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# 4. Data Preprocessing

Knowledge Discovery in Databases with Exercises

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**1. Overview**

**2. Data Cleaning**

**3. Data Integration**

**4. Data Reduction**

**5. Data Transformation and Data Discretization**

**6. Summary**

# Overview

- **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 refs, etc.
  - **Timeliness:** timely updated?
  - **Believability:** how trustworthy is it, that the data is correct?
  - **Interpretability:** how easily can the data be understood?
  - And even many more!

- **Data cleaning:**
  - Fill in missing values.
  - Smooth noisy data.
  - Identify or remove outliers.
  - Resolve inconsistencies.
- **Data integration:**
  - Integration of multiple databases.
  - Data cubes or files.

- **Data reduction:**
  - Dimensionality reduction.
  - Numerosity reduction.
  - Data compression.
- **Data transformation and data discretization:**
  - Normalization.
  - Concept-hierarchy generation.

# Data Cleaning

- **Data in the real world is dirty.**
- **Lots of different kinds of dirty data:**
  - **Incomplete data:** lacking attributes, lacking values or containing aggregate data.
  - **Inconsistencies:** containing discrepancies in codes or names.
  - **Errors:** containing incorrect values.
  - **Noise:** containing small inaccuracies.
  - **Outliers:** containing extreme values.

- **Potential reasons:**

- Data not yet available.
- Technical malfunction.
- Human error.
- etc.

- **Potential solutions:**

- Ignore the tuple.
- Fill in the missing value manually.
  - Often infeasible.
- Fill in automatically with:
  - A global constant.
  - The attribute mean.
  - The class mean.
  - The most probable value.

Mat. Nr.	Age
12345678	23
23061995	25
21241992	
	23
25052025	21
14912780	24

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- **Potential reasons:**

- Merging of data from different sources.
- Missing conventions.
- Human error.
- etc.

- **Potential solutions:**

- Manual data cleaning.
- (Semi-)Automatic data cleaning.
  - Most often common inconsistencies can be detected and solved via rule based approaches.

Applicant	Grade
124	1.0
Michael	2.3
134	3.7
323	A-
174	2.0
123	1.6

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- **Potential reasons:**

- Malfunctions.
- Transmission errors.
- Human error.
- etc.

- **Potential solutions:**

- Ignore the tuple.
- Manual data cleaning.
  - A subject matter expert (SME) is often needed to identify the errors.
- (Semi-)Automatic data cleaning.
  - Errors are often highly case dependent and therefore there is no general solution.

Module	ECTS
EADEIS	5
MoL	5
DL	5
EDB	7.5
KDDmUe	6
POIS	5

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- **Potential reasons:**

- Small sensor inaccuracies.
- Transmission errors.
- etc.

- **Potential solutions:**

- Data smoothing by:
  - Binning.
  - Regression.
  - Clustering.
  - etc.

Time	Temperature
08:01	14.123°C
08:02	14.153°C
08:03	14.163°C
08:04	14.723°C
08:05	14.126°C
08:06	14.463°C

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## Errors $\iff$ Noise

- Noise can be referred to as a special type of error.
- Not every error is noise!

- **Potential reasons:**

- Errors.
- Very rare events.

- **Potential solutions:**

- If an error, treat them as one.
- If a rare event, the outlier is interesting and can be used for further analysis.

Year	Max. Temp.
2026	32°C
2027	34°C
2028	33°C
2029	35°C
2030	61°C
2031	36°C

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## Errors $\iff$ Outliers

- Outliers might indicate errors.
- Not every outlier is an error!

- **Data discrepancy detection:**

- Use **metadata** (e.g. domain, range, dependency, distribution).
- Check field overloading.
- Check uniqueness rule, consecutive rule and null rule.
- Use commercial tools:
  - **Data scrubbing:** use simple domain knowledge (e.g. postal code, spell-check) to detect errors and make corrections.
  - **Data auditing:** by analyzing data to discover rules and relationships to detect violators (e.g. correlation and clustering to find outliers).

- **Data migration and integration:**
  - Data-migration tools: allow transformations to be specified.
  - ETL (Extraction/Transformation>Loading) tools: allow users to specify transformations through a graphical user interface.
- **Integration of the two processes.**
  - Iterative and interactive (e.g. the Potter's Wheel tool).

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# Data Integration

- **Data integration:**
  - Combine data from multiple sources into a coherent store.
- **Schema integration:**
  - E.g. A.cust-id  $\equiv$  B.cust-#.
  - Integrate metadata from different sources.
- **Entity-identification problem:**
  - Identify the same 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 (coding).
    - Different scales, e.g. metric vs. British units.

- **Redundant data often occur when integrating multiple databases.**
  - **Object (entity) 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:**
  - Can be detected by **correlation analysis** and **covariance analysis**.
- **Careful integration of the data from multiple sources:**
  - Helps to reduce/avoid redundancies and inconsistencies and improve mining speed and quality.

- **Example:**

We want to determine if the interests "Reads Books" and "Plays Chess" in the following table correlate with each other:

ID	Reads Books	Plays Chess
1	Y	Y
2	Y	Y
3	Y	N
...	...	...
1499	N	Y
1500	N	N

- **General starting point:**

- **The attributes A and B to be analyzed:**
  - A has  $n$  distinct values:  
 $A := \{a_1, a_2, \dots, a_n\}$ , where  $n \in \mathbb{N}_{>1}$ .
  - B has  $m$  distinct values:  
 $B := \{b_1, b_2, \dots, b_m\}$ , where  $m \in \mathbb{N}_{>1}$ .
- **The set X of all distinct combinations:**
  - X is defined as follows:  
 $X := \{(a, b) \mid a \in A \text{ and } b \in B\}$ .
- **The multi set Y of all tuples:**
  - The multiset Y over the set X is a mapping of X to the set of natural numbers  $\mathbb{N}_0$ . The number  $Y(x)$ ,  $x \in X$  tells how often x is contained in the multiset Y.

- **Starting point in the example:**

- **The attributes A and B to be analyzed:**
  - A ("Reads Books") has 2 distinct values:  
 $A := \{Y, N\}$
  - B ("Plays Chess") has 2 distinct values:  
 $B := \{Y, N\}$
- **The set X of all distinct combinations:**
  - X contains 4 distinct combinations:  
 $X := \{(Y, Y), (Y, N), (N, Y), (N, N)\}$ .
- **The multi set Y of all tuples:**
  - Y contains 1500 tuples:  
 $Y := \{(Y, Y), (Y, Y), \dots, (N, N)\}$ .

- Actual quantity in  $Y$ :

$$c_{ij} = \#\{(a, b) \in Y \mid a = a_i, b = b_j\} = Y((a_i, b_j))$$

- Expected quantity (value of  $c_{ij}$ ) in case of independence, i.e. no correlation:

$$e_{ij} = \frac{\sum_{k=1}^m c_{ik}}{\#Y} \cdot \frac{\sum_{l=1}^n c_{lj}}{\#Y} \cdot \#Y = \frac{\sum_{k=1}^m c_{ik} \cdot \sum_{l=1}^n c_{lj}}{\#Y}$$

## Please note that:

- The sum of all  $c_{ij}$  over an attribute  $a_i$  (or  $b_j$ ) is identical to the sum of all  $e_{ij}$  over  $a_i$  (or  $b_j$ ):

$$\sum_{k=1}^m e_{ik} = \sum_{k=1}^m c_{ik} \text{ and } \sum_{l=1}^n e_{lj} = \sum_{l=1}^n c_{lj}$$

- The values  $c_{ij}$  and  $e_{ij}$  are often presented in a **contingency table**:

	$a_1$	$\dots$	$a_n$	
$b_1$	$c_{11}(e_{11})$	$\dots$	$c_{n1}(e_{n1})$	$\sum_{i=1}^n e_{i1}$
$\dots$	$\dots$	$\dots$	$\dots$	$\dots$
$b_m$	$c_{1m}(e_{1m})$	$\dots$	$c_{nm}(e_{nm})$	$\sum_{i=1}^n e_{im}$
	$\sum_{j=1}^m e_{1j}$	$\dots$	$\sum_{j=1}^m e_{nj}$	$\sum_{i=1}^n \sum_{j=1}^m e_{ij}$

- In our example it would look like this:

	Plays Chess	Doesn't Play Chess	Sum (Row)
Reads Books	250 ( $e_{11}$ )	200 ( $e_{21}$ )	450
Doesn't Read Books	50 ( $e_{12}$ )	1000 ( $e_{22}$ )	1050
Sum (Column)	300	1200	1500

## Expected Quantity for "Plays Chess" & "Reads Books"

$$e_{11} = \frac{\sum_{k=1}^m c_{1k} \cdot \sum_{l=1}^n c_{l1}}{\# Y} = \text{_____} =$$

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## Expected Quantity for "Plays Chess" & "Reads Books"

$$e_{11} = \frac{\sum_{k=1}^m c_{1k} \cdot \sum_{l=1}^n c_{l1}}{\# Y} = \frac{300 \cdot 450}{\# Y} =$$

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## Expected Quantity for "Plays Chess" & "Reads Books"

$$e_{11} = \frac{\sum_{k=1}^m c_{1k} \cdot \sum_{l=1}^n c_{l1}}{\# Y} = \frac{300 \cdot 450}{1500} =$$

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- In our example it would look like this:

	Plays Chess	Doesn't Play Chess	Sum (Row)
Reads Books	250 (90)	200 ( $e_{21}$ )	450
Doesn't Read Books	50 ( $e_{12}$ )	1000 ( $e_{22}$ )	1050
Sum (Column)	300	1200	1500

## Expected Quantity for "Plays Chess" & "Reads Books"

$$e_{11} = \frac{\sum_{k=1}^m c_{1k} \cdot \sum_{l=1}^n c_{l1}}{\# Y} = \frac{300 \cdot 450}{1500} = 90$$

- The values  $c_{ij}$  and  $e_{ij}$  are often presented in a **contingency table**:

	$a_1$	$\dots$	$a_n$	
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$\dots$	$\dots$	$\dots$	$\dots$	$\dots$
$b_m$	$c_{1m}(e_{1m})$	$\dots$	$c_{nm}(e_{nm})$	$\sum_{i=1}^n e_{im}$
	$\sum_{j=1}^m e_{1j}$	$\dots$	$\sum_{j=1}^m e_{nj}$	$\sum_{i=1}^n \sum_{j=1}^m e_{ij}$

- In our example it would look like this:

	Plays Chess	Doesn't Play Chess	Sum (Row)
Reads Books	250 (90)	200 (360)	450
Doesn't Read Books	50 (210)	1000 (840)	1050
Sum (Column)	300	1200	1500

## Expected Quantity for "Plays Chess" & "Reads Books"

$$e_{11} = \frac{\sum_{k=1}^m c_{1k} \cdot \sum_{l=1}^n c_{l1}}{\# Y} = \frac{300 \cdot 450}{1500} = 90$$

- To determine the correlation the  $\chi^2$ -test (Chi-squared test) is applied:

$$\chi^2 = \sum_{i=1}^n \sum_{j=1}^m \frac{(c_{ij} - e_{ij})^2}{e_{ij}}.$$

- Calculation of  $\chi^2$  in our example:

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

## Null hypothesis of the $\chi^2$ -test

- The  $\chi^2$ -test is used to test the null hypothesis  $H_0$  of independence (i.e. no correlation).
- Which  $\chi^2$  value indicates correlation?
  - The  $\chi^2$  value is compared with a critical value from the  $\chi^2$  distribution (see table on the next slide).
  - Before that is done the degrees of freedom (df) must be calculated:

$$df = (n - 1) \cdot (m - 1)$$

Where  $n$  is the count of distinct values in  $A$  and  $m$  of distinct values in  $B$ .

- And a significance level  $\alpha$  must be defined (e.g.  $\alpha = 0.005$ ).

- In our example:

- The degrees of freedom (df) are:

$$df = (2 - 1) \cdot (2 - 1) = 1.$$

<b>df / <math>\alpha</math></b>	<b>0.025</b>	<b>0.010</b>	<b>0.005</b>
<b>1</b>	5.024	6.635	7.879
<b>2</b>	7.378	9.210	10.597
<b>3</b>	9.348	11.345	12.838
<b>4</b>	11.143	13.277	14.860
<b>5</b>	12.833	15.086	16.750
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<sup>1</sup>Good link for a full table: [https://www.hawkeslearning.com/documents/statdatasets/stat\\_tables.pdf](https://www.hawkeslearning.com/documents/statdatasets/stat_tables.pdf)

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- In our example:

- The degrees of freedom (df) are:

$$df = (2 - 1) \cdot (2 - 1) = 1.$$

- We set the significance level to  $\alpha = 0.005$
- The critical value from the  $\chi^2$  distribution<sup>1</sup> is:

$$\chi^2_{0.005,1} = 7.879.$$

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- We set the significance level to  $\alpha = 0.005$
- The critical value from the  $\chi^2$  distribution<sup>1</sup> is:

$$\chi^2_{0.005,1} = 7.879.$$

- Our  $\chi^2$ -value is bigger than the critical value:

$$\chi^2 = 507.93 > 7.879.$$

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<b>1</b>	5.024	6.635	7.879
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<b>3</b>	9.348	11.345	12.838
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<b>5</b>	12.833	15.086	16.750
<b>6</b>	14.449	16.812	18.548
<b>7</b>	16.013	18.475	20.278
<b>8</b>	17.535	20.090	21.955
<b>9</b>	19.023	21.666	23.589

<sup>1</sup>Good link for a full table: [https://www.hawkeslearning.com/documents/statdatasets/stat\\_tables.pdf](https://www.hawkeslearning.com/documents/statdatasets/stat_tables.pdf)

- In our example:

- The degrees of freedom (df) are:

$$df = (2 - 1) \cdot (2 - 1) = 1.$$

- We set the significance level to  $\alpha = 0.005$
- The critical value from the  $\chi^2$  distribution<sup>1</sup> is:

$$\chi^2_{0.005,1} = 7.879.$$

- Our  $\chi^2$ -value is bigger than the critical value:

$$\chi^2 = 507.93 > 7.879.$$

- Therefore we reject the null hypothesis  $H_0$  and conclude that there is correlation between the two attributes.

$df/\alpha$	<b>0.025</b>	<b>0.010</b>	<b>0.005</b>
<b>1</b>	5.024	6.635	7.879
<b>2</b>	7.378	9.210	10.597
<b>3</b>	9.348	11.345	12.838
<b>4</b>	11.143	13.277	14.860
<b>5</b>	12.833	15.086	16.750
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- Numerical correlation can be determined with Pearson's product-moment coefficient:

$$\text{Cor}(A, B) = \frac{\sum_{i=1}^n (a_i - \mu_A)(b_i - \mu_B)}{n \cdot \sigma_A \sigma_B} = \frac{\sum_{i=1}^n a_i b_i - n \cdot \mu_A \mu_B}{n \cdot \sigma_A \sigma_B}.$$

where  $n$  is the number of tuples,  $a_i$  and  $b_i$  are the respective values of  $A$  and  $B$  in tuple  $i$ ,  $\mu_A$  and  $\mu_B$  are the respective mean values of  $A$  and  $B$ ,  $\sigma_A$  and  $\sigma_B$  are the respective standard deviations of  $A$  and  $B$

## Properties of Pearson's product-moment coefficient

- If  $\text{Cor}(A, B) > 0$ :  $A$  and  $B$  are positively correlated (the closer to 1, the stronger the correlation).
- If  $\text{Cor}(A, B) = 0$ :  $A$  and  $B$  are independent.
- If  $\text{Cor}(A, B) < 0$ :  $A$  and  $B$  are negatively correlated (the closer to  $-1$ , the stronger the correlation).

- It is also possible to visually detect numerical correlation:

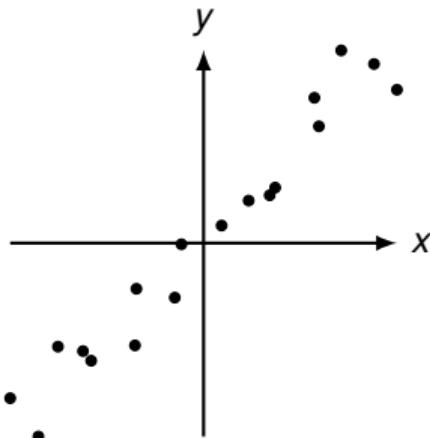


Figure: a) Positive correlation.

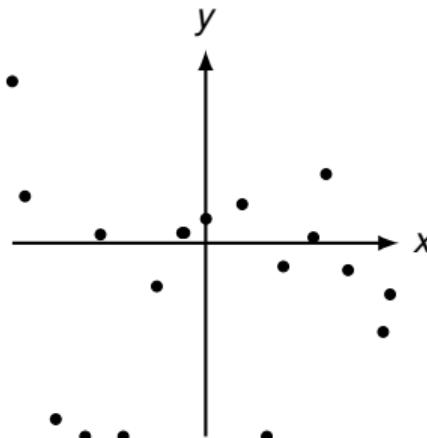


Figure: b) Uncorrelated/no correlation.

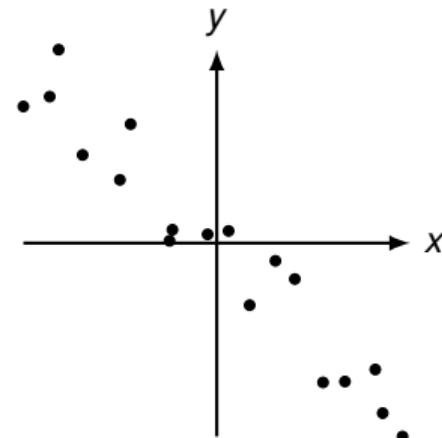
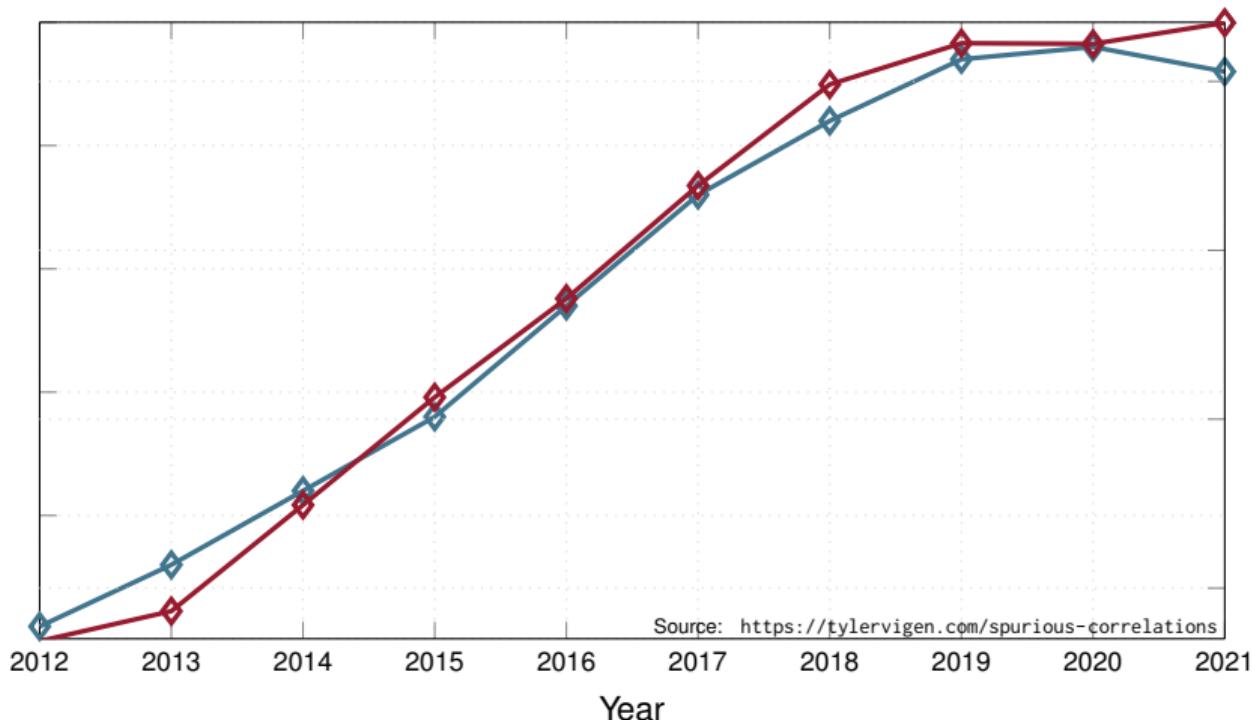
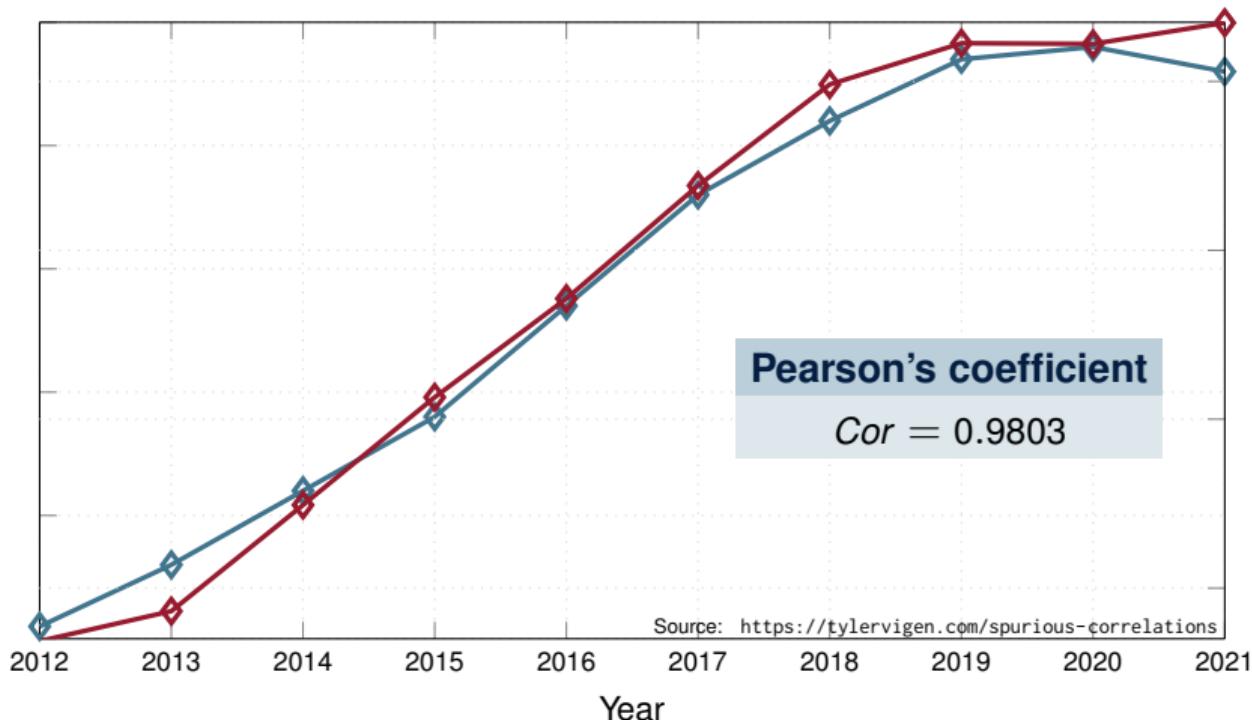


Figure: c) Negative correlation.

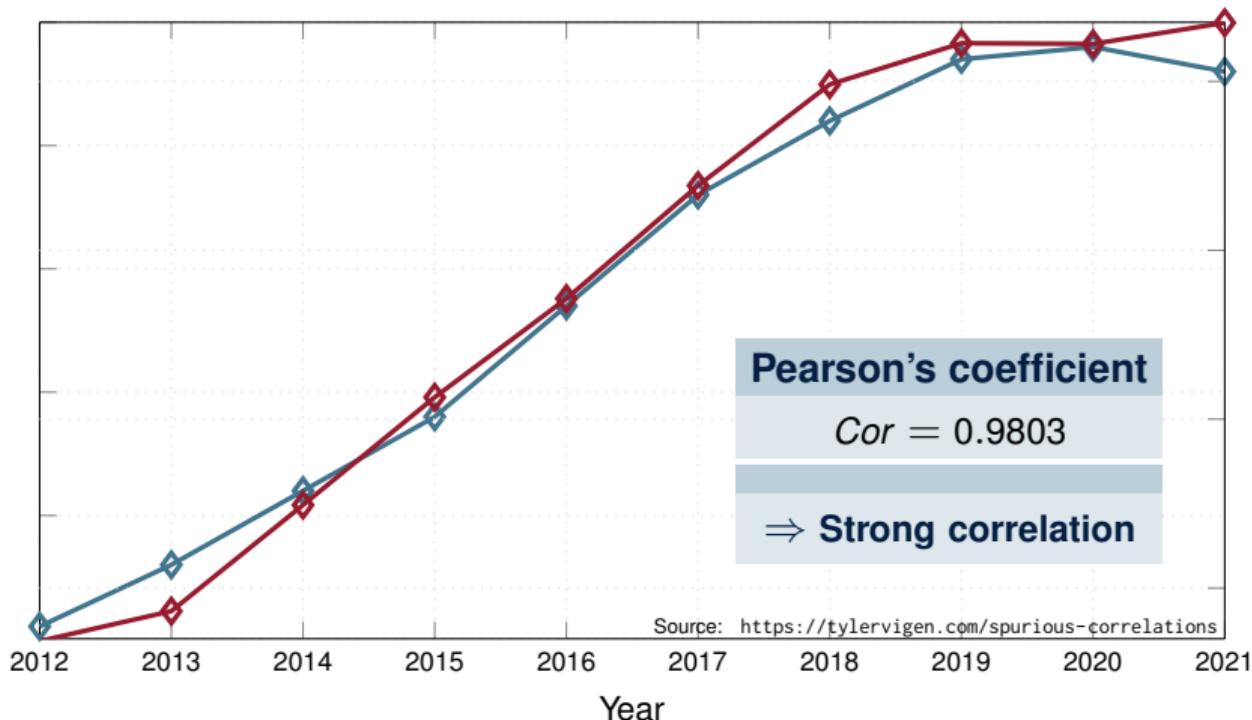
# Correlation vs. Causality



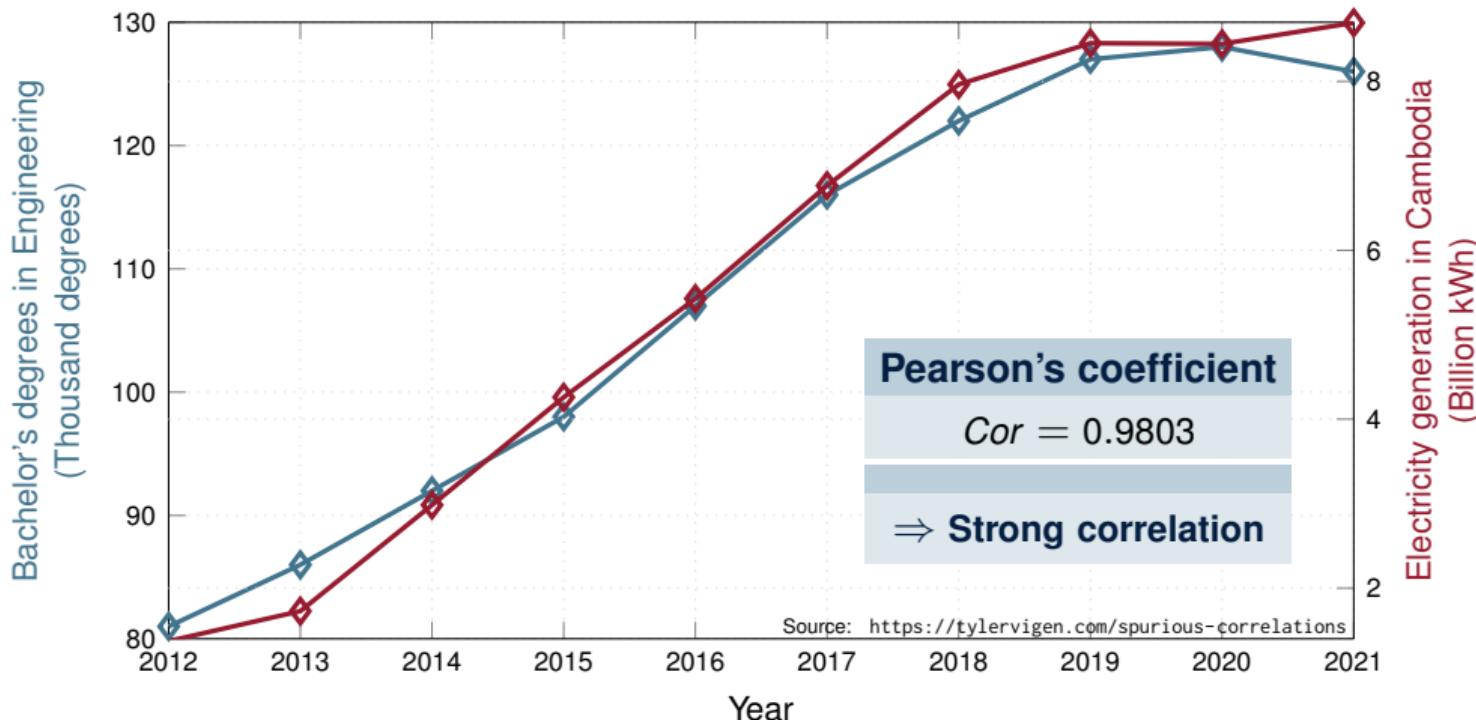
# Correlation vs. Causality



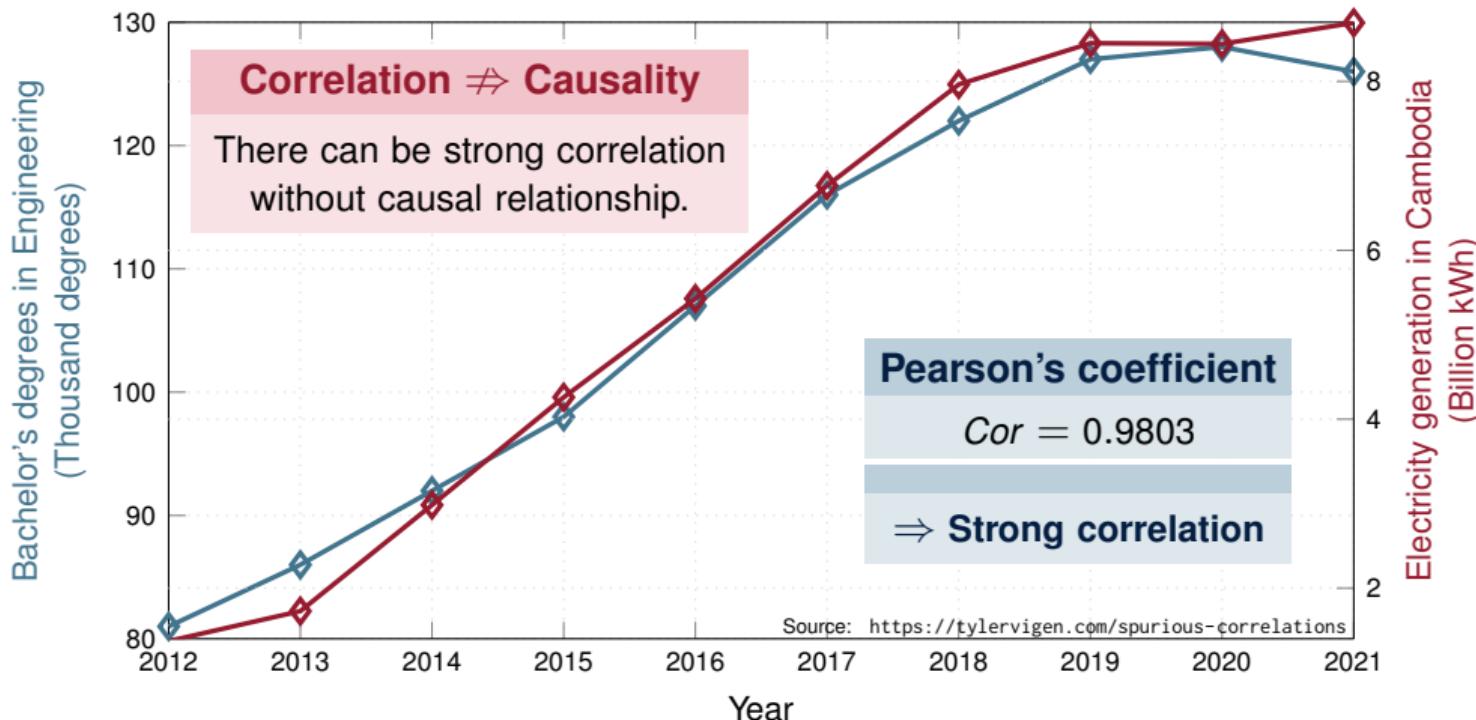
# Correlation vs. Causality



# Correlation vs. Causality



# Correlation vs. Causality



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# Data Reduction

- **What is data reduction?**
  - Obtain a reduced representation of the data set that is much smaller in volume but yet produces the same (or almost the same) results.
- **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.
- **Data reduction strategies:**
  - Dimensionality reduction, i.e. remove unimportant attributes.
    - Wavelet transforms.
    - Principal component analysis.
    - Attribute subset selection or attribute creation.

- **Data reduction strategies (continued):**

- Numerosity reduction:
  - Regression and log-linear models.
  - Histograms, clustering and sampling.
  - Data cube aggregation.
- Data compression.

- **Curse of dimensionality:**
  - When dimensionality increases data becomes increasingly sparse.
  - Density and distance between points become 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.
- **Dimensionality-reduction techniques:**
  - Wavelet transforms.
  - Principal component analysis.
  - Supervised and nonlinear techniques (e.g. feature selection).

- **Discrete wavelet transform:**  
Transforms a vector  $X$  into a different vector  $X'$  of wavelet coefficients with the same length.
- **Compressed approximation:**  
Store only a small fraction of the strongest of the wavelet coefficients.
- **Similar to discrete fourier transform, but better lossy compression, localized in space.**
- **Method:**
  - The length of the vector must be an integer power of 2 (padding with 0's if necessary).
  - Each transform has two functions: smoothing and difference.
  - Applied to pairs of data, resulting in two sets of data with half the length.
  - The two functions are applied recursively until reaching the desired length.

- **Initial vector:**
  - $X = (2, 2, 0, 2, 3, 5, 4, 4)$
- **First step:**
  - $(2, 2) \rightarrow$  Average: 2, Weighted difference: 0
  - $(0, 2) \rightarrow$  Average: 1, Weighted difference: -1
  - $(3, 5) \rightarrow$  Average: 4, Weighted difference: -1
  - $(4, 4) \rightarrow$  Average: 4, Weighted difference: 0
  - $A_1 = (2, 1, 4, 4), D_1 = (0, -1, -1, 0)$
- **Second step:**
  - $(2, 1) \rightarrow$  Average: 1.5, Weighted difference: 0.5
  - $(4, 4) \rightarrow$  Average: 4, Weighted difference: 0
  - $A_2 = (1.5, 4), D_2 = (0.5, 0)$

- **Third step:**

- $(1.5, 4) \rightarrow$  Average: 2.75, Weighted difference: -1.25
- $A_3 = (2.75), D_3 = (-1.25)$

- **Resulting vector:**

- $X' = (2.75, -1.25, 0.5, 0, 0, -1, -1, 0)$

- **Possible compression:**

- Small detail coefficients ( $D_{1,2,3}$ ) can be replaced by 0's, while retaining significant coefficients.

Resolution	Averages	Detail coefficients
8	$(2, 2, 0, 2, 3, 5, 4, 4)$	-
4	$(2, 1, 4, 4)$	$(0, -1, -1, 0)$
2	$(1.5, 4)$	$(0.5, 0)$
1	$(2.75)$	$(-1.25)$

- **Main idea:**
  - Given a data set with  $n$  dimensions.
  - Find  $k \leq n$  orthogonal vectors that capture the largest amount of data.
  - Works only for numeric data.
- **Example data set:**
  - Used on the next few slides to explain the steps of a PCA:

$d_1$	$d_2$	$d_3$
23	6	1
9	9	5
17	5	1
3	6	1

- **Procedure:**

- Each value  $x$  within a dimension  $d_n$  is standardized with the help of the mean ( $\mu_{d_n}$ ) and standard deviation ( $\sigma_{d_n}$ ) of  $d_n$ :

$$x' = \frac{x - \mu_{d_n}}{\sigma_{d_n}}$$

- **Reason:**

- Each dimension should be considered equally in the analysis.
- Dimensions with a wider range of values would dominate without this step.

- **Example:**

- Mean and standard deviation per dimension:

	$d_1$	$d_2$	$d_3$
$\mu$	13.000000	6.500000	2.0
$\sigma$	8.793937	1.732051	2.0

- Standardized data set:

	$d_1$	$d_2$	$d_3$
	+1.137147	-0.288675	-0.5
	-0.454859	+1.443376	+1.5
	+0.454859	-0.866025	-0.5
	-1.137147	-0.288675	-0.5

- **Procedure:**

- A  $n \times n$  covariance matrix is generated that contains the covariance between each possible attribute pairing. When the dimensions are compared with themselves, the variance always replaces the covariance:

$$\begin{bmatrix} \text{Var}(d_1) & \dots & \text{Cov}(d_1, d_n) \\ \dots & \dots & \dots \\ \text{Cov}(d_n, d_1) & \dots & \text{Var}(d_n) \end{bmatrix}$$

- **Reason:**

- Dimensions that are highly correlated contain redundant information.
- This step helps to identify these correlations.

- **Example:**

- The  $3 \times 3$  covariance matrix of our example:

	$d_1$	$d_2$	$d_3$
$d_1$	+1.000000	-0.350150	-0.303239
$d_2$	-0.350150	+1.000000	+0.962250
$d_3$	-0.303239	+0.962250	+1.000000

- **Procedure:**

- The eigenvectors and eigenvalues of the covariance matrix ( $C$ ) are computed by solving the following equation:

$$C\nu = \lambda\nu$$

- If an  $n$  digit vector  $\nu$  satisfies this equation for a  $\lambda \in \mathbb{R}$ , then  $\nu$  is called an eigenvector with associated eigenvalue  $\lambda$

- **Reason:**

- The determined eigenvectors are called **principal components** of the dataset. The eigenvalues indicate which of these principal components has which importance for the significance of the dataset.
- By sorting the eigenvectors in descending order according to their eigenvalues, the principal components that contain the most information can be identified.

- **Example:**

- Eigenvalues and eigenvectors in our example:

$$\lambda_1 = +2.14823654, \nu_1 = \begin{bmatrix} +0.37342507 \\ -0.92684562 \\ -0.03887043 \end{bmatrix}$$

$$\lambda_2 = +0.81530433, \nu_2 = \begin{bmatrix} -0.66009198 \\ -0.23604255 \\ -0.71313568 \end{bmatrix}$$

$$\lambda_3 = +0.03645914, \nu_3 = \begin{bmatrix} -0.6517916 \\ -0.2919608 \\ +0.69994757 \end{bmatrix}$$

- Sorting these three eigenvectors by their significance, we arrive at the order  $\nu_1, \nu_2, \nu_3$

- **Procedure:**

- The top N eigenvectors are selected to create a feature matrix from them.
- There is no fixed rule exactly how many eigenvectors should be selected.
- The dimensionality reduction is larger the fewer eigenvectors are chosen.
- The information loss increases with each eigenvector that is discarded.

- **Reason:**

- It must be considered carefully how much information can be given up in favor of dimensionality reduction.

- **Example:**

- In our example  $\nu_1$  carries approx. 72% of the information:

$$\frac{2.14823654}{2,14823654 + 0,81530433 + 0,03645914} = 0.71607885$$

- It might be interesting to keep only the eigenvector  $\nu_1$  and discard the other two eigenvectors. Our feature matrix therefore looks as follows:

$$\begin{bmatrix} +0.37342507 \\ -0.92684562 \\ -0.03887043 \end{bmatrix}$$

- **Procedure:**

- The original data set ( $D$ ) gets multiplied with the feature matrix ( $F$ ), to create a new data set ( $N$ ) with lower dimensionality:

$$N = D \cdot F$$

- **Reason:**

- This step applies the dimensionality reduction to each tuple.
- The PCA is completed with this step.

- **Example:**

- Our dataset after the transformation and with the PCA completed looks like this:

$$\begin{bmatrix} +0.711632 \\ -1.565948 \\ +0.991963 \\ -0.137647 \end{bmatrix}$$

- It is to be expected that this dataset still contains about 72% of its original information, which can be further used for data mining, while having to deal with a lot less dimensions.

- Another way to reduce dimensionality of data.
- **Redundant attributes:**
  - Duplicate much or all of the information contained in other attributes.
    - E.g. purchase price of a product and the amount of sales tax paid.
- **Irrelevant attributes:**
  - contain no information that is useful for the data-mining task at hand.
    - E.g. students' ID is often irrelevant to the task of predicting students' GPA.

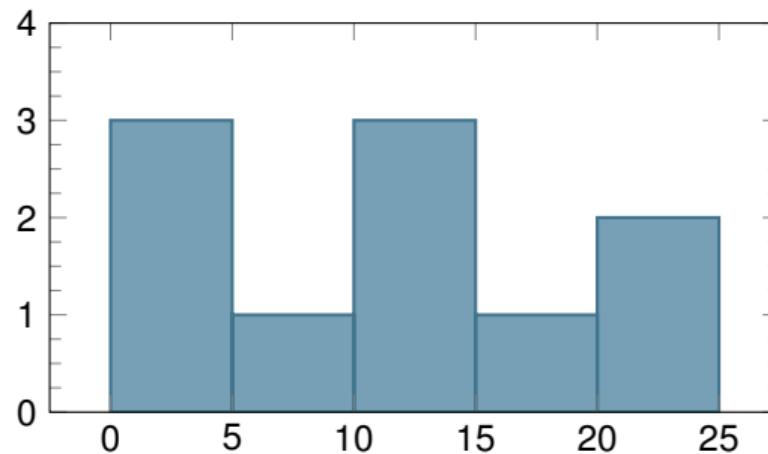
- There are  $2^d$  possible attribute combinations of  $d$  attributes.
- Typical heuristic attribute-selection methods:
  - Best single attribute under the attribute-independence assumption:  
choose by significance tests (e.g. t-test, see Chapter 7 “Classification”).
  - Best step-wise feature selection:
    - The best single attribute is picked first.
    - Then next best attribute condition to the first ...
- Step-wise attribute elimination:
  - Repeatedly eliminate the worst attribute.
- Best combined attribute selection and elimination.
- Optimal branch and bound:
  - Use attribute elimination and backtracking.

- Create new attributes (features) that can capture the important information in a data set more effectively than the original ones.
- Three general methodologies:
  - Attribute extraction.
    - Domain-specific.
  - Mapping data to new space (see: data reduction).
    - E.g. Fourier transformation, wavelet transformation, manifold approaches (not covered).
  - Attribute construction:
    - Combining features (see: discriminative frequent patterns in Chapter 5).
    - Data discretization.

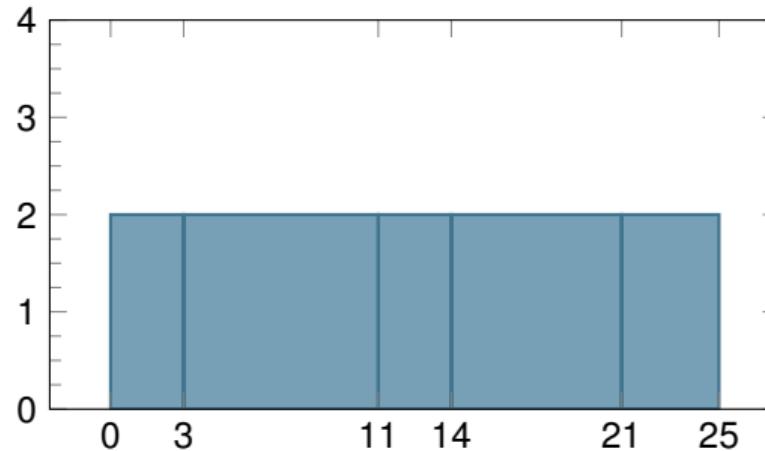
- Reduce data volume by choosing alternative, smaller forms of data representation.
- **Parametric methods** (e.g., regression):
  - Assume the data fits some **model** (e.g. a function).
  - Estimate model parameters.
  - Store only the parameters.
  - Discard the data (except possible outliers):
    - Ex. log-linear models obtain value at a point in  $m$ -dimensional space as the product of appropriate marginal subspaces.
- **Non-parametric methods:**
  - Do not assume models.
  - Major families: histograms, clustering, sampling, ...

- Divide data into buckets and store aggregate (e.g. average) of each bucket.
- Two different partitioning rules:
  - Equal-width: equal width of each bucket.
  - Equal-frequency (or equal-depth): equal number of tuples in each bucket.

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- **Partition data set into clusters based on similarity and store cluster representation (e.g., centroid and diameter) only.**
  - Can be very effective if data points are close to each other under a certain norm and choice of space.
  - Can have hierarchical clustering and be stored in multidimensional index-tree structures.
  - There are many choices of clustering algorithms.
  - Cluster analysis will be studied in depth in Chapter 7.

- Obtain a small sample  $x$  to represent the whole data set  $X$ .
- 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 skew.
  - Develop adaptive sampling methods, e.g. stratified sampling.
- Note: Sampling may not reduce database I/Os.
  - One page at a time.

- **Simple random sampling.**
  - There is an equal probability of selecting any particular item.
- **Sampling without replacement.**
  - Once an object is selected, it is removed from the population.
- **Sampling with replacement.**
  - A selected object is not removed from the population.
- **Stratified sampling:**
  - Partition the data set and draw samples from each partition: Proportionally, i.e. approximately the same percentage of the data.
  - Used in conjunction with skewed data.

- **String compression.**
  - There are extensive theories and well-tuned algorithms.
  - Typically lossless, but only limited manipulation is possible without expansion.
- **Audio/video compression.**
  - Typically lossy compression, with progressive refinement.
  - Sometimes small fragments of signal can be reconstructed without reconstructing the whole.
- **Time sequence is not audio.**
  - Typically short and varies slowly with time.
- **Dimensionality and numerosity reduction may also be considered as forms of data compression.**

# Data Transformation and Data Discretization

- Functions applied to a finite set of samples.
- **Methods:**
  - Smoothing: Remove noise from data.
  - Attribute/feature construction: New attributes constructed from the given ones.
  - Aggregation: Summarization, data-cube construction.
  - Normalization: Scaled to fall within a smaller, specified range.
    - Min-max normalization
    - Z-score normalization.
    - Normalization by decimal scaling.
  - Discretization: concept-hierarchy climbing.

- **Min-max normalization (to some interval [min, max]):**

$$a_{\text{new}} = \frac{a - \min_A}{\max_A - \min_A} (\max - \min) + \min.$$

Example: let income range from \$12,000 to \$98,000 normalized to [0, 1].

Then \$73,600 is mapped to  $\frac{73,600 - 12,000}{98,000 - 12,000} (1 - 0) + 0 = 0.716$ .

- **Z-score normalization:**

$$a_{\text{new}} := z(a) = \frac{a - \mu_A}{\sigma_A}, \text{ with } \mu \text{ being the mean and } \sigma \text{ the standard deviation.}$$

Example: let  $\mu = 54,000$  and  $\sigma = 16,000$ . Then  $\frac{73,000 - 54,000}{16,000} = 1.188$ .

- **Normalization by decimal scaling:**

$$a_{\text{new}} = \frac{a}{10^k}, \text{ where } k \text{ is the smallest integer such that } \max(|a_{\text{new}}|) < 1.$$

- **Three types of attributes:**
  - Nominal – values from an unordered set, e.g. color, profession.
  - Ordinal – values from an ordered set, e.g. military or academic rank.
  - Numerical – numbers, e.g. integer or real numbers.
- **Divide the value range of a continuous attribute into intervals:**
  - **Interval labels** can then be used to replace actual data values.
  - Reduce data size by discretization.
  - Supervised vs. unsupervised.
  - Split (top-down) vs. merge (bottom-up).
  - Discretization can be performed recursively on an attribute.
  - Prepare for further analysis, e.g. classification.

- **Typical methods:**
  - All the methods can be applied recursively.
  - **Binning:**
    - Unsupervised, top-down split.
  - **Histogram analysis:**
    - Unsupervised, top-down split.
  - **Clustering analysis:**
    - Unsupervised, top-down split or bottom-up merge.
  - **Decision-tree analysis:**
    - Supervised, top-down split.
  - **Correlation (e.g.  $\chi^2$ ) analysis:**
    - Unsupervised, bottom-up merge.

- **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:  $W = \frac{(B-A)}{N}$ .
- The most straightforward, but outliers may dominate presentation.
- Skewed data is not handled well.

- **Equal-depth (frequency) partitioning:**

- Divides the range into  $N$  intervals, each containing approximately the same number of samples.
- Good data scaling.
- Managing categorical attributes can be tricky.

- **Sorted data for price (in dollars):**  
4, 8, 9, 15, 21, 21, 24, 25, 26, 28, 29, 34.
- **Partition into equal-frequency (equal-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, 25, 25,  
Bin 3: 26, 26, 26, 34.

- **Classification:**

- E.g. decision-tree analysis.
- Supervised: Class labels given for training set e.g. cancerous vs. benign.
- Using **entropy** to determine split point (discretization point).
- Top-down, recursive split.
- Details will be covered in Chapter 6.

- **Correlation analysis:**

- E.g.  $\chi^2$ -merge:  $\chi^2$ -based discretization.
- Supervised: use class information.
- Bottom-up merge: find the best neighboring intervals (those having similar distributions of classes, i.e., low  $\chi^2$  values) to merge.
- Merge performed recursively, until a predefined stopping condition.

- **Concept hierarchy:**
  - Organizes concepts (i.e. attribute values) hierarchically.
  - Usually associated with each dimension in a data warehouse.
  - Facilitates **drilling and rolling** in data warehouses to view data at multiple granularity.
- **Concept-hierarchy formation:**
  - Recursively reduce the data by collecting and replacing **low-level concepts** (such as numerical values for age) by **higher-level concepts** (such as youth, adult, or senior).
  - Can be explicitly specified by domain experts and/or data-warehouse designers.
  - Can be automatically formed for both numerical and nominal data.
  - For numerical data, use discretization methods shown.

- Specification of a partial/total ordering of attributes explicitly at the schema level by users or experts.
  - $\#(\text{streets}) \prec \#(\text{city}) \prec \#(\text{state}) \prec \#(\text{country})$ .
- Specification of a hierarchy for a set of values by explicit data grouping.
  - $\#(\{\text{"Urbana"}, \text{"Champaign"}, \text{"Chicago"}\}) \prec \#(\text{Illinois})$ .
- Specification of only a partial set of attributes.
  - Only  $\#(\text{street}) \prec \#(\text{city})$ , not others.
- Automatic generation of hierarchies (or attribute levels) by the analysis of the number of distinct values.
  - E.g. for a set of attributes:  $\{\text{street}, \text{city}, \text{state}, \text{country}\}$ .
  - See on the next slides.

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

$$\begin{aligned}\#\text{(streets)} &= 674.339 > \#\text{(city)} = 3567, \\ \#\text{(city)} &= 3567 > \#\text{(province or state)} = 356, \\ \#\text{(province or state)} &= 356 > \#\text{(country)} = 15.\end{aligned}$$

# 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.
- **Data reduction:**
  - Dimensionality reduction.
  - Numerosity reduction.
  - Data compression.
- **Data transformation and data discretization:**
  - Normalization.
  - Concept-hierarchy generation.

## Any questions about this chapter?

Ask them now or ask them later in our forum:



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