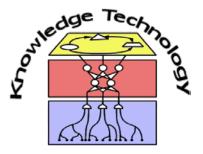
Data Mining

Lecture 3 Preprocessing Methods



http://www.informatik.uni-hamburg.de/WTM/

Data Preprocessing

Data Preprocessing: An Overview



- Data Quality
- Major Tasks in Data Preprocessing
- Data Cleaning
- **Data Integration**
- Data Reduction
- **Data Transformation**
- Data Discretization

Data Quality: why preprocess the Data?

Measures for data quality: A multidimensional view

- Accuracy: correct or wrong, accurate or not
- Completeness: not recorded, unavailable, ...
- Consistency: some modified but some not, dangling, ...
- Timeliness: timely update?
- Believability: how trustable are the data are?
- Interpretability: how easily the data can be understood?

Why We should Clean Dirty Data



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

Data reduction

- Dimensionality reduction
- Data compression
- Data transformation and data discretization
 - Normalization
 - Concept hierarchy generation

Data Preprocessing

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Data Cleaning

- Data in the real world is "dirty" or incorrect, e.g., instrument faulty, human or computer error, transmission error
- Incomplete: lacking attribute values, lacking certain attributes of interest, or only aggregate data available
 - e.g., Occupation=" " (missing data)
- Noisy: containing noise, errors, or outliers
 - e.g., Salary="-10" (an error)
- Inconsistent: containing discrepancies in codes or names, e.g.,
 - Age="42", Birthday="03/07/2012"
 - Was rating "1, 2, 3", now rating "A, B, C"
 - discrepancy between duplicate records
- Intentionally imprecise (e.g., disguised missing data)
 - Jan. 1 as everyone's birthday?

Incomplete (Missing) Data

- Data is not always available
 - E.g., many tuples have no recorded value for several attributes, such as customer income in sales data
- Missing data may be due to
 - equipment malfunction
 - inconsistent with other recorded data and thus deleted
 - data not entered due to misunderstanding
 - certain data may not be considered important at the time of entry
 - not register history or changes of the data
- Missing data may need to be inferred

How to handle missing Data?

- Ignore the tuple: usually done when class label is missing (when doing classification) — not effective when the % of missing values per attribute varies considerably
- Fill in the missing value manually: tedious + infeasible?
- Fill it in automatically with
 - a global constant : e.g., "unknown", a new class?!
 - the attribute mean
 - the attribute mean for all samples belonging to the same class: smarter
 - the most probable value: inference-based such as Bayesian formula or decision tree

Missing Data

 One possible interpretation of missing values – "don't care" values:

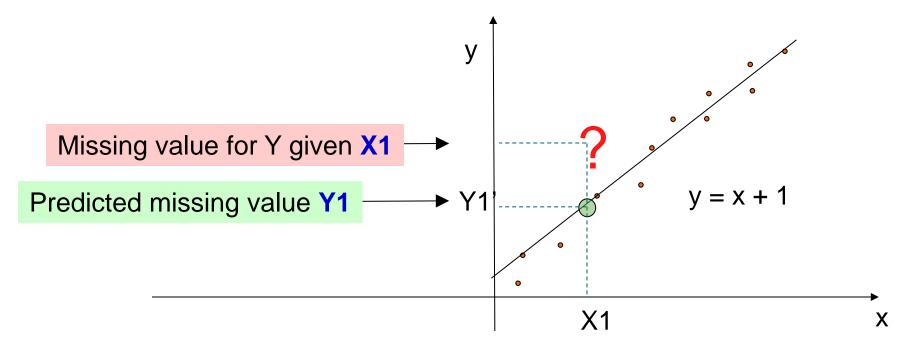
```
X = \{1, ?, 3\}

\rightarrow for the second feature the domain is [0, 1, 2, 3, 4]:

X1 = \{1, 0, 3\}, X2 = \{1, 1, 3\}, X3 = \{1, 2, 3\}, X4 = \{1, 3, 3\}, X5 = \{1, 4, 3\}
```

- Data miner can generate model of correlation between features.
 - Different techniques possible: regression, Bayesian formalism, clustering, or decision tree induction.

Missing Data Replacement with Regression Analysis



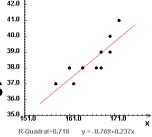
- In general, replacement of missing values using a simple, artificial schema of data preparation is speculative and often misleading.
- It is best to generate multiple solutions of data mining with and without features that have missing values, and then make comparison, analysis and interpretation.

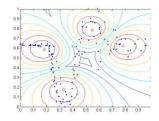
Noisy Data

- Noise: random error or variance in a measured variable
- Incorrect attribute values may be due to
 - technology limitation
 - faulty data collection instruments
 - data entry problems
 - data transmission problems
 - inconsistency in naming convention
- Other data problems which require data cleaning
 - duplicate records
 - incomplete data
 - inconsistent data

How to handle noisy Data?

- Binning
 - first sort data and partition into (equal-frequency) bins
 - then one can smooth by bin means, smooth by bin median, smooth by bin boundaries, etc.
- Regression
 - smooth by fitting the data into regression functions 37.0
- Clustering
 - detect and remove outliers
- Combined computer and human inspection
 - detect suspicious values and check by human (e.g., deal with possible outliers)





Data Quality: why preprocess the Data?



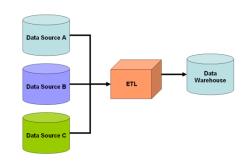
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Data Integration



- Data integration:
 - Combines data from multiple sources into a coherent store
- Schema integration: e.g., A.cust-id ≡ B.cust-#
 - Integrate metadata from different sources
- Entity identification problem:
 - Identify real world entities from multiple data sources, e.g.,
 Bill Clinton = William Clinton
- Detecting and resolving data value conflicts
 - For the same real world entity, attribute values from different sources are different
 - Possible reasons: different representations, different scales, e.g., metric vs. British units

Handling Redundancy in Data Integration

- Redundant data occur often when integrating multiple databases
 - Object identification: The same attribute or object may have different names in different databases
 - Derivable data: One attribute may be a "derived" attribute in another table, e.g., annual revenue
- Redundant attributes may be possible to detect by correlation analysis
- Careful integration of the data from multiple sources may reduce/avoid redundancies and inconsistencies and improve mining speed and quality

Correlation Analysis (Nominal Data)

X² (chi-square) test

$$\chi^2 = \sum \frac{(Observed - Expected)^2}{Expected}$$

- The cells that contribute the most to the X² value are those whose actual count is very different from the expected count
- Correlation does not imply causality
 - # of hospitals and # of car-thefts in a city are correlated
 - Both are causally linked to the third variable: population

Chi-Square Calculation: an Example

• Questionnaire among *N*=1500 participants:

	Play chess	Not play chess	Sum (row)
Like science fiction	250	250	500
Not like science fiction	50	950	1000
Sum (column)	300	1200	1500

• Expected results e_{ij} from the null hypothesis stating that "preferred reading" and "game favour" are uncorrelated):

	Play chess	Not play chess	Sum (row)
Like science fiction	100	400	500
Not like science fiction	200	800	1000
Sum (column)	300	1200	1500

$$e_{ij} = sum(col\ i) \cdot sum(row\ j) / N$$

Chi-Square Calculation: an Example

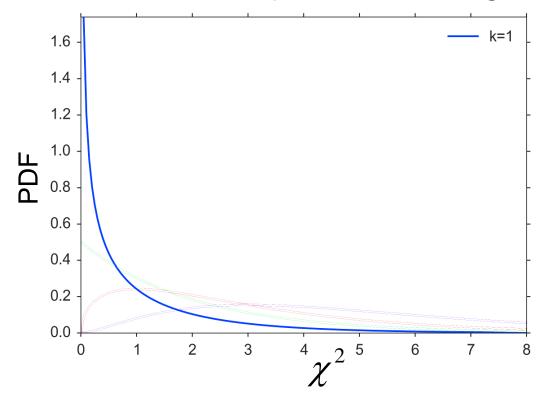
X² (chi-square) calculation:

$$\chi^2 = \frac{(250 - 100)^2}{100} + \frac{(50 - 200)^2}{200} + \frac{(250 - 400)^2}{400} + \frac{(950 - 800)^2}{800} = \underline{421.9}$$

What does it show?

Chi-square Test

Small deviations are more expected than large deviations:

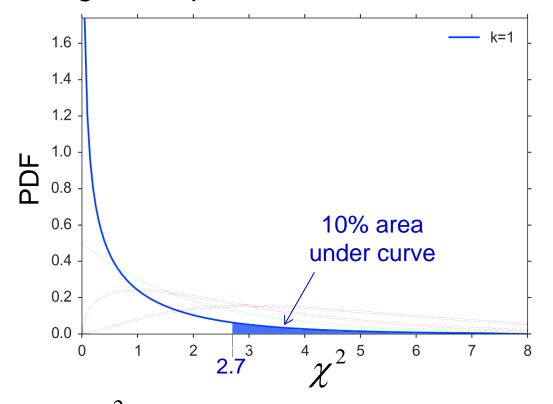


- Random variable Y has a normal distribution
 - \rightarrow Y² has a chi-square distribution (with 1 degree of freedom)

Chi-square Test

Some percentage of expected deviations is over a

threshold



- E.g.: 10% of all χ^2 values are larger than critical value 2.7
- Values can be looked up in a chi-square distribution table

Chi-square Distribution Table

probability level

α	0.5	0.1	0.05	0.02	0.01	0.001
	0.455	2.706	3.841	5.412	6.635	10.827

- α=0.1: 90% of values are below critical value 2.706
- α=0.05: 95% of values are below critical value 3.841

. . .

- α: significance level
- Earlier, we have found a value of 421.9
- → Our data are extremely unlikely given the null hypothesis

Chi-Square Calculation: an Example

X² (chi-square) calculation:

$$\chi^2 = \frac{(250 - 100)^2}{100} + \frac{(50 - 200)^2}{200} + \frac{(250 - 400)^2}{400} + \frac{(950 - 800)^2}{800} = \underline{421.9}$$

- It shows that "preferred reading" and "game favour" are correlated in the group (since X² larger than 10.827, from X² table – a statistical measure for significance of 2x2 table)
- What if all numbers were 10x smaller (N=150 participants)?

$$\rightarrow \chi^2 = ... = 42.19$$

What if all numbers were 50x smaller (N=30 participants)?

$$\chi^{2} = \frac{(5-2)^{2}}{2} + \frac{(1-4)^{2}}{4} + \frac{(5-8)^{2}}{8} + \frac{(19-16)^{2}}{16} = 8.4375$$

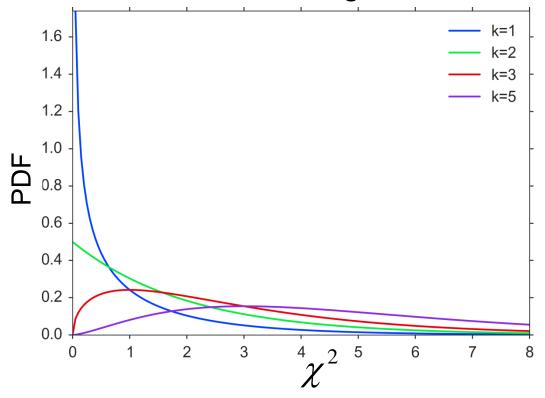
Chi-Square Calculation: Degrees of Freedom

		Category 1 Levels			Sum	
		L1	L2	•••	LJ	(row)
Category 2 Levels	L1					
Leveis						
	LI					
Sum	(col.)					N

- If the two categories have several levels (J levels for category 1 and I levels for category 2), then there are more degrees of freedom in which the entries can differ
- I x J contingency table
- Number of degrees of freedom: (I-1) x (J-1)

Chi-Square Calculation: Degrees of Freedom

More degrees of freedom make larger deviations probable:



- Sum of k independ. random variables Y_i with normal distrib.
 - $\rightarrow \Sigma_i Y_i^2$ has a χ^2 distribution with k degrees of freedom

Chi-square Distribution Table

probability level

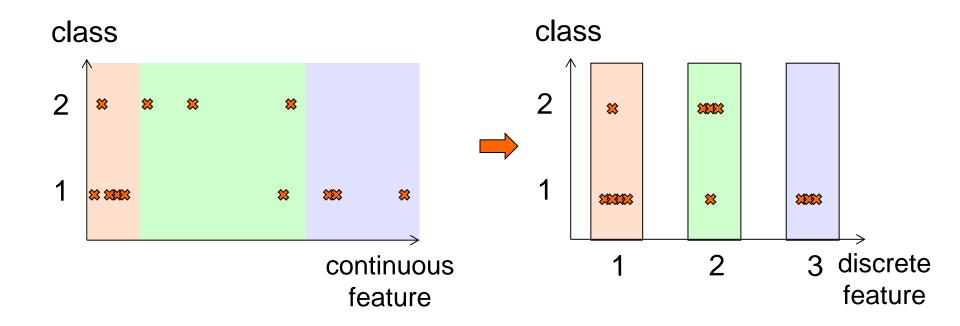
DF	0.5	0.1	0.05	0.02	0.01	0.001
1	0.455	2.706	3.841	5.412	6.635	10.827
2	1.386	4.605	5.991	7.824	9.210	13.815
3	2.366	6.251	7.815	9.837	11.345	16.268
4	3.357	7.779	9.488	11.668	13.277	18.465
5	4.351	9.236	11.070	13.388	15.086	20.517



Degrees of freedom

Values Reduction

Data often have continuous features



Many classification algorithms require data with discrete attributes

- 1. Sort the data for the given feature in ascending order
- 2. Define initial intervals so that every value of the feature is in a separate interval

3. Repeat.

- 3.1 Compute X^2 tests for each pair of adjacent intervals
- 3.2 Merge two adjacent intervals with the lowest X^2 value, if calculated X^2 is less than threshold
- **Until** no X^2 test of any two adjacent intervals is less than threshold value

Values Reduction – Contingency Table

• A ChiMerge requires computation of X^2 test for the contingency table 2 x 2 of categorical data:

	Class 1	Class 2	Σ
Interval-1	A ₁₁	A ₁₂	R_1
Interval-2	A ₂₁	A ₂₂	R ₂
Σ	C ₁	C_2	N

 X^2 test is:

$$\chi^{2} = \sum_{i=1}^{2} \sum_{j=1}^{k} \frac{\left(A_{ij} - E_{ij}\right)^{2}}{E_{ij}}$$

where:

k = number of classes,

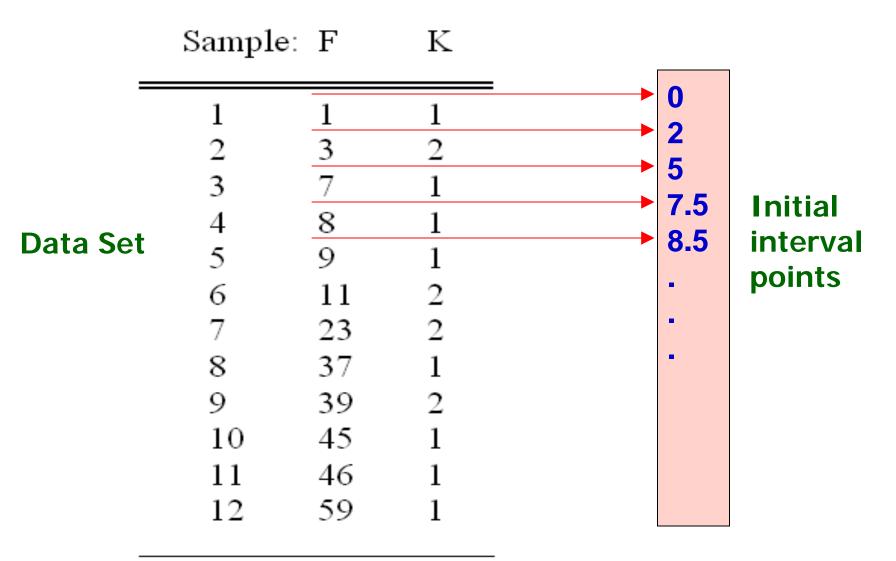
A_{ii} = number of instances in the i-th interval, j-th class,

 $E_{ij} = expected frequency of A_{ij}$, which is computed as $(R_i \cdot C_j) / N$,

 R_i = number of instances in the i-th interval = $\sum A_{ij}$, j = 1,...k,

 C_{j} = number of instances in the j-th class = $\sum A_{ij}$, i = 1,2,

N = total number of instances = $\sum R_i$, i = 1,2.



• X^2 was minimum for intervals: [7.5,8.5] and [8.5,10]

Sample:	F	K
1	1	1
2	3	2
3	7	1
4	8	1
5	9	1
6	11	2 2
7	23	2
8	37	1
9	39	2
10	45	1
11	46	1
12	59	1

	Class 1	Class 2	Σ
Interval [7.5,8.5]	A ₁₁ =1	A ₁₂ =0	$R_1=1$
Interval [8.5,10]	A ₂₁ =1	A ₂₂ =0	R ₂ =1
Σ	C ₁ =2	C ₂ =0	N=2

Based on the table's values, we can calculate expected values:

E11 =
$$1*2/2 = 1$$
, E12 = $1*0/2 = 0$, E21 = $1*2/2 = 1$, & E22 = $1*0/2 = 0$

and corresponding X^2 test:

$$X^2 = (1-1)^2/1 + (0-0)^2/0 + (1-1)^2/1 + (0-0)^2/0 = \mathbf{0}$$

For d=1 degree of freedom: $X^2 = 0 < 2.706 \rightarrow merge!$ (a=0.1)

... one of the following iterations:

Sample:	F	K
1	1	1
2	3	2
2 3 4 5 6 7	7	1
4	8 9	1
5	9	1
6	11	2
	23	2
8	37	1
9	39	2
10	45	1
11	46	1
12	59	1

	Class 1	Class 2	Σ
Interval [0.0 , 7.5]	A ₁₁ =2	A ₁₂ =1	R ₁ =3
Interval [7.5,10]	A ₂₁ =2	A ₂₂ =0	R ₂ =2
Σ	C ₁ =4	C ₂ =1	N=5

E11 =
$$3*4/5$$
 = 2.4, E12 = $3*1/5$ = 0.6,
E21 = $2*4/5$ = 1.6, E22 = $2*1/5$ = 0.4
 $X^2 = (2-2.4)^2/2.4 + (1-0.6)^2/0.6 + (2-1.6)^2/1.6 + (0-0.4)^2/0.4 =$ **0.834**

For d=1 degree of freedom:
$$X^2 = 0.834 < 2.706 \rightarrow merge!$$
(a=0.1)

... One of the additional iterations:

Sample:	F	K
1	1	1
2	3	2
3	7	1
4	8	1
5	9	1
6	11	2 2
7	23	2
8	37	1
9	39	2
10	45	1
11	46	1
12	59	1

	Class 1	Class 2	Σ
Interval [0.0,10]	A ₁₁ =4	A ₁₂ =1	R ₁ =5
Interval [10 ,42]	A ₂₁ =1	A ₂₂ =3	R ₂ =4
Σ	C ₁ =5	C ₂ =4	N=9

$$E11 = 2.78$$
, $E12 = 2.22$, $E21 = 2.22$, $E22 = 1.78$

$$X^2 = 2.72 > 2.706 \longrightarrow NO merge!$$

Final discretization:

Interval representatives: 5 (low) 26 (medium) 51 (high)

Sample: F K			Sampl	ample: F		
1	1	1	=	1	5	1
2	3	2	with reduced	2	5	2
3	7	1	set of values F:	3	5	1
4	8	1		4	5	1
5	9	1		5	5	1
6	11	2	Original set	6	26	2
7	23	2		7	26	2
8	37	1		8	26	1
9	39	2		9	26	2
10	45	1		10	51	1
11	46	1		11	51	1
12	59	1		12	51	1

Data Preprocessing

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- Data Reduction



- Data Transformation
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Feature Reduction

Which features to select, and how?

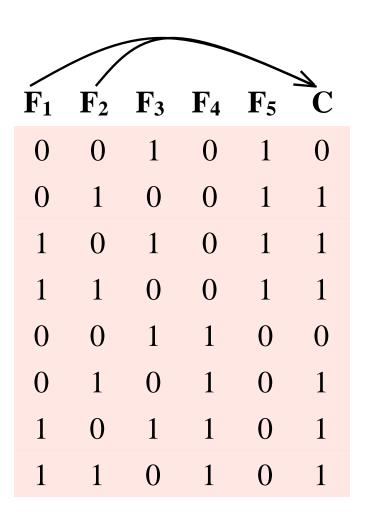
TRS_DT	TRS_TYP_CD	REF_DT	REF_NUM	CO_CD	GDS_CD	QTY	UT_CD	UT_PRIC
21/05/93	00001	04/05/93	25119	10002J	001M	10	CTN	22.000
21/05/93	00001	05/05/93	25124	10002J	032J	200	DOZ	1.370
21/05/93	00001	05/05/93	25124	10002J	033Q	500	DOZ	1.000
21/05/93	00001	13/05/93	25217	10002J	024K	5	CTN	21.000
21/05/93	00001	13/05/93	25216	10026H	006C	20	CTN	69.000
21/05/93	00001	13/05/93	25216	10026H	008Q	10	CTN	114.000
21/05/93	00001	14/05/93	25232	10026H	006C	10	CTN	69.000
21/05/93	00001	14/05/93	25235	10027E	003A	5	CTN	24.000
21/05/93	00001	14/05/93	25235	10027E	001M	5	CTN	24.000
21/05/93	00001	22/04/93	24974	10035E	009F	50	CTN	118.000
21/05/93	00001	27/04/93	25033	10035E	015A	375	GRS	72.000
21/05/93	00001	20/05/93	25313	10041Q	010F	10	CTN	26.000
21/05/93	00001	12/05/93	25197	10054R	002E	25	CTN	24.000

Features Reduction

Two standard approaches:

- Feature selection: A process that chooses an optimal subset of features according to an objective function:
 - feature ranking algorithms,
 - minimum subset algorithms.
- Feature extraction: refers to the mapping of the original high-dimensional data onto a lower-dimensional space.
 - Descriptive setting: minimizes the information loss
 - Predictive setting: maximizes the class discrimination

Feature selection – Example for Optimal Features' Subset



- Data set (whole set)
 - Five Boolean features
 - $C = F_1 \vee F_2$
 - $F_3 = \neg F_2$, $F_5 = \neg F_4$
 - Optimal subset:

$$\{F_1, F_2\}$$
 or $\{F_1, F_3\}$

 Combinatorial nature of searching for an optimal subset

Feature Selection – Complexity

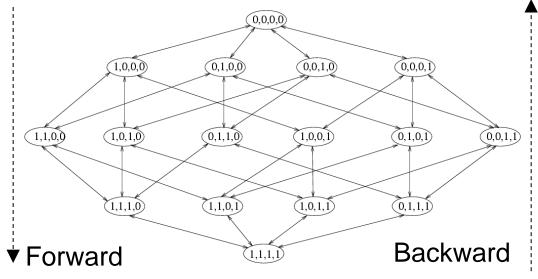
Feature selection in general can be viewed as a search problem.

 For practical methods, an optimal search is not feasible, and simplifications are made to produce acceptable and timely reasonable results:

heuristic criteria

bottom-up approach

top-down approach



Methods of Feature Selection

- Univariate methods
 - Considers one variable (feature) at a time.
- Filter methods
 - Separating feature selection from classifier learning
 - Relying on general characteristics of data (information, distance, dependence, consistency)
 - No bias toward any learning algorithm, fast
- Wrapper methods
 - Relying on a predetermined classification algorithm.
 - Using predictive accuracy as goodness measure
 - High accuracy, computationally expensive
- Embedded methods
 - Combine Filter and Wrapper approaches

Features Selection: Univariate Methods

Comparison of means and variances:

Samples of two classes (A and B) can be examined:

$$SE(A-B) = \sqrt{\frac{var(A)}{n_1} + \frac{var(B)}{n_2}}$$

TEST:

$$\frac{\left| \operatorname{mean}(A) - \operatorname{mean}(B) \right|}{\operatorname{SE}(A - B)} > threshold-value$$

where n_1 and n_2 are the corresponding number of samples for classes A and B.

Features Selection: Univariate Methods

Comparison of means and variances – Example:

X	Y	С
0.3 0.2 0.6 0.5 0.7 0.4	0.7 0.9 0.6 0.5 0.7 0.9	A B A A B

Threshold value is 0.5

$$X_A = \{0.3, 0.6, 0.5\},\$$

$$Y_A = \{0.7, 0.6, 0.5\},\$$

$$X_{B} = \{0.2, 0.7, 0.4\},\$$

$$Y_B = \{0.9, 0.7, 0.9\}$$

Features Selection: Univariate Methods

Comparison of means and variances – Example:

$$SE(X_A - X_B) = \sqrt{\frac{\text{var}(X_A)}{n_1} + \frac{\text{var}(X_B)}{n_2}} = \sqrt{\frac{0.0233}{3} + \frac{0.6333}{3}} = 0.4678$$

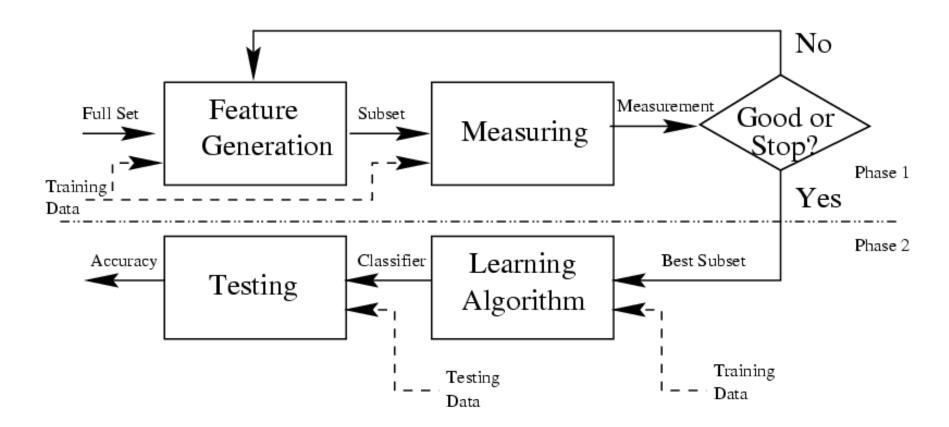
$$SE(Y_A - Y_B) = \sqrt{\frac{\text{var}(Y_A)}{n_1} + \frac{\text{var}(Y_B)}{n_2}} = \sqrt{\frac{0.0133}{3} + \frac{0.0133}{3}} = 0.0875$$

Tests:

X:
$$\frac{\left| \operatorname{mean}(A) - \operatorname{mean}(B) \right|}{\operatorname{SE}(A - B)} = \frac{\left| 0.4667 - 0.4333 \right|}{0.4678} < 0.5$$
Y:
$$\frac{\left| \operatorname{mean}(A) - \operatorname{mean}(B) \right|}{\operatorname{SE}(A - B)} = \frac{\left| 0.6 - 0.8333 \right|}{0.0875} > 0.5$$

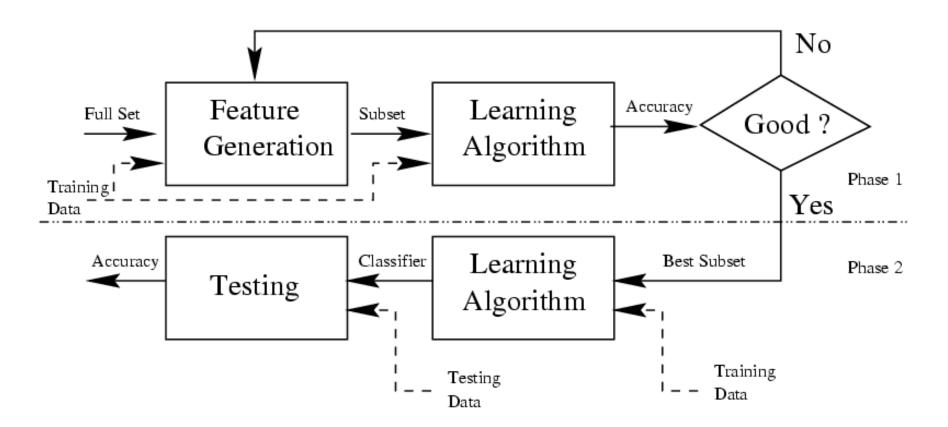
X is a candidate feature for reduction because its mean values are close, and therefore the final test is below threshold value.

Filter Model



- Example filter algorithm for feature selection:
 - Relief (Kira & Rendell 1992)

Wrapper Model



- Example wrapper algorithm for Feature Selection:
 - SVM

Data Reduction: Dimensionality Reduction

Curse of dimensionality

- When dimensionality increases, data becomes increasingly sparse
- Density and distance between points, which is critical to clustering, outlier analysis, becomes less meaningful
- The possible combinations of subspaces will grow exponentially

Dimensionality reduction

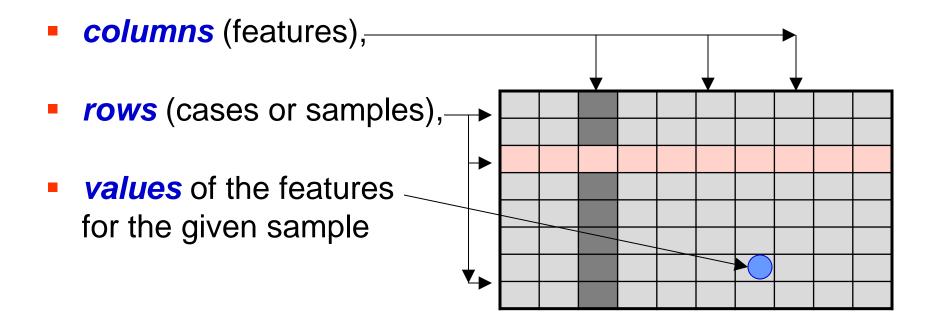
- Avoid the curse of dimensionality
- Help eliminate irrelevant features and reduce noise
- Reduce time and space required in data mining
- Allow easier visualization

Data Reduction Strategies

- Why data reduction? A database/data warehouse may store terabytes of data. Complex data analysis may take a very long time to run on the complete data set.
- Obtain a reduced representation of the data set that is much smaller in volume but yet produces the (almost) same analytical results
- Data reduction strategies
 - Dimensionality reduction, e.g., remove unimportant attributes
 - Feature subset selection, feature creation
 - Wavelet transforms
 - Principal Components Analysis (PCA)
 - Numerosity reduction (some simply call it: Data Reduction)
 - Regression and Log-Linear Models
 - Histograms, clustering, sampling
 - Data cube aggregation
 - Data compression

Dimensions Reduction of Large Data Sets

Main dimensions:



Data Preprocessing

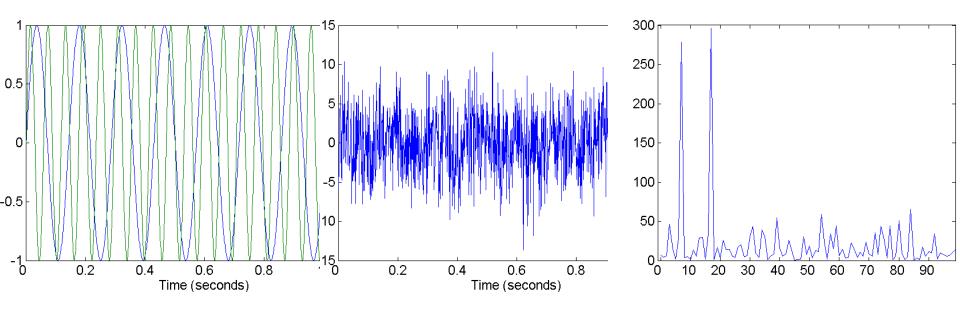
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Data Discretization

Mapping Data to a New Space

- Fourier transform: mapping from time to frequency domain
- Wavelet transform



Two Sine Waves

Two Sine Waves + Noise

Frequencies

Data Transformation

- A function that maps the entire set of values of a given attribute to a new set of replacement values so that each old value can be identified with one of the new values
- Methods
 - Smoothing: Remove noise from data
 - Attribute/feature construction
 - New attributes constructed from the given ones
 - Aggregation: Summarization
 - Normalization: Scaled to fall within a smaller, specified range
 - min-max normalization
 - z-score normalization
 - normalization by decimal scaling
 - Discretization: Concept hierarchy climbing

Normalization

Min-max normalization: to [new_min_A, new_max_A]

$$v' = \frac{v - min_A}{max_A - min_A} (new _ max_A - new _ min_A) + new _ min_A$$

- Ex. Let income range \$12,000 to \$98,000. Normalize to [0.0, 1.0]. Then \$73,600 is mapped to $\frac{73,600-12,000}{98,000-12,000}(1.0-0)+0=0.716$
- **Z-score normalization** (μ: mean, σ: standard deviation):

$$v' = \frac{v - \mu_A}{\sigma_A}$$

- Ex. Let $\mu = 54,000$, $\sigma = 16,000$. Then $\frac{73,600-54,000}{16,000} = 1.225$
- Normalization by decimal scaling

$$v' = \frac{v}{10^{j}}$$
 where j is the smallest integer such that Max(|v'|) < 1

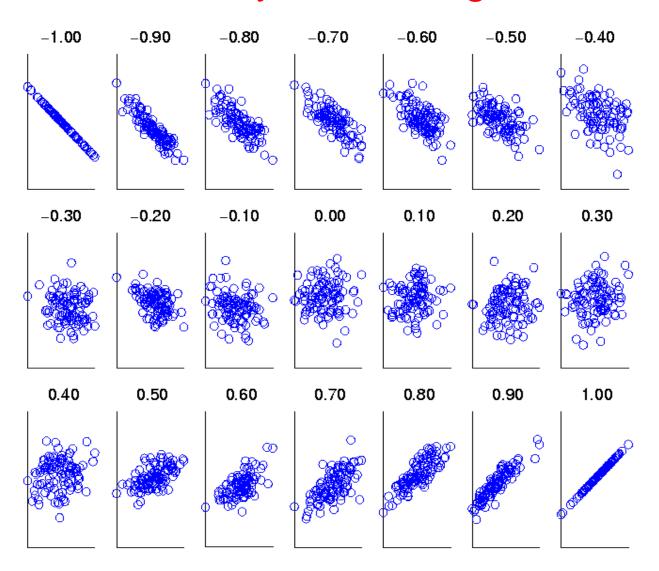
Correlation Analysis (Numeric Data)

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

$$r_{A,B} = \frac{\sum_{i=1}^{N} (a_i - \overline{A})(b_i - \overline{B})}{N\sigma_A \sigma_B}$$

- N = number of tuples
- \overline{A} and \overline{B} = means of attributes A and B
- σ_A and σ_B = standard deviations of A and B
- $r_{A,B} > 0$: A and B are positively correlated
 - A's values increase as B's. Larger $r_{AB} \rightarrow$ stronger correlation.
- $r_{A,B} = 0$: independent
- $r_{AB} < 0$: negatively correlated

Visually Evaluating Correlation



Scatter plots

Correlation coefficients range from –1 to 1.

Covariance (Numeric Data)

Covariance:

$$Cov(A, B) = E((A - \bar{A})(B - \bar{B})) = \frac{\sum_{i=1}^{n} (a_i - \bar{A})(b_i - \bar{B})}{n}$$

Related to correlation coefficient: $r_{A,B} = \frac{Cov(A,B)}{\sigma_A \sigma_B}$

- n = number of tuples
- \overline{A} and \overline{B} = mean or **expected values** of A and B
- σ_A and σ_B = standard deviation of A and B.
- **Positive covariance**: If $Cov_{A,B} > 0$, then A and B both tend to be larger than their expected values.
- **Negative covariance**: If $Cov_{A,B} < 0$ then if A is larger than its expected value, B is likely to be smaller than its expected value.
- Independence: $Cov_{A,B} = 0$

Co-Variance: an Example

$$Cov(A, B) = E((A - \bar{A})(B - \bar{B})) = \frac{\sum_{i=1}^{n} (a_i - \bar{A})(b_i - \bar{B})}{n}$$

It can be simplified in computation as

$$Cov(A, B) = E(A \cdot B) - \bar{A}\bar{B}$$

- Suppose two stocks A and B have the following values in one week: (2, 5), (3, 8), (5, 10), (4, 11), (6, 14).
- Question: If the stocks are affected by the same industry trends, will their prices rise or fall together?
 - E(A) = (2 + 3 + 5 + 4 + 6)/5 = 20/5 = 4
 - E(B) = (5 + 8 + 10 + 11 + 14)/5 = 48/5 = 9.6
 - Cov(A,B) = (2.5 + 3.8 + 5.10 + 4.11 + 6.14)/5 4.9.6 = 4
- Thus, A and B rise together since Cov(A, B) > 0.

Principal Component Analysis (Steps)

- Given N data vectors from n-dimensions, find m << n orthogonal vectors (principal components) to represent the data
 - Subtract mean from the input data, each attribute has mean zero
 - Compute *m* orthonormal (unit) vectors, i.e., *principal components*
 - Each input data (vector) is a linear combination of the m principal component vectors
 - The principal components are sorted in order of decreasing "significance" (1st component: data has maximum variance)
 - Since the components are sorted, the size of the data can be reduced by eliminating the insignificant components, i.e., those with low variance
 - Thus, using only the most significant principal components, it is possible to reconstruct a good approximation of the original data.
- Works for numeric data; reduction of higher dimensions to lower

Principal Components Analysis Algorithm

1. Compute the *n×n* covariance matrix S

$$S_{ij} = \frac{1}{N-1} \cdot \sum_{d=1}^{N} \left(x_i^d - \overline{x}_i \right)^T \cdot \left(x_j^d - \overline{x}_j \right) \quad \text{ where } \ \overline{x} = \frac{1}{N-1} \cdot \sum_{d=1}^{N} x^d$$

 Calculate the eigenvalues of the covariance matrix S for the given data and sort the eigenvalues of S:

$$\{\lambda_1, \lambda_2, ..., \lambda_n\}$$
 where $\lambda_1 \ge \lambda_2 \ge ... \ge \lambda_n \ge 0$.

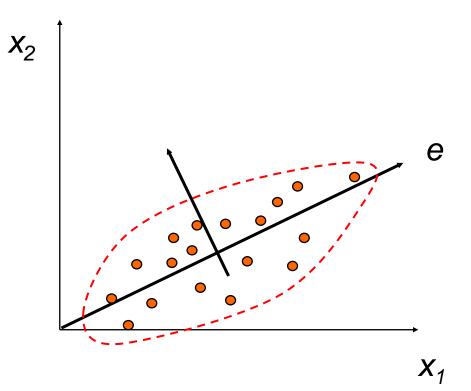
The eigenvalues are the variances of the data in the directions of ``their'' eigenvectors.

3. The **eigenvectors** $e_1, e_2, ..., e_n$ correspond to ``their' eigenvalues $\lambda_1, \lambda_2, ..., \lambda_n$,

The eigenvectors are called the *principal axes*.

Principal Component Analysis (PCA)

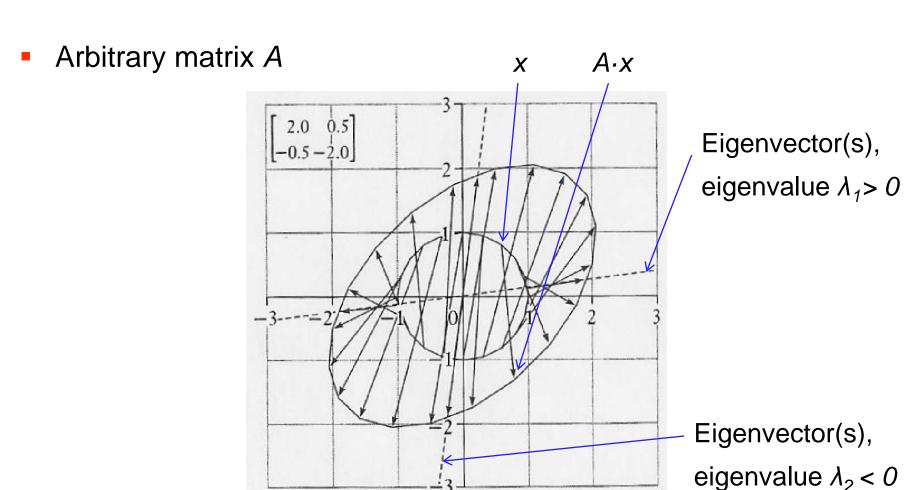
- Find a projection that captures the largest amount of variation in data
- The original data are projected onto a much smaller space, resulting in dimensionality reduction. We find the eigenvectors of the covariance matrix, and these eigenvectors define the new space



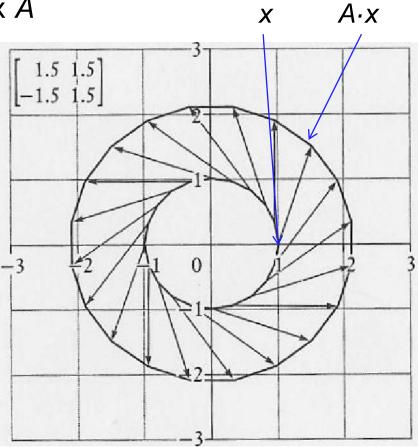
Symmetric matrix A $A \cdot x$ X [2.5 0.5] Eigenvector(s), $A \cdot e_1 = \lambda_1 \cdot e_1$ large eigenvalue λ_1 Eigenvector(s) $A \cdot e_2 = \lambda_2 \cdot e_2$

small eigenvalue λ_2

Non-symmetric matrix A $A \cdot x$ X 2.6 0.5 Eigenvector(s) $[-0.5 \ 1.4]$ Eigenvector(s)



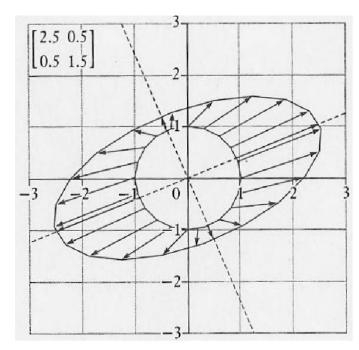
Rotation matrix A



No eigenvectors

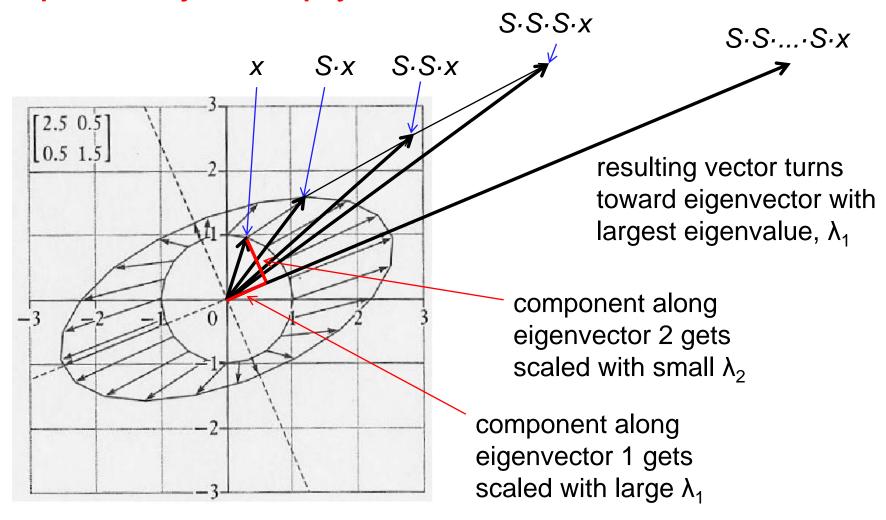
Multiply a Vector with a Covariance Matrix

- The covariance matrix S is a symmetric matrix
 - It always has eigenvectors
 - Eigenvectors are orthogonal to each other, if their eigenvalues differ
- Its eigenvalues are ≥ 0
 - S is positive (semi-) definite
- Eigenvalues are variances of the data in the directions of the corresponding eigenvectors

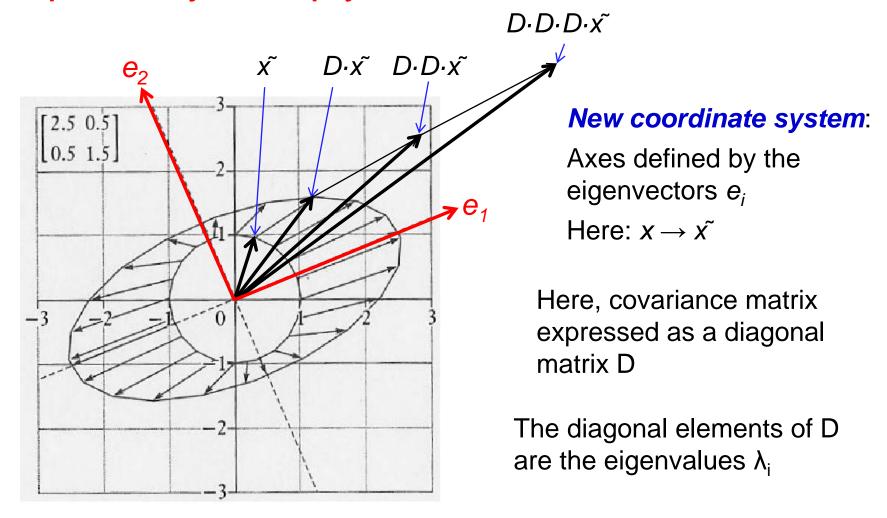


Multiplying by S does not rotate any vector by more than 90°

Repeatedly Multiply with the Covariance Matrix



Repeatedly Multiply with the Covariance Matrix



Easy to see: individual dimensions/components i of \tilde{x} are multiplied by λ_i

Obtain Largest Eigenvalue and Eigenvector

Iterative method

- Choose an initial random vector x
- Repeat
 - $x \leftarrow S \cdot x$
 - Normalize x to length 1
- Until converged. Then *normalized eigenvector* $e_1 = x$

Compute corresponding eigenvalue as the norm:

$$\lambda_1 = / S \cdot e_1 /$$

 Other linear algebra methods, e.g. SVD, compute all eigenvectors and corresponding eigenvalues.

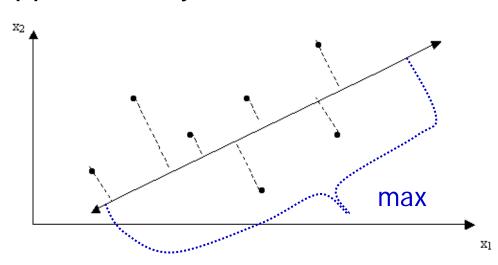
Principal Components Analysis

The data can be expressed in the new coordinate system

$$x = \overline{x} + \sum_{j=1}^{m} w_j \cdot e_j$$

- w_i are the data coordinates along the eigenvector axes
- m < n: data are only approximately reconstructed</p>

The first principal component is an axis in the direction of maximum variance.



Principal Components Analysis

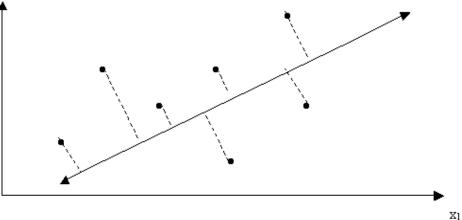
The criterion for features selection is based on the ratio R of the sum of the m largest eigenvalues (m≤n) of S to the trace of S (for example R>90%):

$$R = \frac{\displaystyle\sum_{i=1}^{m} \lambda_{i}}{\displaystyle\sum_{i=1}^{n} \lambda_{i}}$$
 sum over all explained variances

Trace of S

= sum over all variances

- Benefit:
 - non-explained variance $\sum_{i=m+1}^{n} \lambda_i$ may be small, even if m << n.



Principle Components Example: Eigenfaces

Eigenfaces are the (first few) principle components of a face image database.

reconstructed face eigenfaces

Sampling

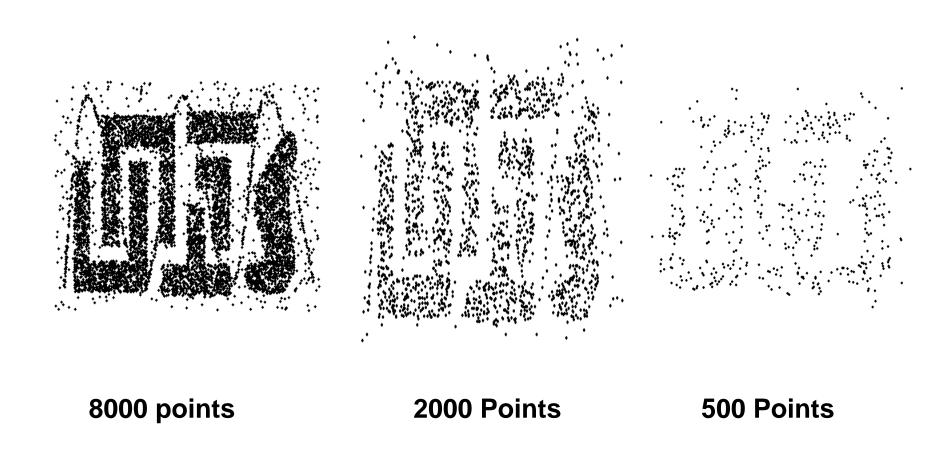
- Sampling: obtaining a small sample s to represent the whole data set N
- Allow a mining algorithm to run in complexity that is potentially sub-linear to the size of the data
- Key principle: Choose a representative subset of the data
 - Simple random sampling may have very poor performance in the presence of skewed data
 - Develop adaptive sampling methods, e.g., stratified sampling

Cases Reduction: Sampling ...

Key principle for effective sampling:

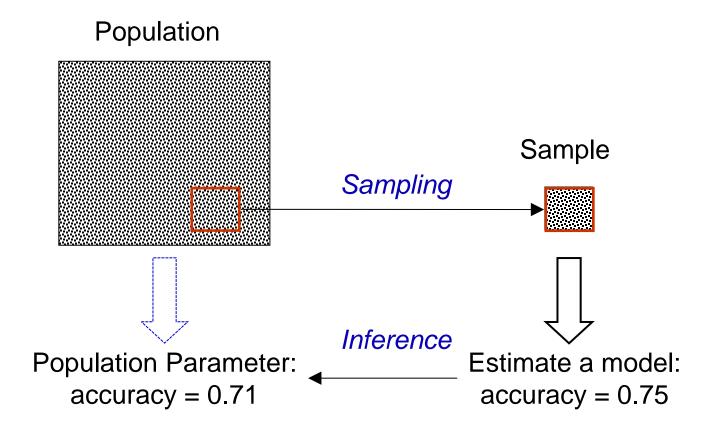
- Using a sample will work almost as well as using the entire data sets, if the sample is representative.
- A sample is representative if it has approximately the same property (of interest) as the original set of data.

Cases Reduction: Sample Size



Cases Reduction: Accuracy Parameter Estimation

 Challenging task: Infer the value of a population parameter based on a sample model.



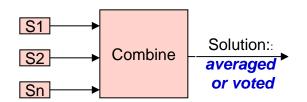
General-purpose Sampling Methods

Systematic sampling:

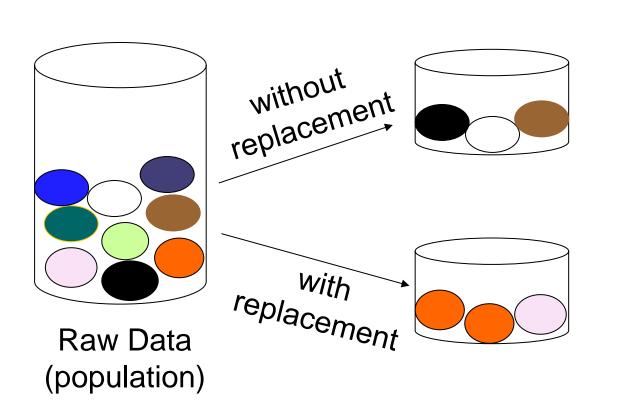
- For example 50% of a data set (every second sample)
- Simplest.
- Built in most of Data Mining tools
- Problem: regularities in data set!

Random sampling

- Random sampling without replacement,
- Random sampling with replacement.
- Average sampling: Combined solution from several subsets (randomly selected)
 - Stratified sampling



Sampling With or Without Replacement

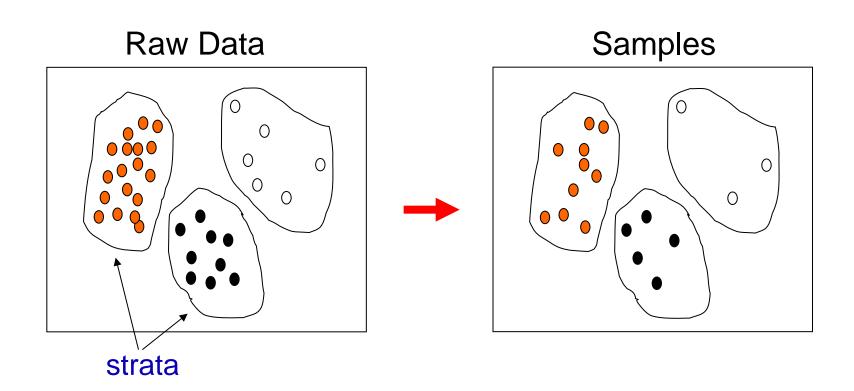


Once an object is selected, it is removed from the population

A selected object is not removed from the population

Stratified Sampling

- Partition the data set into strata (non-overlapping)
- Draw samples from each partition proportionally to its percentage in the data



Data Preprocessing

- Data Preprocessing: An Overview
 - Data Quality
 - Major Tasks in Data Preprocessing
- Data Cleaning
- **Data Integration**
- **Data Reduction**
- Data Transformation
- Data Discretization



Data Discretization Methods

- Reduce number of values for given continuous attribute by dividing into intervals
 - Binning: equal width binning and replacing bin by mean
 - Top-down split, unsupervised, no class information used
 - Histogram analysis
 - Top-down split, unsupervised, no class information used
 - Clustering analysis (unsupervised, top-down split or bottom-up merge)
 - Decision-tree analysis (supervised, top-down split)
 - Correlation (e.g., χ²) analysis (unsupervised, bottom-up merge)

Simple Discretization: Binning

- Equal-width (distance) partitioning
 - Divides the range into N intervals of equal size: uniform grid
 - If A and B are the lowest and highest values of the attribute, the width of intervals will be: W = (B A)/N.
 - Simple, but outliers may dominate presentation
 - Skewed data is not handled well
- Equal-depth (frequency) partitioning
 - Divides the range into N intervals, each containing approximately same number of samples
 - Good data scaling

Binning Methods for Data Smoothing

Sorted data for price (in dollars): 4, 8, 9, 15, 21, 21, 24, 25, 26, 28, 29, 34

Partition into equal-frequency (equi-depth) bins:

- Bin 1: 4, 8, 9, 15
- Bin 2: 21, 21, 24, 25
- Bin 3: 26, 28, 29, 34

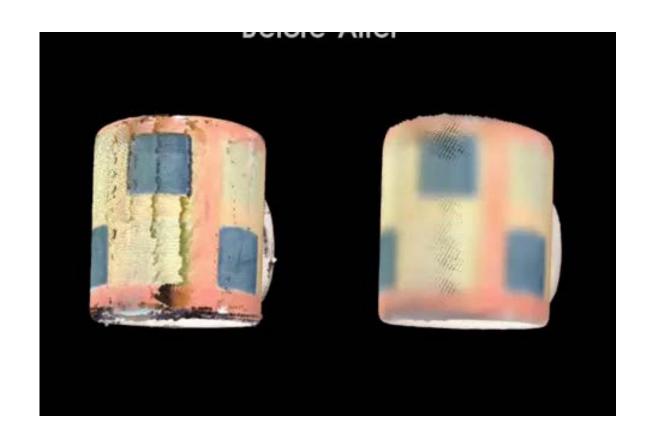
Smoothing by bin means:

- Bin 1: 9, 9, 9, 9
- Bin 2: 23, 23, 23, 23
- Bin 3: 29, 29, 29, 29

Smoothing by **bin boundaries**:

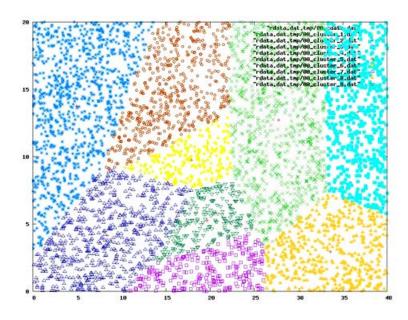
- Bin 1: 4, 4, 4, 15
- Bin 2: 21, 21, 21, 25
- Bin 3: 26, 26, 26, 34

Example: Data Resampling and Smoothing in Point Cloud Application



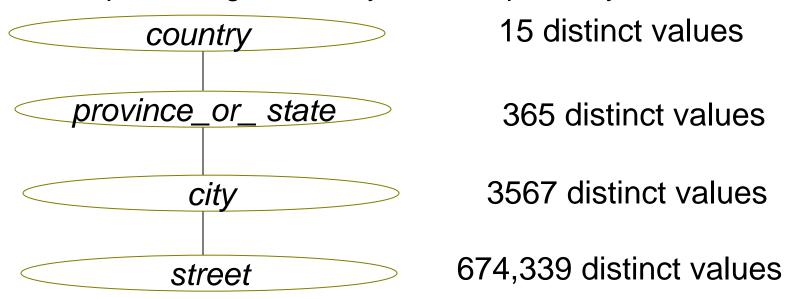
Clustering

- Partition data set into clusters based on similarity, and store cluster representation (e.g., centroid and diameter) only
- Can have hierarchical clustering and be stored in multidimensional index tree structures
- There are many choices of clustering definitions and clustering algorithms



Automatic Concept Hierarchy Generation

- Some hierarchies can be automatically generated based on the analysis of the number of distinct values per attribute in the data set
 - The attribute with the most distinct values is placed at the lowest level of the hierarchy
 - Exceptions, e.g., weekday, month, quarter, year



The time series of values can be expressed as a list:

```
X = \{t(1), t(2), t(3), ..., t(n)\},
where t(n) is the most recent value.
```

- For many problems based on time series the goal is to:
 - **forecast** t(n+1) from previous n values of the feature (or more general forecast t(n+j)), where these values are directly related to the predicted value, or
 - find patterns in time series.
- The most important step in preprocessing of row timedependent data is specification of a window or a time lag

Example: time series consisting of eleven measurements:

$$\mathbf{X} = \{t(0), t(1), t(2), t(3), t(4), t(5), t(6), t(7), t(8), t(9), t(10)\}$$

window size:

$$w=5$$

next value:

Sample	W M1	I N M2	D O	W M4	M5	Next Value
1	<u>t(0)</u>	t(1)	t(2)	t(3)	t(4)	t(5)
2	t(1)	t(2)	t(3)	t(4)	t(5)	t(6)
3	t(2)	t(3)	t(4)	t(5)	t(6)	t(7)
4	t(3)	t(4)	t(5)	t(6)	t(7)	t(8)
5	<u>t(4)</u>	t(5)	t(6)	t(7)	t(8)	t(9)
6	<u>t(</u> 5)	t(6)	t(7)	t(8)	t (9)	t(10)

Example: time series consisting of eleven measurements:

$$\mathbf{X} = \{t(0), t(1), t(2), t(3), t(4), t(5), t(6), t(7), t(8), t(9), t(10)\}$$

window size:

$$w=5$$

next value:

$$j=3$$

Sample	\mathbf{W}	I N	Next Value			
	M1	M2	М3	M4	M5	
1	<u>t(0)</u>	t(1)	t(2)	t(3)	t(4)	t(7)
2	<u>t(1)</u>	t(2)	t(3)	t(4)	t(5)	t(8)
3	$\widetilde{\mathfrak{t}}(2)$	t(3)	t(4)	t(5)	t(6)	t(9)
4	<u>t(3)</u>	t(4)	t(5)	t(6)	t(7)	t(10)

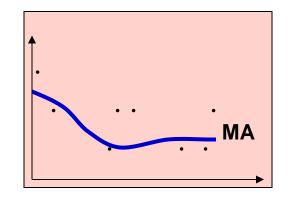
Time-dependent **2D** data

Samples prepared for window w = 3

Time	а	b		Sample	а	а	a(n)	b	b	b(n)
1	5	117			(n-2)	(n-1)		(n-2)	(n-1)	
2	8	113		1	5	8	4	117	113	116
3	4	116	-	2	8	4	9	113	116	118
4	9	118		3	4	9	10	116	118	119
5	10	119		3	7	9	10	110	110	119
6	12	120		4	9	10	12	118	119	120

 One way of summarizing features in the data set is to average them, producing "moving averages" (MA):

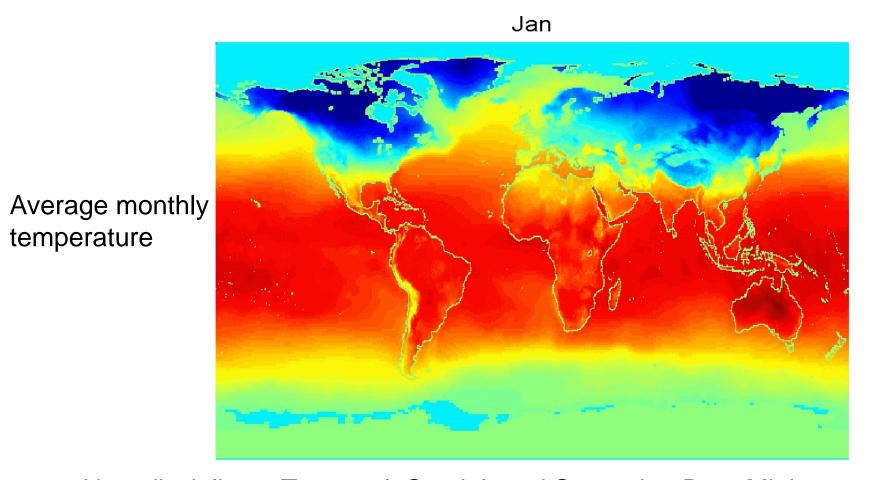
$$MA(i,m) = \frac{1}{m} \cdot \sum_{j=i-m+1}^{i} t(j)$$



The objective is to smooth neighboring time points by a moving average to reduce the random variation and noise components:

$$MA(i,m) = t(i) = mean(i) + error$$

Spatial-Temporal Data



New disciplines: Temporal, Spatial, and Streaming Data Mining

Noise example in real world from the WTM lab

<u>www.informatik.uni-hamburg.de/WTM</u> or <u>www.knowledge-technology.info</u>



Summary

- Data quality: accuracy, completeness, consistency, timeliness, believability, interpretability
- Data cleaning: e.g. missing/noisy values, outliers
- Data integration from multiple sources:
 - Entity identification problem; Remove redundancies; Detect inconsistencies
 - Chi-square for correlation analysis

Data reduction

Dimensionality reduction, e.g. by comparison of means and variances;
 Numerosity reduction; Data compression

Data transformation

- Normalization
- PCA

Data discretization

Binning, Clustering, Concept hierarchy generation

HiWi Position at WTM Available

- 20 40 hours (one-off, approx. €400-€500)
- Immediate availability
- Task: Organize the Publications on the WTM Web Page