

Lecture notes
Knowledge Discovery in Databases
Summer Semester 2012

Lecture 2: Data Preprocessing / Feature spaces

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[http://www.dbs.ifi.lmu.de/cms/Knowledge_Discovery_in_Databases_I_\(KDD_I\)](http://www.dbs.ifi.lmu.de/cms/Knowledge_Discovery_in_Databases_I_(KDD_I))

- Data preprocessing
- Features
- Basic data descriptors
- Feature spaces and distance functions
- Feature transformation for text data
- Things you should know
- Homework/tutorial

Why data preprocessing?

- Real world data are noisy, incomplete and inconsistent:
 - Noisy: errors/ outliers
 - o erroneous values : e.g. salary = -10K
 - o unexpected values: e.g. salary=100K when the rest dataset lies in [30K-50K]
 - Incomplete: missing data
 - o missing attributes of interest: e.g. no information on occupation
 - o missing values: e.g., occupation=" "
 - Inconsistent: discrepancies in the data
 - o e.g. student ratings between different universities might differ
- “Dirty” data → poor mining results
- Data preprocessing is necessary for improving the quality of the mining results !!!

Major tasks in data preprocessing

- Data cleaning:
 - Fill in missing values, smooth noisy data, identify or remove outliers, and resolve inconsistencies
- Data integration:
 - Integration of multiple databases, data cubes, or files (Entity identification, Value resolution)
- Data transformation:
 - e.g. normalization in a given range
 - Generalization through e.g. some concept hierarchy
- Data reduction:
 - Aggregation
 - Dimensionality reduction
 - Duplicate elimination

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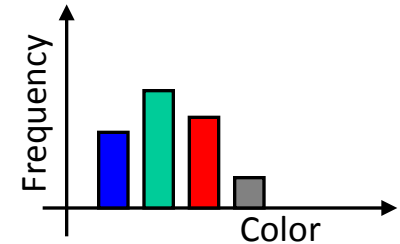
Data objects/ Examples/Instances

- Datasets consists of objects /examples / instances
 - e.g., in a movie database: movies, actors, director,...
 - e.g., in a library database: books, users, loans, publishers,
 - e.g., in a university database: students, professors, courses, grades,...
- Objects are described through features/ attributes/ variables
 - E.g. in a database table, the rows are the objects, the columns are the attributes/features/variables.

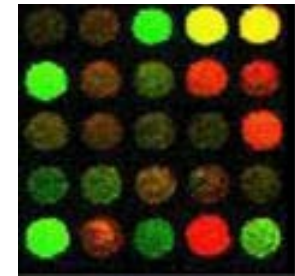
id	person	name	web	bio	location	following	followers
8	1	Justin Bieber	http://www.youtube.com/justinbieber	www.BieberFever.comRequest my NEW...	on the MY WORLD TOUR!!!	88045	5792472
9	2	Perez Hilton	http://www.PerezHilton.com	Perez Hilton is the creator and writer of 	...	Hollywood, California	341	2566369
10	3	Paris Hilton	http://www.parishilton.com	Hugel!	UT: 35.975487;-115.141709	842	2915057
11	4	Britney Spears	http://www.britneyspears.com	It's Britney Bitch!	Los Angeles, CA	417405	6168589
12	5	Kim Kardashian	http://kimkardashian.celebuzz.com/	business woman, exec producer, fashion...	on a plane...	96	5139761
13	6	Mariah Carey				0	3400111
14	7	Shakira	http://www.shakira.com	Welcome to Shakira's Official Twitter pag...	Bahamas	33	3318367
15	8	Justin Timberlake	http://www.justintimberlake.com	Official Justin Timberlake Twitter.	Memphis, TN	19	3339151
16	9	Gov. Schwarzenegger	http://gov.ca.gov	As California's 38th Governor I look forwa...	Sacramento, California	110763	1824274
17	10	Serena Williams	http://www.serenawilliams.com	Living, Loving, and working to help you...	Paris	84	1819778
18	11	Larry King	http://www.CNN.com/LarryKing	CNN's Larry King Live	LA	183	1721681
19	12	Panos Ipeirotis	http://behind-the-enemy-lines.blogspot.com/	Associate Professor at Stern School of B...	New York, NY	156	547
..	..	..	..	..	..	..	..

ID	title	URL	unknc	Action	Adventure	Animation	Childrens	Comedy	Crime	Documentar	Drama	Fantasy	FilmNoir	Horror	Musical	Mystery	Romance	SciFi	Thrille
1	Toy Story (1995)	http://us.imdb	0	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0
2	GoldenEye (1995)	http://us.imdb	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3	Four Rooms (1995)	http://us.imdb	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
4	Get Shorty (1995)	http://us.imdb	0	1	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0
5	Copycat (1995)	http://us.imdb	0	0	0	0	0	0	1	0	1	0	0	0	0	0	0	0	0
6	Shanghai Triad (Yao a yao dao waipo qiao	http://us.imdb	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
7	Twelve Monkeys (1995)	http://us.imdb	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	0
8	Babe (1995)	http://us.imdb	0	0	0	0	1	1	0	0	1	0	0	0	0	0	0	0	0
9	Dead Man Walking (1995)	http://us.imdb	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
10	Richard III (1995)	http://us.imdb	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
11	Seven (Se7en) (1995)	http://us.imdb	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
12	Usual Suspects, The (1995)	http://us.imdb	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0

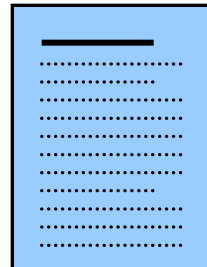
Image databases:
Color histograms



Gene databases:
gene expression level



Text databases:
Word frequencies



Data	25
Mining	15
Feature	12
Object	7
...	

The feature-approach allows uniform treatment of objects of different classes of applications.

- **Binary/ Dichotomous variables:** the attribute can take 2 values 0 or 1
 - usually, 0 means absence, 1 means presence
 - Symmetric binary: both outcomes equally important:
 - e.g., gender
 - Asymmetric binary: outcomes not equally important.
 - e.g., medical test (positive vs. negative)
 - Convention: assign 1 to most important outcome (e.g., HIV positive)
- **Categorical/ Nominal variables:** the attribute can take values within a set of M categories/ states.
 - e.g., eye color = {brown, green, blue}, occupation = {engineer, doctor, teacher}
 - There is *no ordering* in the categories/ states.
 - Only *equal/ different* relationships apply.
- **Ordinal variables:** similar to categorical variables, but the M states are ordered/ ranked in a meaningful way.
 - e.g., movie rating = {dislike, indifferent, like}, medal = {bronze, silver, gold}
 - There is an *ordering* between the values. But, we don't know the magnitude between successive values. The distance between these values has no meaning.

Basic feature types

Numeric variables: quantitative (integer or real)

- **Interval-scale variables**

- continuous measurements of a roughly linear scale
 - e.g., weight, height, latitude, longitude, temperature C°, citation counts
- Measured on a scale of equal-sized units
 - It is assumed that the intervals keep the same importance throughout the scale.
- Except for ordering, we can quantify and compare the magnitudes of differences between them.
 - e.g. $40^{\circ}\text{C} > 30^{\circ}\text{C}$, 40°C is twice as high as 20°C , and an increase from 20°C to 40°C is twice as much as the increase from 30°C to 40°C .

- **Ratio-scale variables**

- continuous positive measurements on a nonlinear scale, e.g. in an exponential scale.
 - e.g., the growth of bacterial population (say, with a growth function Ae^{Bt}). In this model, equal time intervals multiply the population by the same ratio.
 - e.g., the decay of a radioactive element

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Let $x_1, \dots, x_n$ be a random sample of an attribute X . Measures of central tendency of X include:

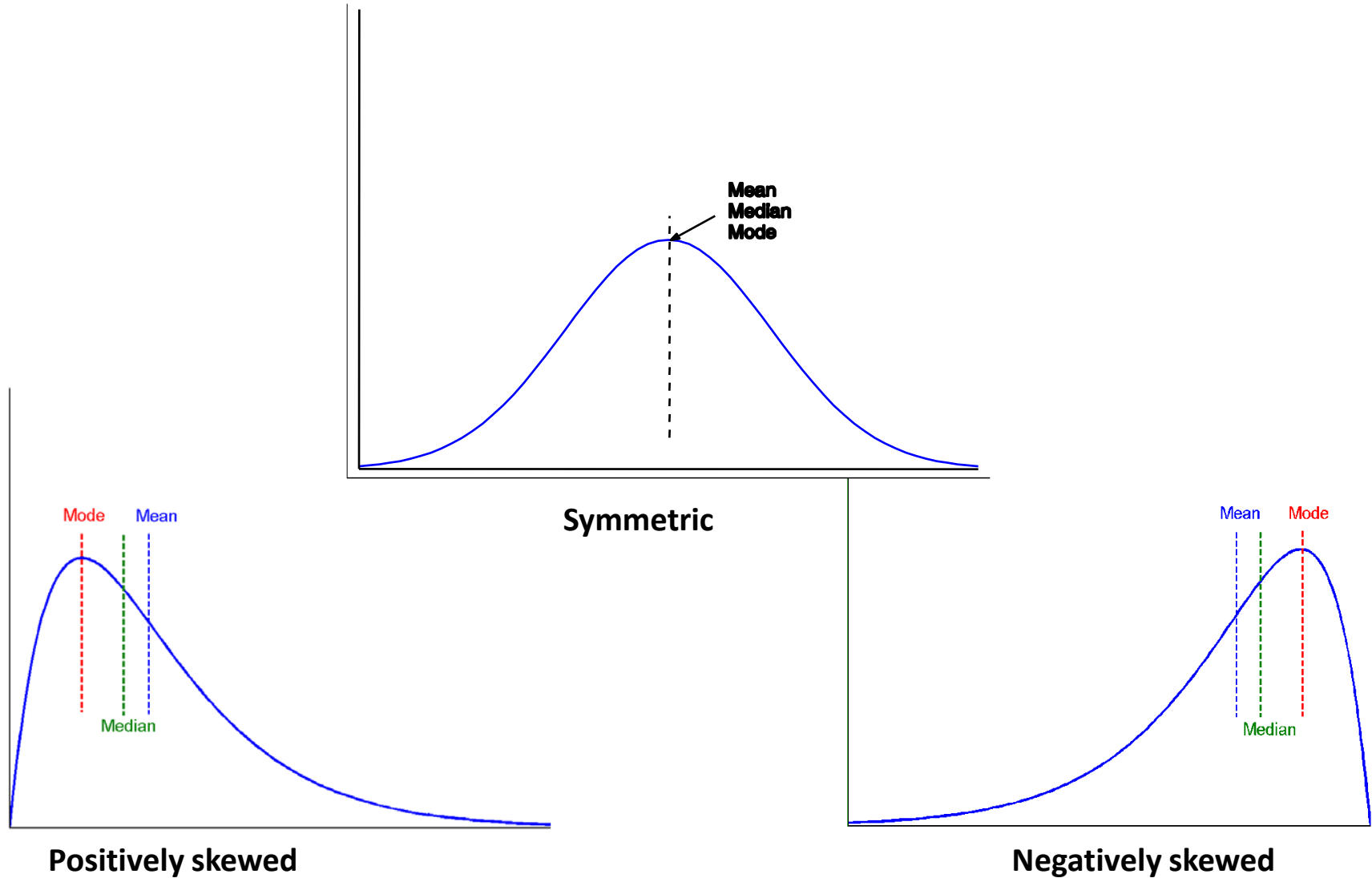
- (Arithmetic) mean/ center/ average: $\longleftarrow$ Algebraic measure

$$\bar{x} = \frac{1}{n} \cdot \sum_{i=1}^n x_i$$

- Weighted average: $\bar{x} = \frac{\sum_{i=1}^n w_i x_i}{\sum_{i=1}^n w_i}$

- Median: the central element in ascending ordering $\longleftarrow$ Holistic measure
 - Middle value if odd number of values, or average of the middle two values otherwise
- Mode: Value that occurs most frequently in the data $\longleftarrow$ Holistic measure
 - Unimodal, bimodal, trimodal

Symmetric vs Skewed data



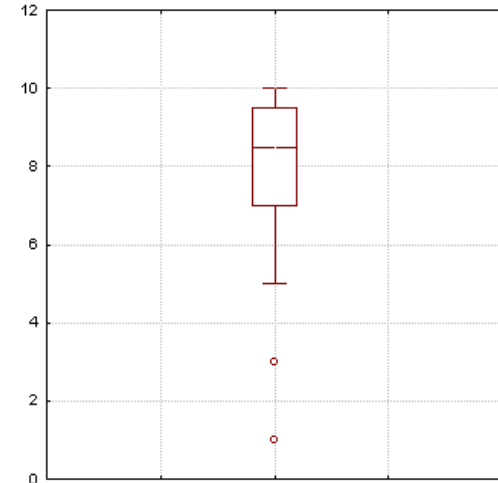
Let $x_1, \dots, x_n$ be a random sample of an attribute X . The degree to which X values tend to spread is called dispersion or variance of X :

- Range: max value – min value
- Q_1 (25th percentile), Q_3 (75th percentile)
 - Median is the 50th percentile
- 5 number summary: min, Q_1 , median, Q_3 , max
 - Boxplots to visualize them
- Variance σ^2 :

$$\sigma^2 = \frac{1}{n} \sum_{i=1}^n (x_i - \bar{x})^2 = \frac{1}{n} \left[\sum_{i=1}^n x_i^2 - \frac{1}{n} \left(\sum_{i=1}^n x_i \right)^2 \right]$$

- Standard deviation σ :

$$\sigma = \sqrt{\frac{1}{n} \cdot \sum_{i=1}^n (x_i - \bar{x})^2}$$

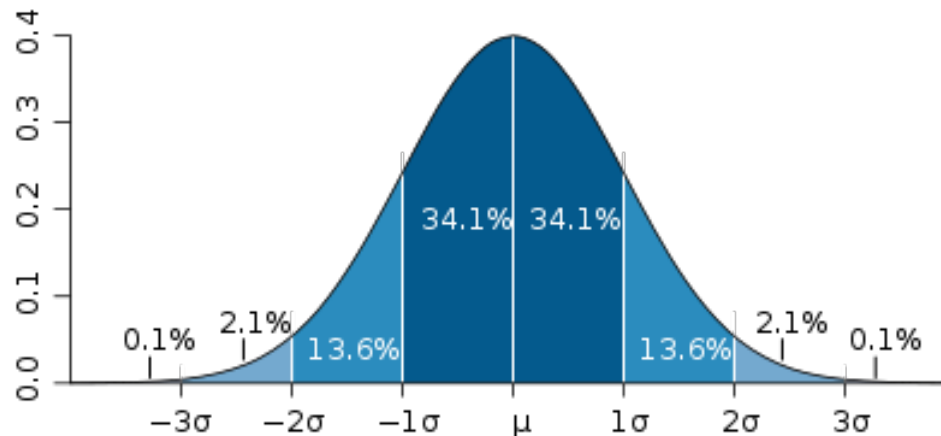


Source:
<http://de.wikipedia.org/wiki/Boxplot>

Example: Normal distribution

The normal distribution curve

- ~68% of values drawn from a normal distribution are from $\mu - \sigma$ to $\mu + \sigma$
- ~95% of the values lie from $\mu - 2\sigma$ to $\mu + 2\sigma$
- ~99.7% of the values are from $\mu - 3\sigma$ to $\mu + 3\sigma$

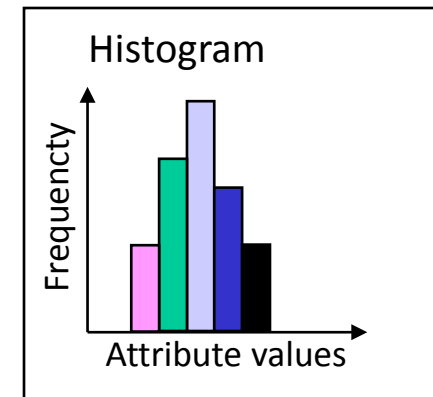


Source: http://en.wikipedia.org/wiki/Normal_distribution

Let $x_1, \dots, x_n$ be a random sample of an attribute X .

For the visual inspection of the data, several types of graphs are useful, e.g.:

- Boxplots
 - 5 number summary
- Histograms:
 - Summarizes the distribution of X
 - X axis: attribute values, Y axis: frequencies
 - Absolute frequency: for each value a , # occurrences of a in the sample
 - Relative frequency: $f(a) = h(a)/n$
- Different types of histograms, e.g.:
 - Equal width:
 - It divides the range into N intervals of equal size
 - Equal frequency/ depth:
 - It divides the range into N intervals, each containing approximately same number of samples



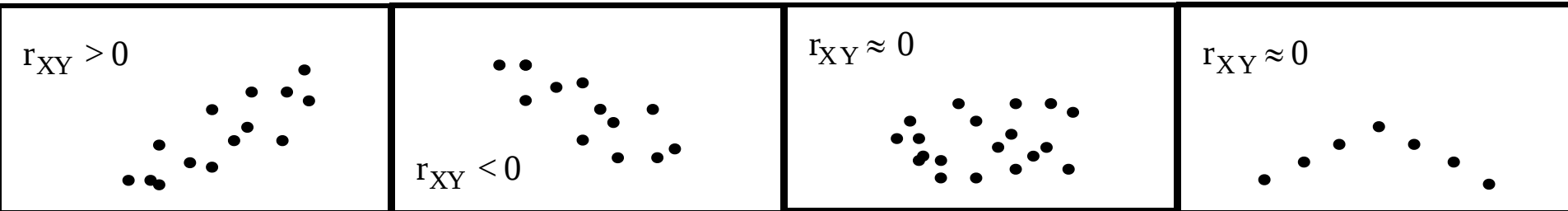
- Given two attributes X , Y one can measure how strongly they are correlated
- For numerical data $\rightarrow$ correlation coefficient
- For categorical data $\rightarrow \chi^2$ (chi-square)

Correlation coefficient (also called Pearson's product moment coefficient) :

$$r_{XY} = \frac{\sum_{i=1}^n (x_i - \bar{x}) \cdot (y_i - \bar{y})}{n\sigma_X\sigma_Y} = \frac{\sum_{i=1}^n (x_i y_i) - n\bar{x}\bar{y}}{n\sigma_X\sigma_Y}$$

– n: # tuples; x_i, y_i : the values in the i^{th} tuple for X, Y

- $-1 \leq r_{XY} \leq 1$
- the higher r_{XY} the stronger the correlation
 - $r_{XY} > 0$ positive correlation
 - $r_{XY} < 0$ negative correlation
 - $r_{XY} \sim 0$ no correlation/ independent



Visual inspection of correlation

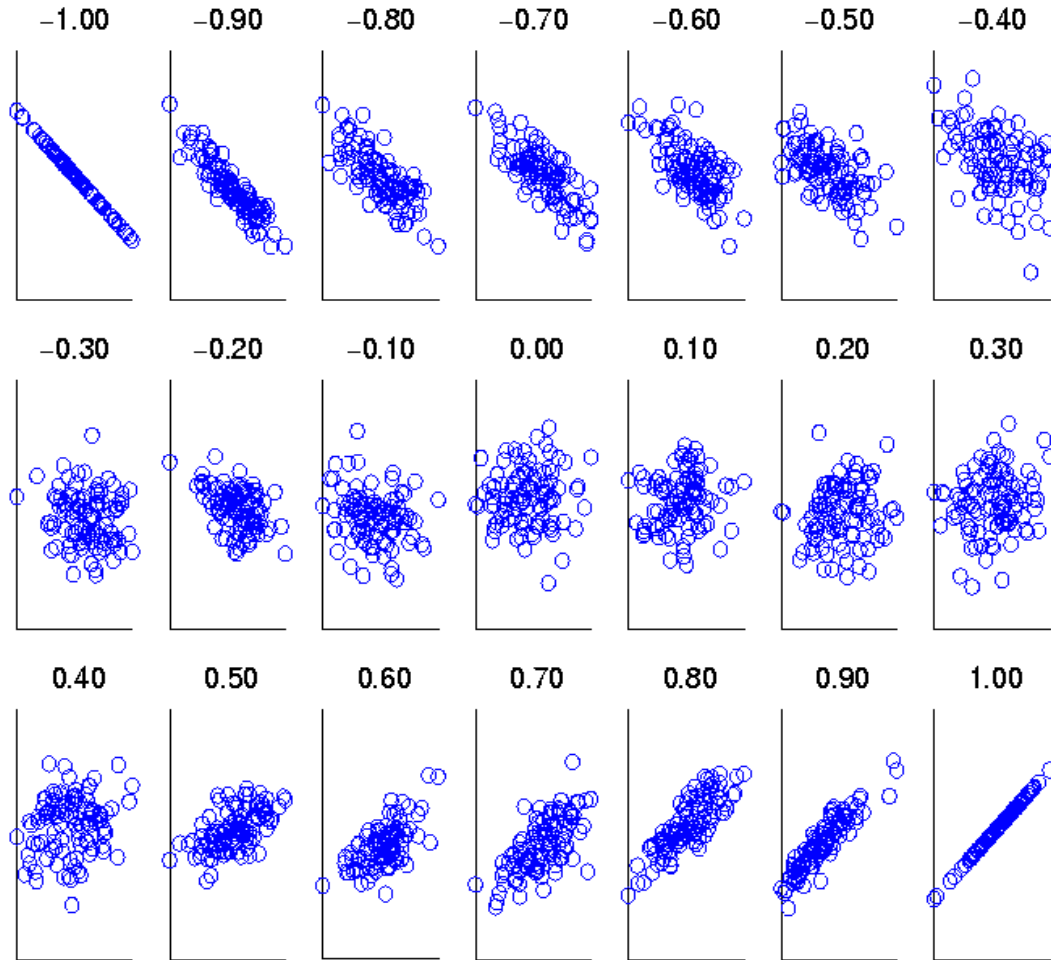


Figure 5.11. Scatter plots illustrating correlations from -1 to 1.

Contingency table

- For categorical/ nominal features $X=\{x_1, \dots, x_c\}$, $Y=\{y_1, \dots, y_r\}$
- Represents the absolute frequency h_{ij} of each combination of values (x_i, y_j) and the marginal frequencies h_i, h_j of X, Y .

Attribute X	Attribute Y		
	Medium-term unemployment	Long-term unemployment	Total
No education	19	18	37
Teaching	43	20	63
Total	62	38	100

Chi-square χ^2 test

$$\chi^2 = \sum_{i=1}^c \sum_{j=1}^r \frac{(o_{ij} - e_{ij})^2}{e_{ij}}$$

o_{ij} : observed frequency
 e_{ij} : expected frequency

$$e_{ij} = \frac{h_i h_j}{n}$$

Chi-square example

	Play chess	Not play chess	Sum (row)
Like science fiction	250 (90)	200 (360)	450
Not like science fiction	50 (210)	1000 (840)	1050
Sum(col.)	300	1200	1500

- χ^2 (chi-square) calculation (numbers in parenthesis are expected counts calculated based on the data distribution in the two categories)

$$\chi^2 = \frac{(250 - 90)^2}{90} + \frac{(50 - 210)^2}{210} + \frac{(200 - 360)^2}{360} + \frac{(1000 - 840)^2}{840} = 507.93$$

- It shows that like_science_fiction and play_chess are correlated in the group

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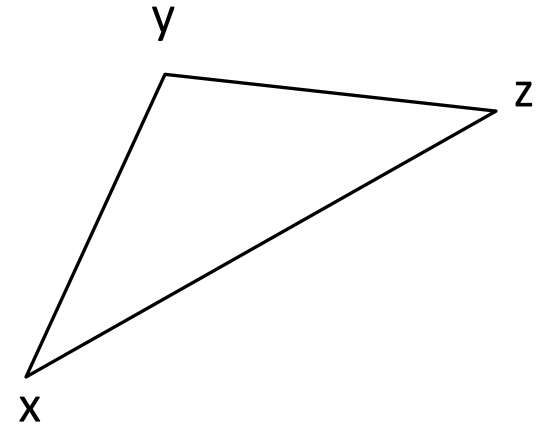
Feature spaces and distance functions

Feature space

- Intuitively: a domain with a distance function
- Formally: feature space $\mathbf{F} = (Dom, dist)$
- Dom is a set of attributes / features
- $dist : Dom \times Dom \rightarrow \mathbb{R}_0^+$
- For all x, y in Dom , $x \neq y$, $dist$ is required to satisfy the following properties:
 - $dist(x, y) > 0$ (strictness)
 - $dist(x, x) = 0$ (reflexivity)
 - $dist(x, y) = dist(y, x)$ (symmetry)

Metric space

- Formally: Metric space $\mathbf{M} = (Dom, dist)$ with the following property:
 - $\mathbf{M}$ is a feature space
 - $\forall x, y, z \in Dom : dist(x, z) \leq dist(x, y) + dist(y, z)$ (triangle inequality)
- Famous example: Euclidean vector space
 - Formally: Euclidean vector space $\mathbf{E} = (Dom, dist)$:
 - $(Dom, dist)$ is a metric space
 - $Dom = \mathbb{R}^d$
- Sprechweise:**
 - Euclidean vector space = “Feature space”
 - Vectors (Objects in the Euclidean feature space) = “Feature vectors”
 - The d dimensions of the vector space = “Features”



Feature spaces and distance functions

Similarity/ distance between feature vectors (Euclidean vectors)

- Manhattan-Norm (L_1): $dist_2 = |p_1 - q_1| + |p_2 - q_2| + \dots$

The sum of the absolute differences of their coordinates

- Euclidean Norm (L_2): $dist_1 = ((p_1 - q_1)^2 + (p_2 - q_2)^2 + \dots)^{1/2}$

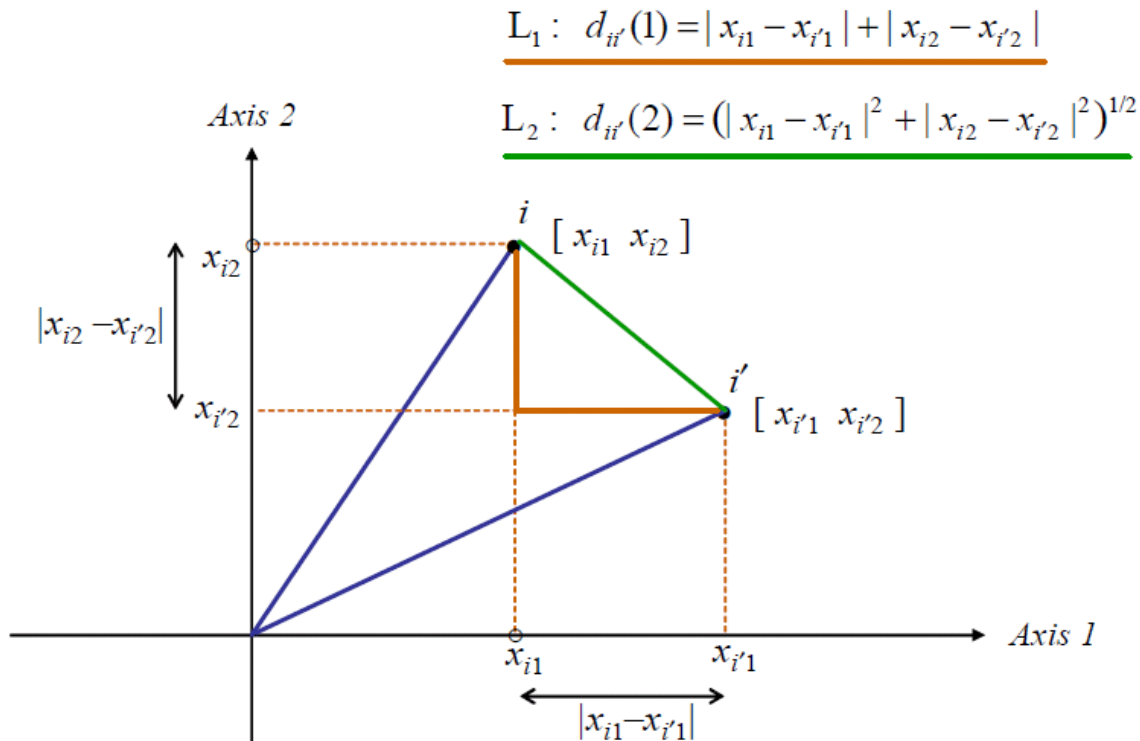
The length of the line segment connecting p and q

- Maximums-Norm (L_∞): $dist_\infty = \max\{|p_1 - q_1|, |p_2 - q_2|, \dots\}$

The distance between the least similar coordinates counts.

- Generalization of L_p -distance: $dist_p = (|p_1 - q_1|^p + |p_2 - q_2|^p + \dots)^{1/p}$

Feature spaces and distance functions



Source: <http://www.econ.upf.edu/~michael/stanford/maeb5.pdf>

- Attributes with large ranges outweigh ones with small ranges
 - e.g. income [10K-100K]; age [10-100]
- To balance the “contribution” of each attribute in the resulting distance, the attributes are scaled to fall within a small, specified range
- min-max normalization: to $[new_min_A, new_max_A]$

$$v' = \frac{v - min_A}{max_A - min_A} (new_max_A - new_min_A) + new_min_A$$

- e.g. normalize age=30 in [0-1], when min=10,max=100. $new_age = (30-10)/(100-10) = 2/9$
- z-score normalization

$$v' = \frac{v - mean_A}{stand_dev_A}$$

e.g. normalize 70,000 iff $\mu=50,000$, $\sigma=15,000$.
 $new_value = (70,000 - 50,000) / 15,000 = 1.33$

- Normalization by decimal scaling

$$v' = \frac{v}{10^j} \quad \begin{array}{l} j \text{ is the smallest integer} \\ \text{such that } \text{Max}(|v'|) < 1 \end{array}$$

e.g., if v lies in $[1, 999]$, if we set $j=3$ then v' will lie in $[0.001, 0.999]$

Dissimilarity between binary attributes

- A binary attribute has two states: 0 (absence), 1 (presence)
- A contingency table for binary data

		Object j		
		1	0	sum
Object i	1	q	r	$q + r$
	0	s	t	$s + t$
sum		$q + s$	$r + t$	p

- Simple matching coefficient
(for symmetric binary variables)

$$d(i, j) = \frac{r + s}{q + r + s + t}$$

- for asymmetric binary variables:

$$d(i, j) = \frac{r + s}{q + r + s}$$

- Jaccard coefficient

(for *asymmetric* binary variables)

$$sim_{Jaccard}(i, j) = \frac{q}{q + r + s}$$

Example

Name	Fever	Cough	Test-1	Test-2	Test-3	Test-4
Jack	1	0	1	0	0	0
Mary	1	0	1	0	1	0
Jim	1	1	0	0	0	0

$$d(jack, mary) = \frac{0 + 1}{2 + 0 + 1} = 0.33$$

$$d(jack, jim) = \frac{1 + 1}{1 + 1 + 1} = 0.67$$

$$d(jim, mary) = \frac{1 + 2}{1 + 1 + 2} = 0.75$$

Dissimilarity measures between categorical/ nominal attributes

- A categorical/ nominal attribute has >2 states (generalization of a binary attribute)
 - e.g. color={red, blue, green}
- Method 1: Simple matching
 - m: # of matches, p: total # of variables

$$d(i, j) = \frac{p - m}{p}$$

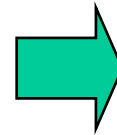
- Method 2: Map it to binary variables
 - create a new binary attribute for each of the M nominal states of the attribute

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Feature transformations for text data

- *Text represented as set of terms (“Bag-Of-Words” model)*
 - Term:
 - Single words (“cluster“, “analysis“..)
or
 - bigrams, trigrams, ...n-grams (“cluster analysis“..)
 - Transformation of a document D in a vector $r(D) = (h_1, ..., h_d)$
 $h_i \geq 0$: the frequency of term t_i in D

Schnee und Eis haben die Straßen in weiten Teilen Deutschlands in Rutschbahnen verwandelt. Lastwagen gerieten ins Schleudern, zahlreiche Fahrzeuge kamen an Steigungen nicht mehr vorwärts. Die Streudienste waren im Dauereinsatz...



...	...
Schnee	1
Eis	1
Fahrzeug	1
Politik	0
...	...

h_{Eis} points to the value 1 in the second column.

Feature transformations for text data

- Challenges in Text Mining:
 1. Common words (“e.g.”, “the”, “and”, “for”, “me”)
 2. Words with the same root (“fish”, “fisher”, “fishing”,...)
 3. Very high-dimensional space (dimensionality $d > 10.000$)
 4. Not all terms are equally important
 5. Most term frequencies $h_i = 0$ (“sparse feature space”)
- More challenges due to language:
 - Different words have same meaning (synonyms)
 - “freedom” – “liberty”
 - Words have more than one meanings
 - e.g. “java”, “mouse”

Feature transformations for text data

- Problem 1: Common words (“e.g.”, “the”, “and”, “for”, “me”)
 - Solution: ignore these terms (Stopwords)
- Problem 2: Words with the same root (“fish”, “fisher”, “fishing”,...)
 - Solution: Stemming

Map the words to their root

- "fishing", "fished", "fish", and "fisher" to the root word, "fish".

For English, the Porter stemmer is widely used.

(Porters Stemming Algorithms: <http://tartarus.org/~martin/PorterStemmer/index.html>)

Stemming solutions exist for other languages also.

The root of the words is the output of stemming.

Feature transformations for text data

- Problem 3: Too many features/ terms
 - Solution: Select the most important features (“Feature Selection”)
 - Example: average document frequency for a term
 - Very frequent items appear in almost all documents
 - Very rare terms appear in only a few documents

Ranking procedure:

1. Compute document frequency for all terms t_i :
2. Sort terms w.r.t. $DF(t_i)$ and get $rank(t_i)$
3. Sort terms by $score(t_i) = DF(t_i) \cdot rank(t_i)$
 e.g. $score(t_{23}) = 0.82 \cdot 1 = 0.82$
 $score(t_{17}) = 0.75 \cdot 2 = 1.5$
4. Select the k terms with the larger $score(t_i)$

$$DF(t_i) = \frac{|Dok_t_i|}{|ALL_Doks|}$$

Rank	Term	DF
1.	t_{23}	0.82
2.	t_{17}	0.65
3.	t_{14}	0.52
4.	...	...

Feature transformations for text data

- Problem 4: Not all terms are equally important
 - Idea: Very frequent terms are less informative than less frequent words. Define such a term weighting schema.
 - Solution: TF-IDF (Term Frequency · Inverse Document Frequency)
Consider both the importance of the term in the document and in the whole DB.

$$TF(t, d) = \frac{n(t, d)}{\sum_{w \in d} n(w, d)}$$

The frequency of term t in d

$$IDF(t) = \log\left(\frac{|DB|}{|\{d \mid d \in DB \wedge t \in d\}|}\right)$$

Inverse frequency of term t in all DB

$$TF \times IDF = TF(t, d)IDF(t)$$

Feature vector with TF IDF : $r(d) = (TF(t_1, d) \cdot IDF(t_1), \dots, TF(t_n, d) \cdot IDF(t_n))$

Feature transformations for text data

- Problem 5: for most of the terms $h_i = 0$
 - Euclidean distance is not a good idea: it is influenced by vectors lengths
 - Idea: use more appropriate distance measures

Jaccard Coefficient: Consider common terms

$$d_{Jaccard}(D_1, D_2) = 1 - \frac{|D_1 \cap D_2|}{|D_1 \cup D_2|}$$

Cosine Coefficient: Consider terms values (e.g. TFIDF)

$$d_{cosinus}(D_1, D_2) = 1 - \frac{\langle D_1, D_2 \rangle}{\|D_1\| \cdot \|D_2\|} = 1 - \frac{\sum_{i=0}^n (d_{1,i} \cdot d_{2,i})}{\sqrt{\sum_{i=0}^n d_{1,i}^2} \cdot \sqrt{\sum_{i=0}^n d_{2,i}^2}}$$

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Things you should know

- Feature types: binary, categorical/nominal, ordinal, numeric
- Basic univariate descriptors
- Basic bivariate descriptors
- Feature spaces/ metric spaces
- Distance functions for numeric data
- Distance functions for binary data
- Distance functions for categorical data
- Text transformation

- **Tutorial:** 1st tutorial on Thursday on:
 - basic data mining tasks
 - feature types
- **Homework:** Have a look at the tutorial in the website and try to solve the exercises.
 - Bring your questions
- **Suggested reading:**
 - Han J., Kamber M., Pei J. *Data Mining: Concepts and Techniques* 3rd ed., Morgan Kaufmann, 2011 (Chapter 2) (Section 7.2)
 - Tutorial on Text Mining and Link Analysis for Web and Semantic Web
 - Video: http://videolectures.net/ess07_grobelnik_twdml/
 - Slides are also there (Download slides)
 - Natural language processing course: <https://www.coursera.org/course/nlp>

Feature Transformation für räumliche Objekte (CAD-Daten, Proteine, ...)

– Invarianzen

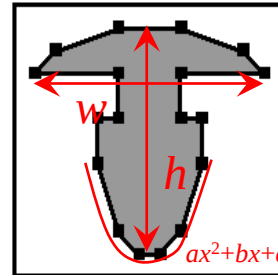
- Gleichheit (oder Ähnlichkeit) von Formen unabhängig von Lage und Orientierung im Raum
- Beispiele gleicher Formen im 2D und im 3D:



• Erwünscht:

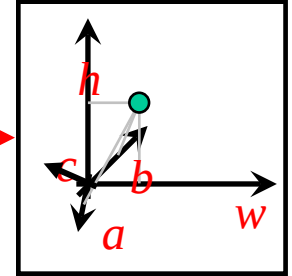
- Kanonische Darstellung, d.h. ohne Lage- und Orientierungsinformation
- Verallgemeinerung auf andere Objekteigenschaften

Objekt-Raum



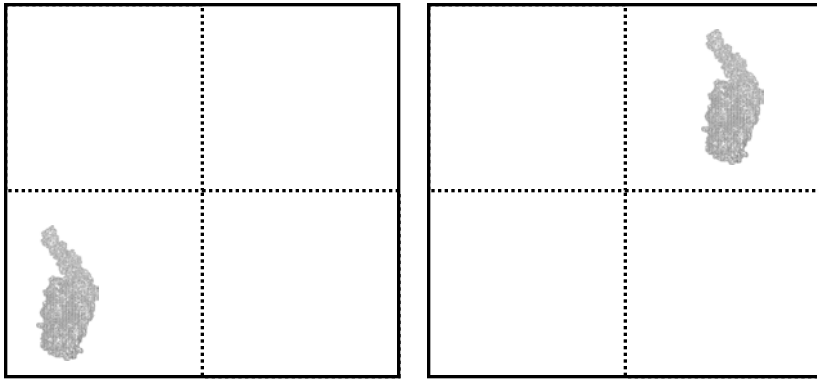
(h, w, a, b, c)

Merkmals-Raum

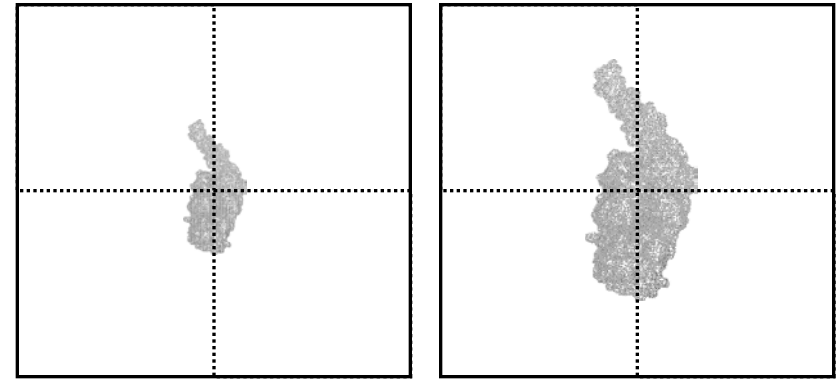


Die wichtigsten Invarianzen

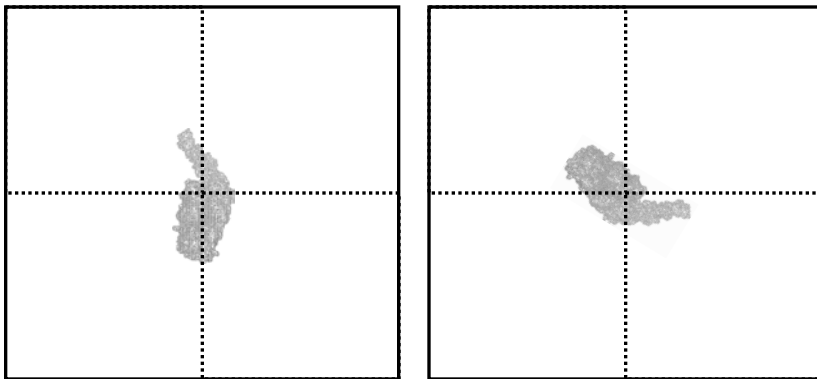
Translation



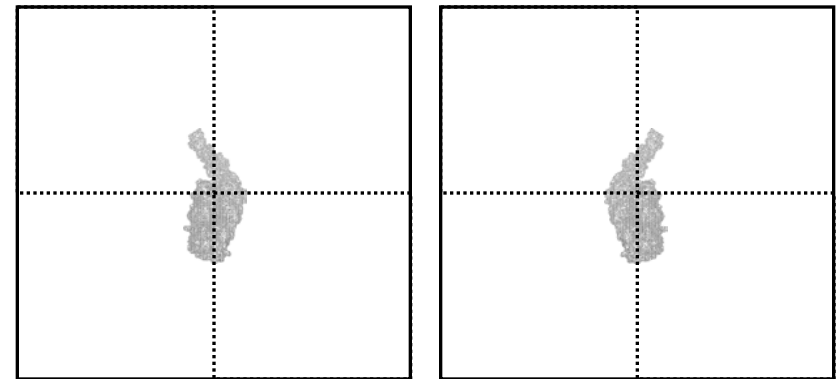
Skalierung



Rotation

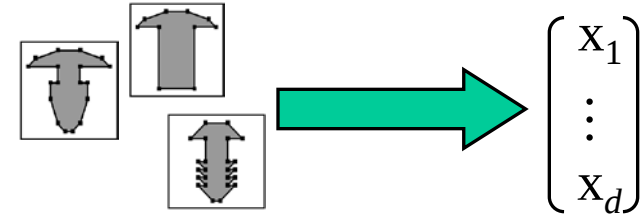


Spiegelung



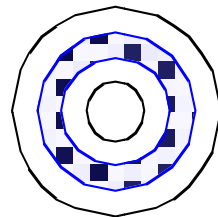
Volume Model [Ankerst, Kastenmüller, Kriegel, Seidl 99]

- Applikationen: CAD, Protein 3D-Strukturen
- Idee: *Formhistogramme* für 3D Objekte

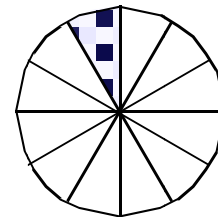


- Partitioniere den 3D-Raum in Zellen (Histogramm-Bins).
- Bestimme den Anteil an Punkten des Objektes pro Zelle (normiertes Histogramm).
- Durch die Normierung werden die Histogramme unabhängig von der Punktedichte.

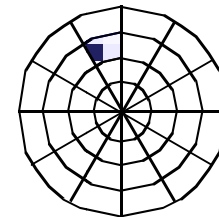
- Partitionierungen



Schalenmodell



Sektorenmodell

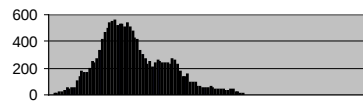


kombiniertes Modell

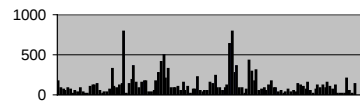
- Beispiel



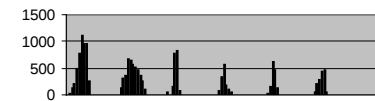
Seryl-tRNA
Synthetase



Schalenmodell
(120 Schalen)



Sektorenmodell
(122 Sektoren)

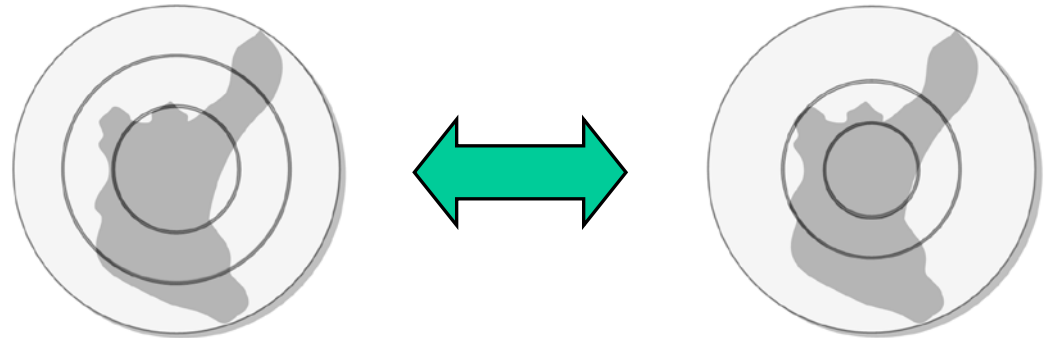


kombiniertes Modell
(20 Schalen, 6 Sektoren)

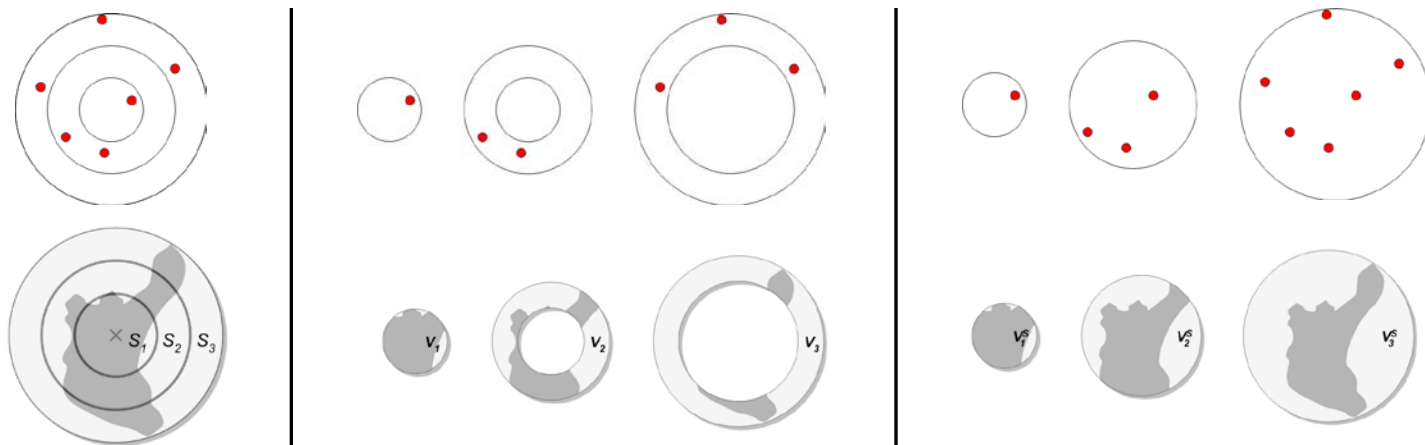
- Formale Definition der Histogramme
 - *Schalenmodell*: Definiere die Bins über den Abstand zum Mittelpunkt, d.h. Anzahl der Punkte auf der jeweiligen Schale.
 - *Sektorenmodell*: Anzahl der Punkte im jeweiligen Sektor.
 - *Kombiniertes Modell*: Synthese aus Schalen- und Sektorenmodell.
- Invarianzen
 - Translationsinvarianz durch Lagenormierung:
Verschiebung des Schwerpunkts eines Objektes in den Ursprung.
 - Rotationsinvarianz durch Hauptachsentransformation:
 - Drehung der Objekte, so dass die Hauptachsen auf den Koordinatenachsen liegen.
 - unnötig beim Schalenmodell, dieses ist inhärent rotationsinvariant.

- Verbesserung der Formhistogramme [Aßfalg, Kriegel, Kröger, Pötke 05]

- Proportionale Aufteilung



- Redundante Zuordnung zu den Bins



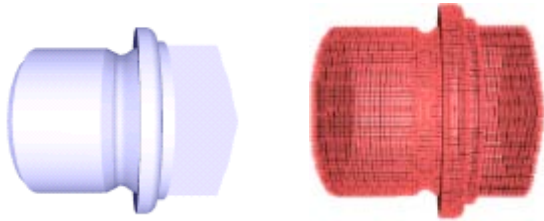
Objekt

Bisheriger Ansatz

Redundante Zuordnung

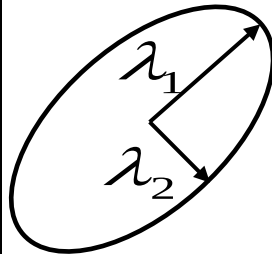
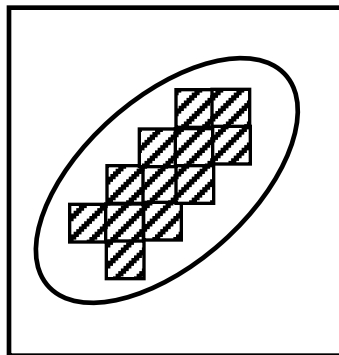
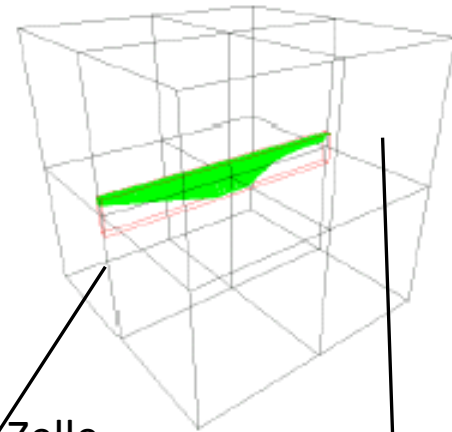
Eigenvalue Model [Kriegel, Kröger, Mashaël, Pfeifle, Pötke, Seidl 03]

- Volumen-Diskretisierung durch Voxel (3dimensionale Pixel)



- Würfelförmige Partitionierung der Bounding Box

- Bestimmung der Eigenwerte des Voxelinhaltes jeder Zelle



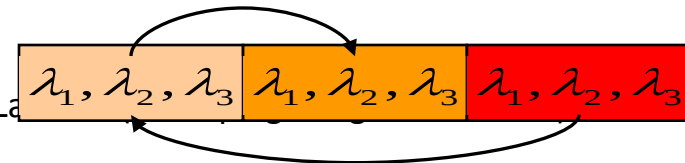
$\lambda_1, \lambda_2, \lambda_3$

.....

$\lambda_1, \lambda_2, \lambda_3$

Invarianzen

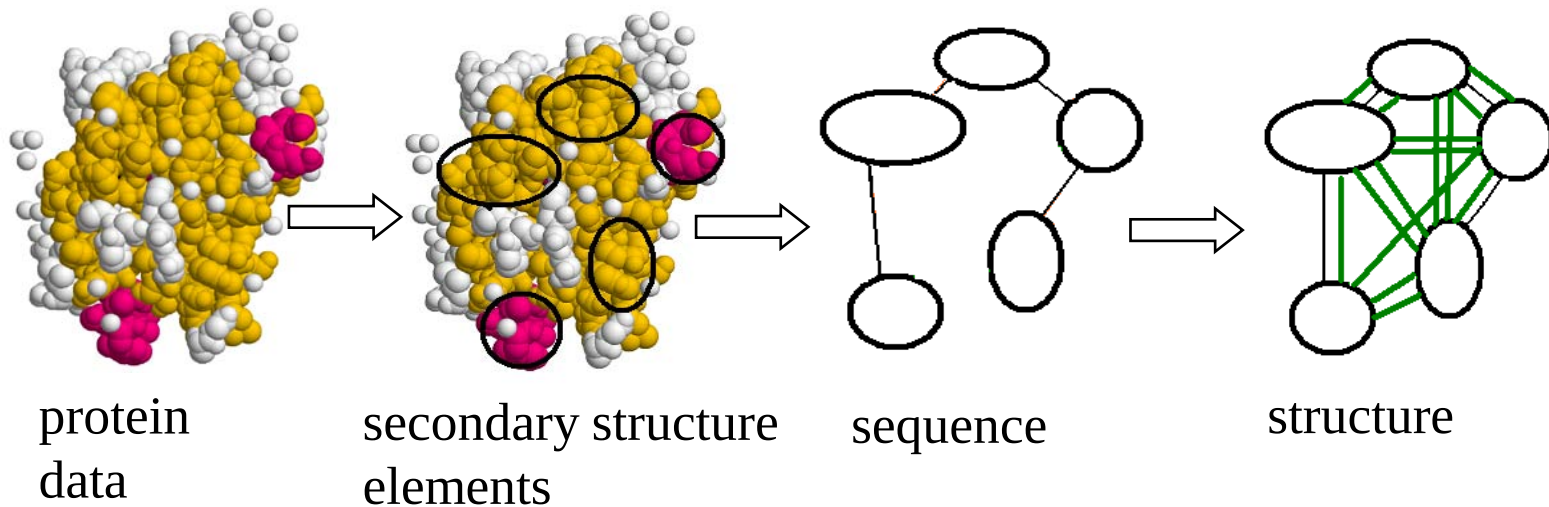
- Translationsinvarianz durch Lagenormierung:
Verschiebung des Schwerpunkts eines Objektes in den Ursprung.
- Skalierungsinvarianz durch Voxelisierung der Bounding Box/Bounding Cube des Objekts mit immer gleicher Voxelauflösung
- Rotationsinvarianz
 - Hauptachsentransformation (völlig rotationsinvariant, aber bei manchen Objekten sensitiv gegenüber kleinen Änderungen)
 - CAD Objekte oft in „vernünftiger“ Lage durch Konstrukteur abgespeichert, dann besser 90-Grad-Rotationsinvarianz: Zur Laufzeit werden die 24 Würfelpositionen durch Permutation der Merkmalsvektor-Elemente simuliert, die Distanz zweier Objekte ist das Minimum über 24 Distanzen
- Reflektionsinvarianz
 - Betrachte 48 statt 24 Permutationen zur La



Protein Datenbanken [Borgwardt, Ong, Schönauer, Vishwanathan, Smola, Kriegel 05]

Idee:

- Graphmodel für Protein 3D-Strukturen
- Knoten: Untereinheiten von Proteinen (secondary structure elements)
- Kanten: Nachbarschaft von Untereinheiten innerhalb der 3D-Struktur und entlang der Aminosäure Sequenz.

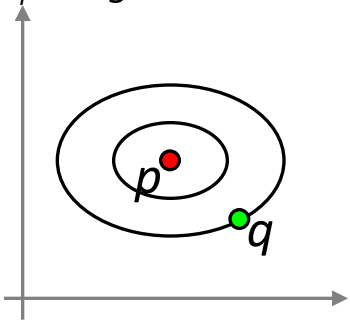


Feature spaces and distance functions

Weighted Euclidean Norm:

$$\text{dist} = (w_1(p_1 - q_1)^2 + w_2(p_2 - q_2)^2 + \dots)^{1/2}$$

w_i : weight in coordinate i



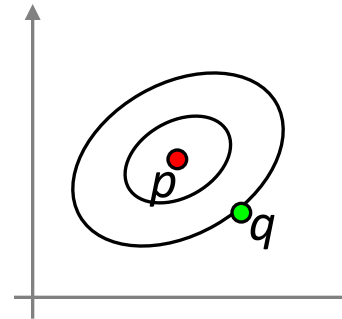
Häufig sind die Wertebereiche der Merkmale deutlich unterschiedlich.

Example: feature $M_1 \hat{I}[0.01 \dots 0.05]$
feature $M_2 \hat{I}[3.1 \dots 22.2]$

Since M_1 überhaupt berücksichtigt wird, should be weighted higher

Quadratic form:

$$\text{dist} = ((p - q) \mathbf{M} (p - q)^T)^{1/2}$$



In the previous case, the features are separately weighted.

Besonders bei Farbhistogrammen müssen auch verschiedene Merkmale gemeinsam gewichtet werden.

Instead of distance measures, which measure the distance between two objects, we can use sometimes similarity measures

Descriptors for feature vectors

Descriptors for feature vectors

- Consider a set of feature vectors/ objects DB
- Centroid (**vgl.** Arithmetic mean):

$$\mu_{DB} = \frac{1}{DB} \cdot \sum_{o \in DB} o$$

- **Achtung: bei allgem. Metrischen Räumen muss Centroid nicht notwendigerweise existieren!!!**

- Medoid m_{DB} :
 - The most centrally located object in DB
 - **Bei allgem. Metrischen Räumen:** The one with the smallest average distance to all the other objects in DB

- Variance:
$$Var_{DB} = \frac{1}{DB} \cdot \sum_{o \in DB} dist(o, \mu_{DB})$$

- Standard deviation: square root of variance

Feature spaces and distance functions

Hauptachsenanalyse eine Menge DB von *Euklidischen Vektoren*

- Covariance matrix:
$$\Sigma_{DB} = \frac{1}{|DB|} \sum_{o \in DB} (o - \mu_{DB})(o - \mu_{DB})^T$$
- The matrix is decomposed in:
 - an orthonormal matrix $V = [e_1, \dots, e_d]$ (Eigen vectors)
 - and a diagonal matrix $\Lambda = \text{diag}(\lambda_1, \dots, \lambda_d)$ (Eigen values)
 - so dass gilt: $\Sigma_{DB} = V \Lambda V^T$
- Interpretation:
 - Eigenvektoren:
Hauptausrichtung der Datenpunkte in DB
 - Eigenwerte:
Varianz der Datenpunkte in DB entlang der entspr. Eigenvektoren

