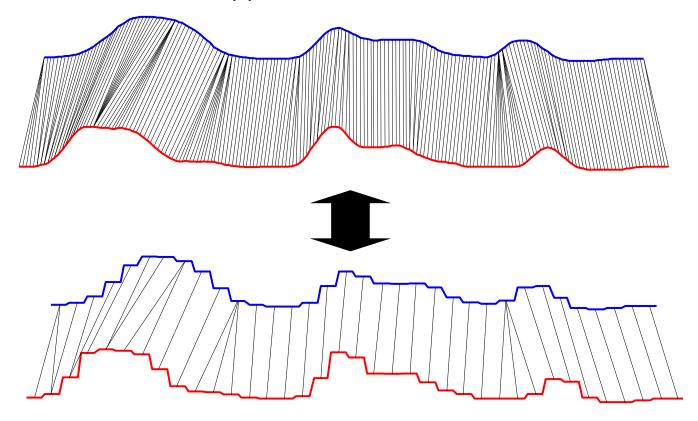




# Approximate Dynamic Time Warping Distance

### idea:

- approximate the time series (compressed representation, Sampling, ...)
- calculate DTW for the approximates

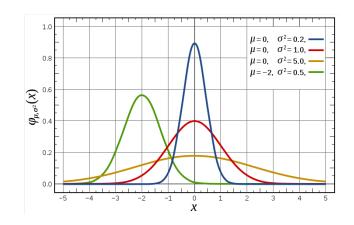


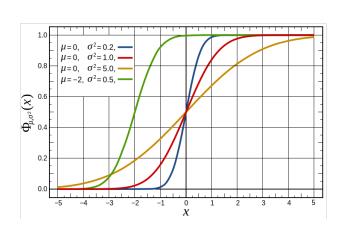
## Statistic Models for Time

### problem:

How is the time of the next action modeled?

- ⇒ statistic models for the time between two events is necessary.
- ⇒ time is a continuous variable => probability density function
- $\Rightarrow$  task: compute the probability for the next event *e* occurring within the time frame t+t.
- ⇒ the cumulative probability density function describes this probability

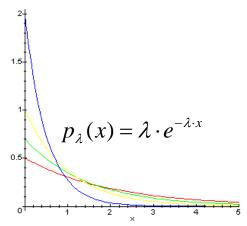




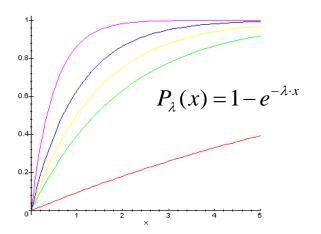
# Homogeneous Poisson Processes

- simplest process to model time
- points in time between 2 events are exponentially distributed
- probability density of the exponential distribution:  $p_{\lambda}(x) = \lambda \cdot e^{-\lambda \cdot x}$
- integration yields the cumulative density function describing the probability of the next action happening in the time interval between 0 ... x.

$$P_{\lambda}(x) = \int_{0}^{x} p_{\lambda}(t)dt = 1 - e^{-\lambda \cdot x}$$



Density function of the exponential distribution



Accumulated density function of the exponential distribution

## Parameter assessment

**given**: A training set of points in time  $X=\{x_1, ..., x_n\}$ , which are distributed exponentially.

task: The most likely value for the intensity parameter.

Approximation with Maximum Likelihood

=> Search the value of  $\lambda$  with the highest probability of generating X. Likelihood function L for Sample X:

$$L_X(\lambda) = \prod_{i=1}^n \lambda \cdot e^{-\lambda \cdot x_i} = \lambda^n \cdot e^{-\lambda \cdot \sum_{i=1}^n x_i} = \lambda^n \cdot e^{-\lambda \cdot n \cdot E(X)} \qquad mit \quad E(X) = \frac{\sum_{i=1}^n x_i}{n}$$

Differentiate the log-likelihood for  $\lambda$  and set the gradient to zero:

$$\frac{d}{d\lambda} \ln L(\lambda) = \frac{d}{d\lambda} (n \cdot \ln(\lambda) - \lambda \cdot n \cdot E(X)) = \frac{n}{\lambda} - n \cdot E(X)$$

$$\Rightarrow \lambda^* = \frac{1}{E(X)}$$

# Learning Goals

- Sequences and time series
- Frequent Subsequence Mining with Suffix-Trees
- Distance measuring sequences
  - Hamming Distance
  - Levenshtein Distance
- Markov-Chains
- Hidden Markov chains
- Time series and preprocessing steps
- Dynamic Time Warping
- Poisson processes

### Literature

- Kyong Jin Shim, Jaideep Srivastava: Sequence Alignment Based Analysis of Player Behavior in Massively Multiplayer Online Role-Playing Games (MMORPGs), in Proceedings of the 2010 IEEE International Conference on Data Mining Workshops, 2010.
- Ben G. Weber, Michael Mateas: A data mining approach to strategy prediction, in Proceedings of the 5th International Conference on Computational Intelligence and Games, 2009.
- K.T. Chen, J.W. Jiang, P. Huang, H.H. Chu, C.L. Lei, W.C. Chen: Identifying MMORPG bots: A traffic analysis approach, In Proceedings of the 2006 ACM SIGCHI International Conference on Advances in Computer Entertainment Tsechnology, 2006.