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Chair of Business Information Systems, esp. Intelligent Systems and Services

Data Science: Advanced Analytics

Segmentation Methods

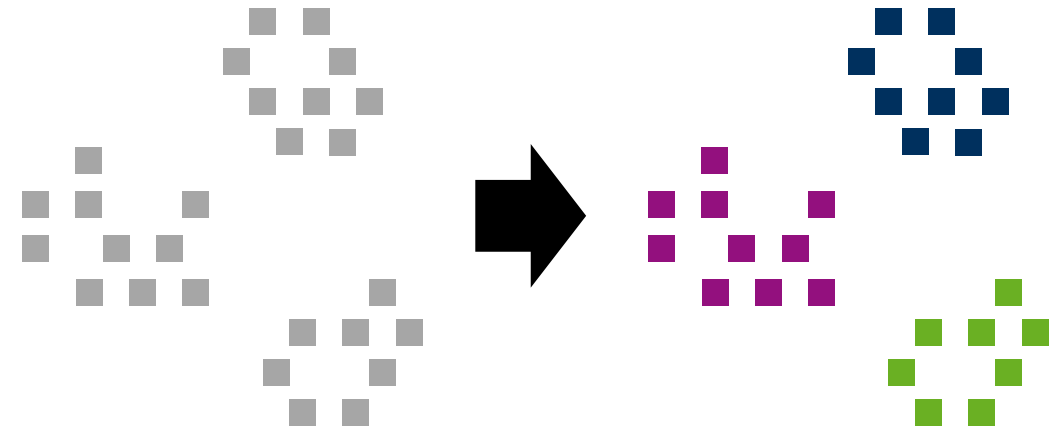
Dresden // 26.04.2023
Sommersemester 2023



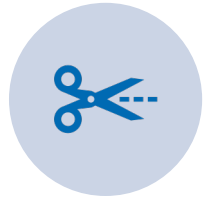
Recap: Aims of the cluster analysis

Segmentation or cluster analysis is used to group objects and/or characteristics into classes or groups so that

- between the elements of the same classes the greatest possible similarity,
- between the elements of different classes the greatest possible diversity is achieved

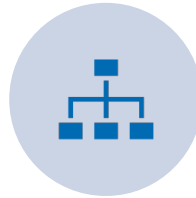


Clustering Methods



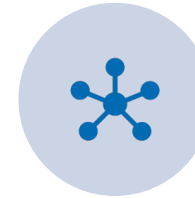
Partitional Clustering

K-Means & K-Medoids
For n objects or data tuples, a partitioning method constructs k partitions of the data, where each partition represents a cluster $k \leq n$.



Hierarchical Clustering

Agglomerative, Divisive
Grouping data objects into a tree (dendrogram) of clusters



Density-Based Clustering

DBSCAN
Grouping data tuples along density-connected points.



Grid-Based Clustering

STING, WaveCluster & CLIQUE
Method uses a multi-resolution grid data structure. E.g. spatial area is divided into rectangular cells (STING)



Others

Model-Based Clustering, Fuzzy, Evolutionary, Simulated Annealing

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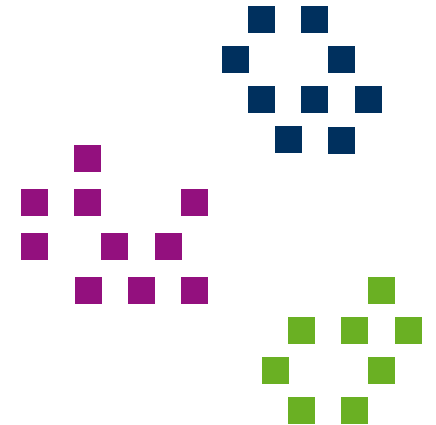
Partional Clustering Methods for Segmentation



Partional Clustering Methods

Partitioning methods are segmentation methods which divide the set of **objects N** on the basis of a fixed number of **classes s** in such a way that the computed segmentation or **partition K** minimizes a given **quality index b(K)**:

$$\min_{\in \wp(\wp(N))} \left\{ b(\mathbf{K}) : \quad = \{K_1, K_2, \dots, K_s\}, \quad \bigcup_{i=1}^s K_i = N, \quad K_i \cap K_j = \emptyset \right\}$$



The exchange principle

- (1) Choose start partition $\mathbf{K}^0 = \{K_1^0, \dots, K_s^0\}$ (start heuristic).
Choose $b(\mathbf{K}^0)$.
- (2) Search Object(s), so that a transfer reduces b .
- (3) Change Object(s) from the current to the best new class.
- (4) Repeat (2) and (3) until no other change is possible.
 \Rightarrow **local optimum** found

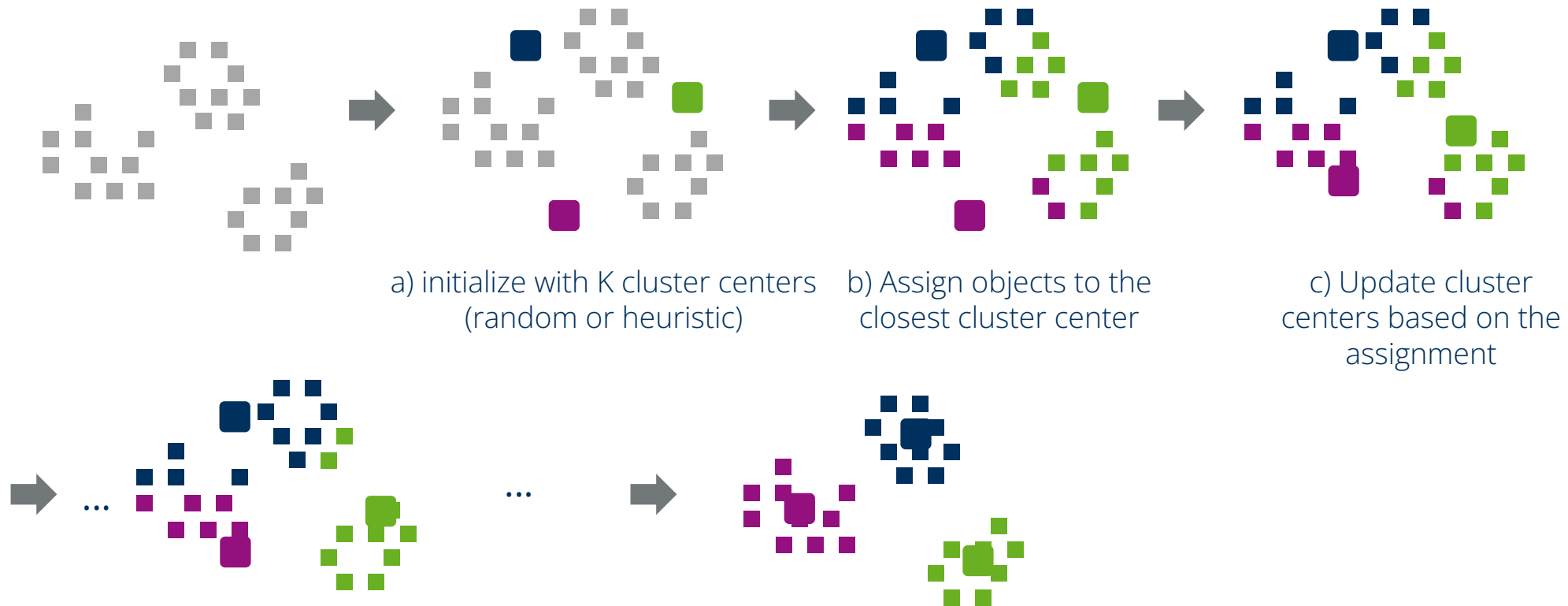
- The procedures break off after a finite number of steps.
- The procedures usually reach only a **suboptimum** (global optima are usually reached only if several objects can be exchanged simultaneously, taking into account all exchange possibilities).
- The result depends i.a. on the selected start partition (use several start partitions).

Error Minimization Algorithms

e.g. K-means

Partitions the data into K clusters represented by their centers or means.

Example: 2-dimensional, K=3

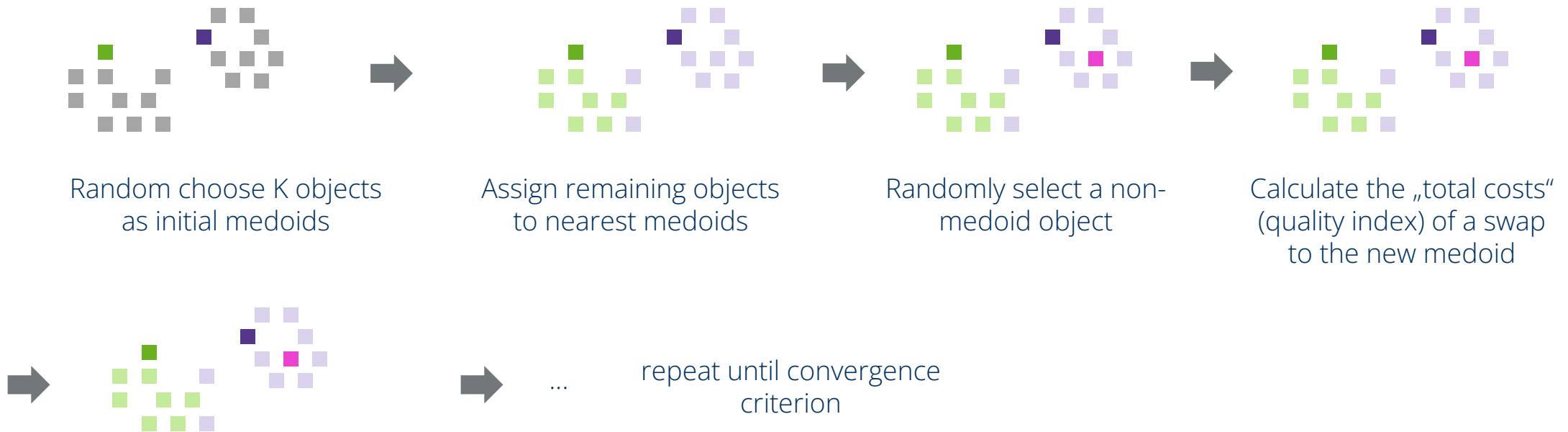


Error Minimization Algorithms

e.g. K-medoids or PAM (partition around medoids)

Find representative objects (medoids) in clusters

Example: 2-dimensional, K=2



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Hierarchical Clustering Methods for Segmentation

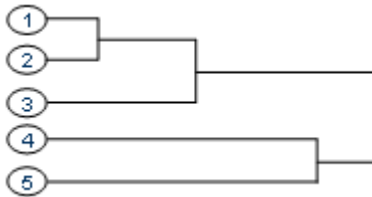


Hierarchical Clustering Methods

Segmentation methods that construct a sequence of partitions on the basis of a set of **objects N**.

Agglomerative Clustering

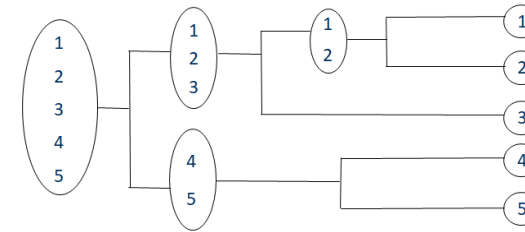
Each object initially represents a cluster of its own. Then clusters are successively merged until the desired cluster structure is obtained.



- Starting point are $n = |N|$ one-element classes.
- Successive transition to coarser decompositions
- Termination as soon as given criterion is fulfilled
- Low computation times, good practical suitability

Diversive Clustering

All objects initially belong to one cluster. Then the cluster is divided into sub-clusters, which are successively divided into their own sub-clusters. This process continues until the desired cluster structure is obtained.



- Starting point is the class of all objects.
- Successive transition to finer decompositions
- Termination as soon as given criterion is fulfilled

> Kaufman, L. and Rousseeuw, P.J. (1990) Partitioning around Medoids (Program PAM). In: Kaufman, L. and Rousseeuw, P.J., Eds., Finding Groups in Data: An Introduction to Cluster Analysis, John Wiley & Sons, Inc., Hoboken, 68-125.

Similarity measures

are based on the different **recalculation** of the interclass differences:

Single Linkage

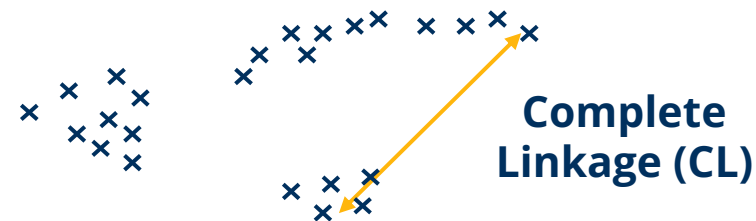
Nearest Neighbour



$$v(K,L) = \min_{i \in K, j \in L} d(i,j)$$

Complete Linkage

Furthest Neighbour



$$v(K,L) = \max_{i \in K, j \in L} d(i,j)$$

Average Linkage

Group Average



$$v(K,L) = \frac{1}{|K| \cdot |L|} \sum_{i \in K, j \in L} d(i,j)$$

Single Linkage

An Example

D	2	3	4	5
1	4,48	2,91	2,08	1,78
2		5,07	3,92	3,70
3			4,33	4,03
4				2,63

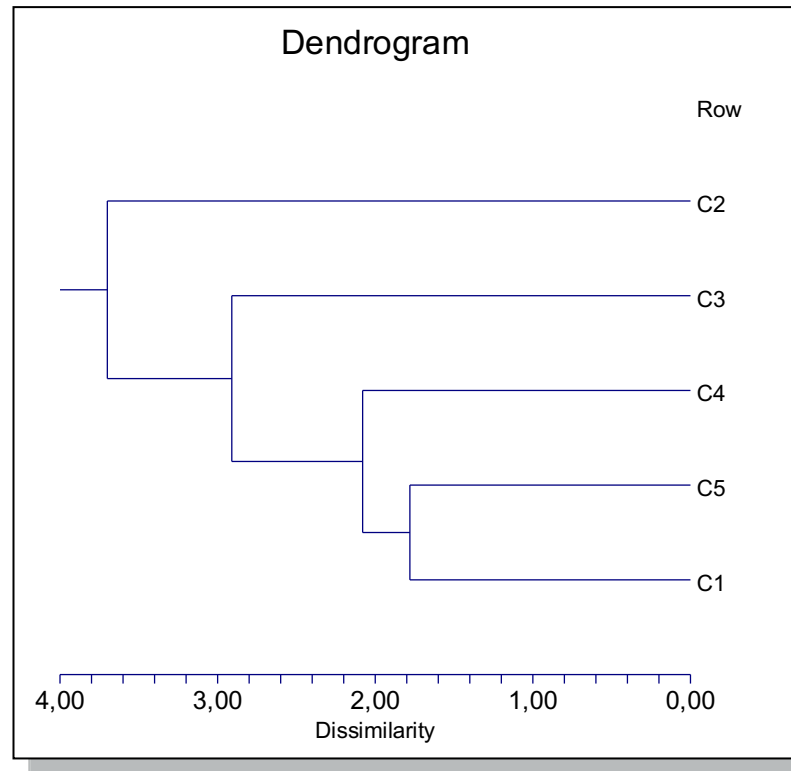
First fusion of objects 1 and 5 ($\min d_{ij} = 1.78$).

$\Rightarrow K^1 = \{\{1,5\}, \{2\}, \{3\}, \{4\}\}$

Fusion levels :

1.78 2.08 2.91 3.70

What we are looking for is a hierarchy with
 $v(i,j) = d(i,j)$ and $v(K,L) = \min_{i \in K, j \in L} d(i,j)$



Group Average Linkage

An Example

D	2	3	4	5
1	4,48	2,91	2,08	1,78
2		5,07	3,92	3,70
3			4,33	4,03
4				2,63

First fusion of objects 1 and 5 (min $d_{ij} = 1.78$).

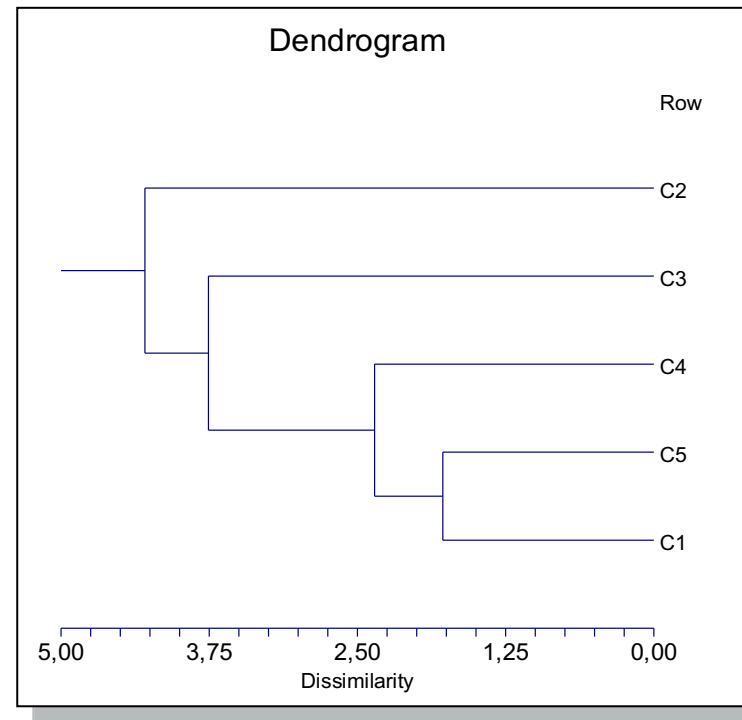
$\Rightarrow K^1 = \{\{1,5\}, \{2\}, \{3\}, \{4\}\}$

Fusion levels :

1.78 2.36 3.76 4.29

What we are looking for is a hierarchy with

$$v(i,j) = d(i,j) \text{ and } v(K,L) = \frac{1}{|K| \cdot |L|} \sum_{\substack{i \in K \\ j \in L}} d(i,j)$$



Complete Linkage

An Example

D	2	3	4	5
1	4,48	2,91	2,08	1,78
2		5,07	3,92	3,70
3			4,33	4,03
4				2,63

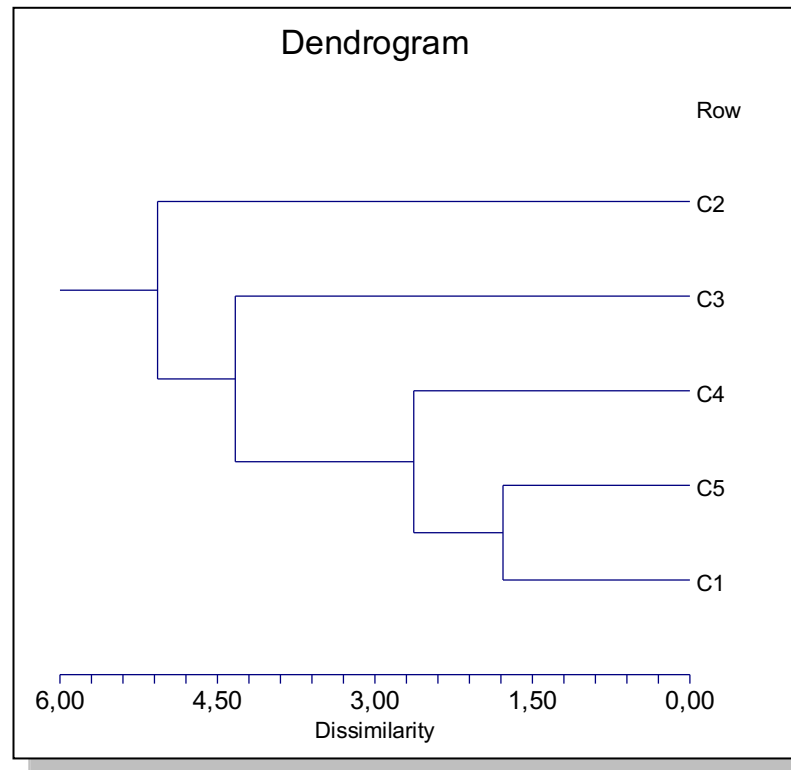
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Fusion levels :

1.78 2.63 4.33 5.07

What we are looking for is a hierarchy with
 $v(i,j) = d(i,j)$ and $v(K,L) = \max_{i \in K, j \in L} d(i,j)$



Interpretation of a Dendrogram

Abrupt changes in the value of the quality criterion allow conclusions to be drawn about the appropriate **number of classes (EB)**.

Similar objects are merged early, dissimilar objects later; **outliers** are assigned to a large cluster only at the end.

The class structure is

- **stable**, if different methods lead to similar results,
- **intensive**, if classes of comparable size are successively merged and
- **weak**, if successively only neighboring single objects are added.

Assessment of a hierarchy

To answer the question, which of the determined hierarchies "best" reproduces the given **distance matrix D**, one first calculates the (ultrametric) **distance matrix D***, which can be uniquely determined from the dendrograph, according to

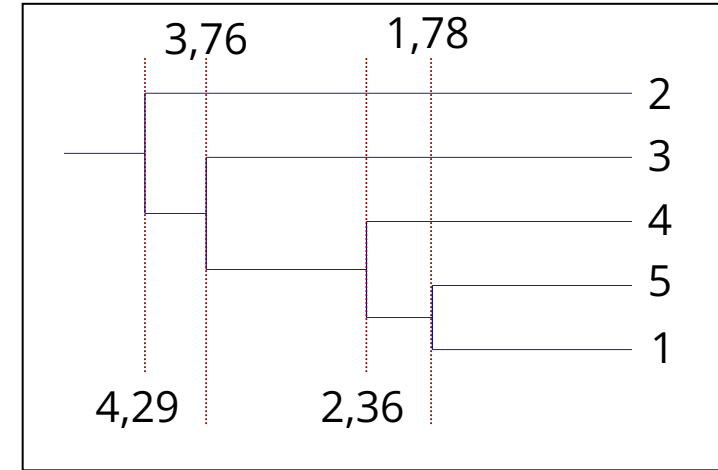
$$d^*(i,j) = \min_{\substack{i \in K, j \in L \\ K, L \in \mathcal{K}}} v(K,L)$$

and compares this matrix suitably with the original distance matrix D.

Distance matrix D^* of the average linkage

An Example

D	2	3	4	5
1	4,48	2,91	2,08	1,78
2		5,07	3,92	3,70
3			4,33	4,03
4				2,63



D^*	2	3	4	5
1	4,29	3,76	2,36	1,78
2		4,29	4,29	4,29
3			3,76	3,76
4				2,36

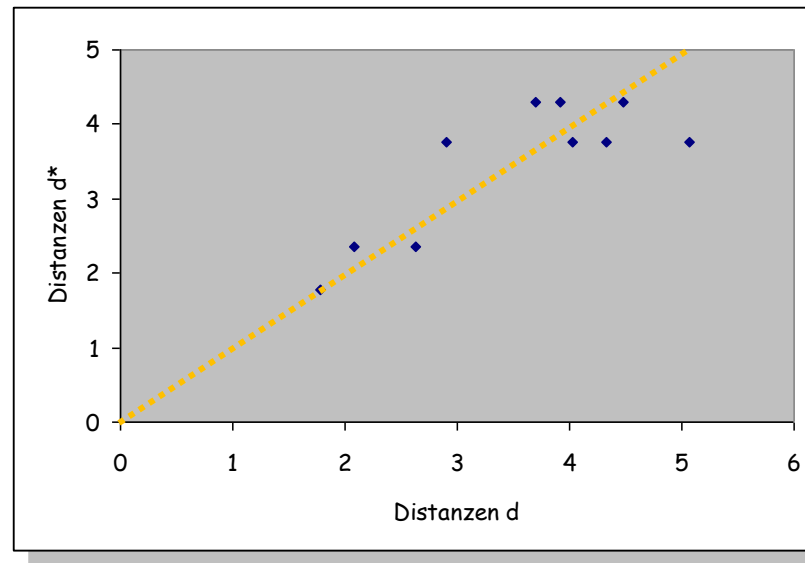


Calculation of the dissimilarity
using the **dendrogram**

The Shepard-Diagram

The simplest way to evaluate the different distance matrices D and D^* is the so-called Shepard diagram, in which the true **distances d** and the **calculated distances d^*** are compared in a **coordinate system**.

Example: Average Linkage



Variance-Accounted-For-Criteria

The **VAF criterion** can be calculated to assess the loss of information in procedures that explicitly use distances:

$$\text{VAF} = 1 - \frac{\sum_{i=2}^n \sum_{j=1}^{i-1} (d(i,j) - d^*(i,j))^2}{\sum_{i=2}^n \sum_{j=1}^{i-1} (d(i,j) - \bar{d})^2} \quad \text{mit} \quad \bar{d} = \frac{2}{n(n-1)} \sum_{i=2}^n \sum_{j=1}^{i-1} d(i,j)$$

For procedures that use variance to measure heterogeneity, the VAF cannot be meaningfully interpreted.

A value close to 1 is desired.

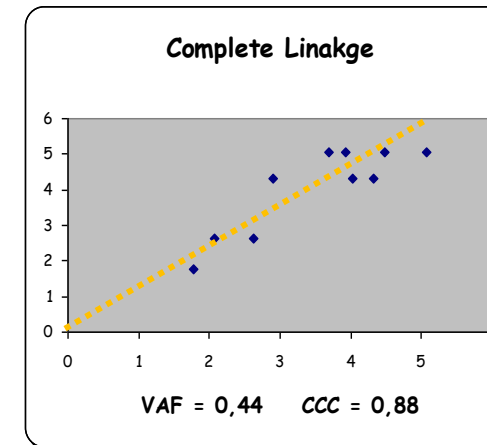
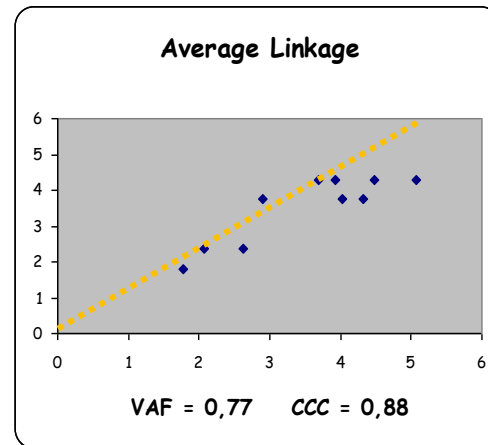
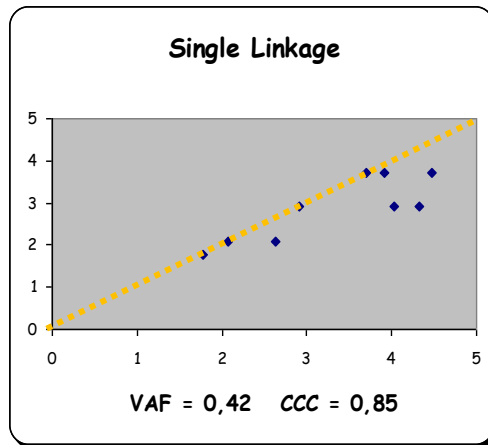
Cophenetic correlation coefficient

To assess the existence of a linear relationship between the true distances d and the calculated distances d^* according to:

$$CCC = \frac{\sum_{i=2}^n \sum_{j=1}^{i-1} (d(i,j) - \bar{d})(d^*(i,j) - \bar{d}^*)}{\sqrt{\sum_{i=2}^n \sum_{j=1}^{i-1} (d(i,j) - \bar{d})^2 \sum_{i=2}^n \sum_{j=1}^{i-1} (d^*(i,j) - \bar{d}^*)^2}} \quad \text{mit}$$
$$\bar{d} = \frac{2}{n(n-1)} \sum_{i=2}^n \sum_{j=1}^{i-1} d(i,j)$$
$$\bar{d}^* = \frac{2}{n(n-1)} \sum_{i=2}^n \sum_{j=1}^{i-1} d^*(i,j)$$

Values close to 1 indicate a small loss of information.

Example: Assessment of a hierarchy



It can be seen that the **Average Linkage** solution is judged the best.

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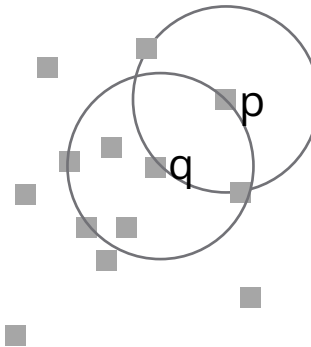
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Density Based Clustering Methods for Segmentation



Density Based Clustering Methods

Designed for discovering clusters of arbitrary shape. It is also used to handle noise in the data clusters



ϵ ... maximum radius of the neighborhood

MinPts ... minimum number of points in an ϵ -neighborhood

„p is directly *density-reachable* from q“

Idea: continue growing a given cluster as long as the density (number of objects or data points) in the neighborhood exceeds some threshold. The neighborhood within a radius ϵ has to contain at least a minimum number of objects. It needs density parameters as a termination condition.

> Ester, M., Kriegel, H.P., Sander, J. and Xu, X. (1996) A Density-Based Algorithm for Discovering Clusters in Large Spatial Databases with Noise. Proceedings of the 2nd International Conference on Knowledge Discovery and Data mining, 226-231

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Thank you for your attention

