#### **Cluster analysis**

Petr Ocelík

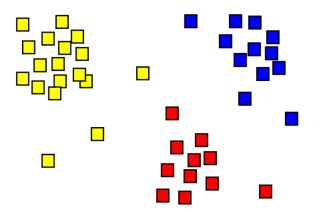
MVPd002 Quantitative Research in International and European Politics

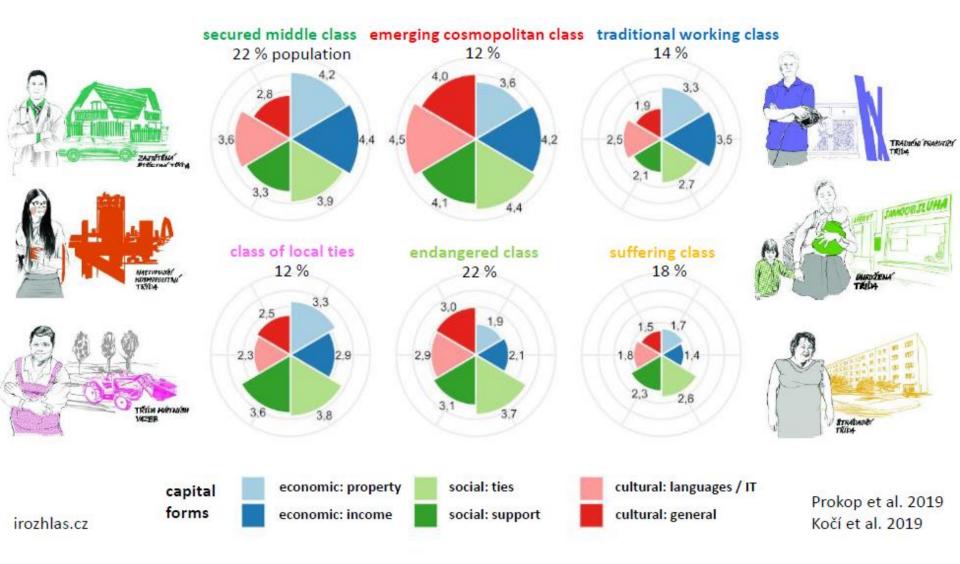
# Plan for today

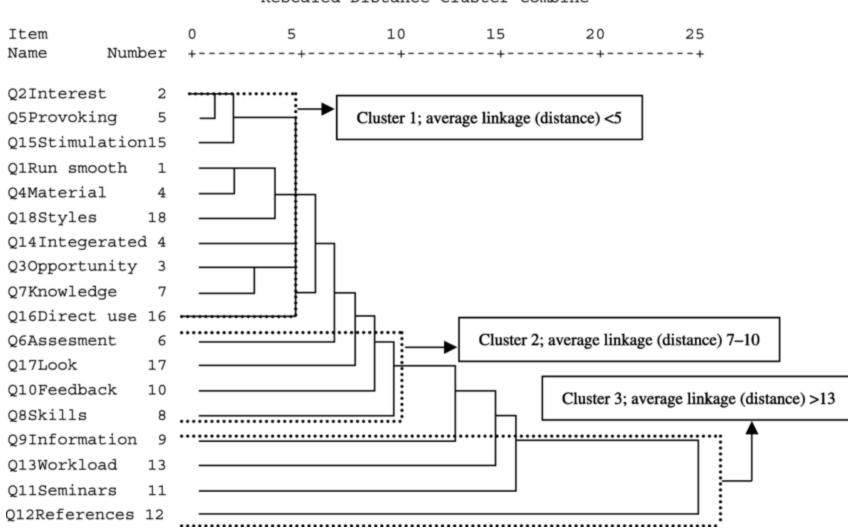
- Intuition
- Cluster analysis step-by-step
- Exercise

## Cluster analysis

- Data reduction technique
- **Cluster:** a grouping of similar objects
- Basic idea: identifying groups of mutually **similar objects** based on particular variable(s)
- Unsupervised technique





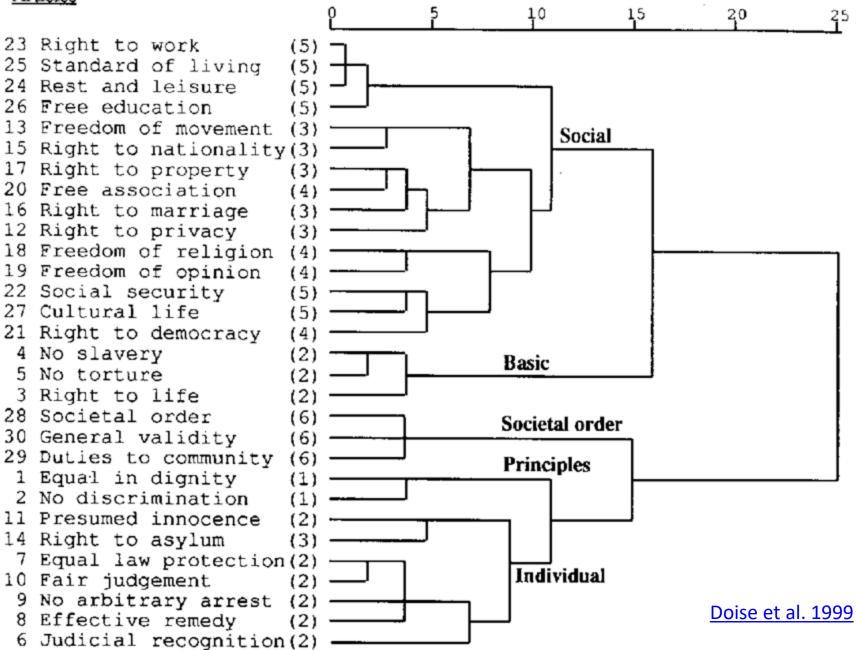


Rescaled Distance Cluster Combine

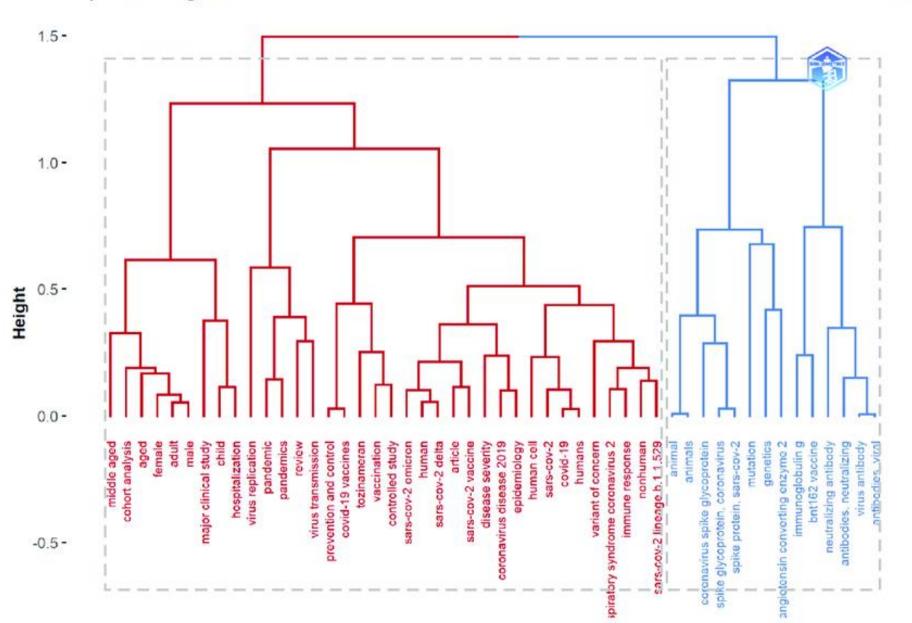
Ansari & Oskrochi 2006

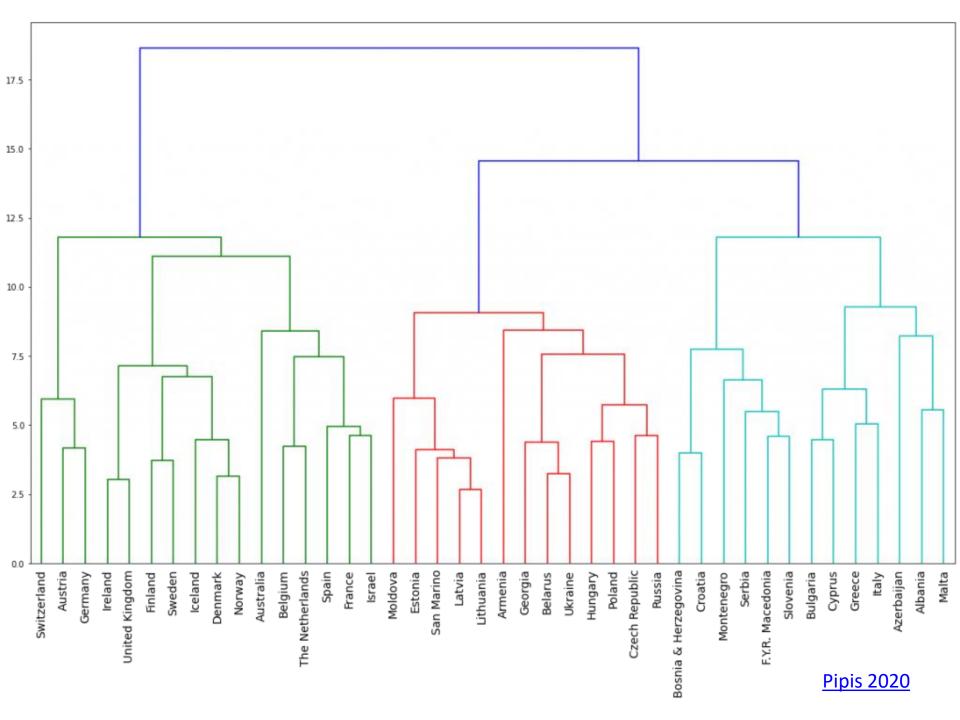
#### Rescaled dissimilarity coefficient

#### <u>Articles</u>



#### **Topic Dendrogram**





### Cluster analysis: process

- 1. Sampling and data collection
- 2. Similarity measures
- 3. Clustering methods
- 4. Cluster solution interpretation
- 5. Cluster solution diagnostics

# 1. Sampling and data collection

- What is a **target population**?
- What is the **level of analysis**?
- What is the **unit of observation**?
- What **set of variables** are we interested in?
- Practical considerations

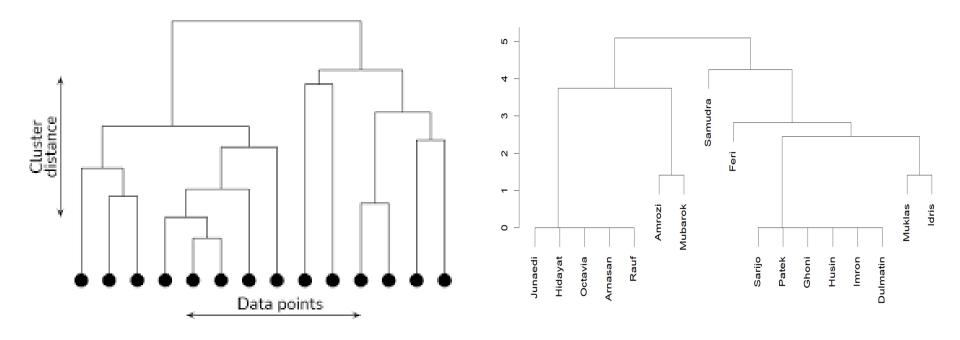
# 2. Similarity measures

- (Dis)similarity of objects is quantified by using **similarity measures**
- Choice of similarity measure needs to consider:
- 1. level of measurement: categorical vs continuous
- 2. data dimensionality: number of variables (vars)
- **3.** scale sensitivity: small vs large data, vars scales
- We distinguish between **association-based** and **distance-based** similarity measures (not exhaustive)

# 2. Similarity measures: representations

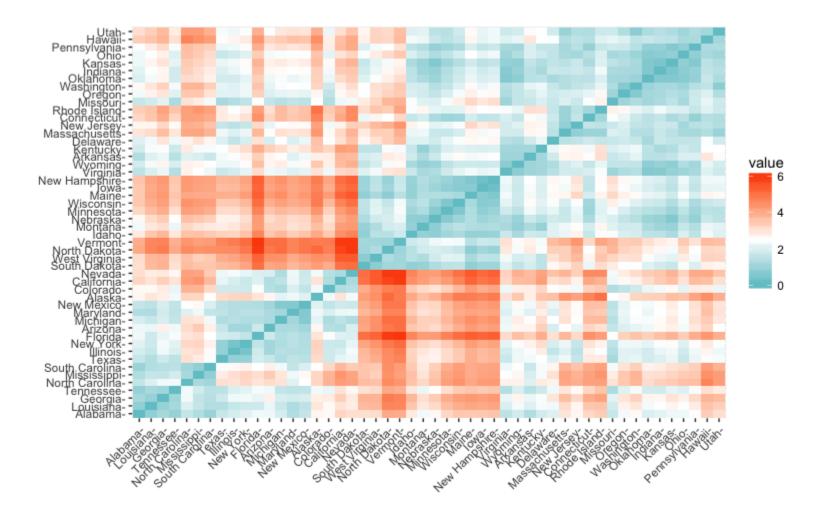
- The degree of (dis)similarity can be captured numerically or **graphically**.
- Dendrogram
- Heatmap
- Cluster profile

# Dendrogram



<u>Pai 2021</u>

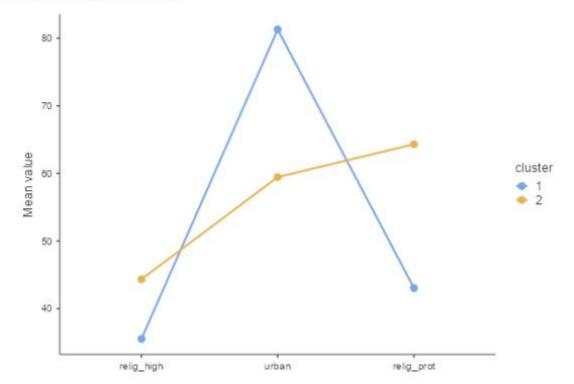
#### Heatmap



UC Business 2024

## Cluster profile

#### Plot of means across clusters



# 2.1 Pearson's coefficient

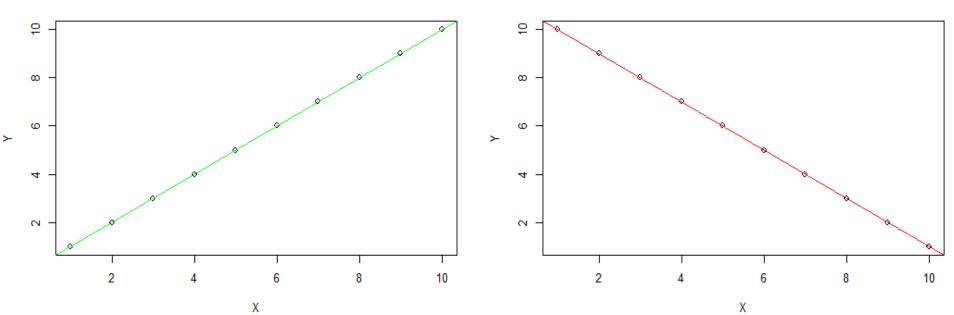
 Pearson's r measures the existence, strength and direction of the linear relationship between two variables

measurement level	number of values	range	coefficient
continuous-continuous	many-many	<-1, 1>	Pearson's r
			Soukup et al. 2022

• Not suitable for heterogeneous data and/or nonlinear relationships between variables

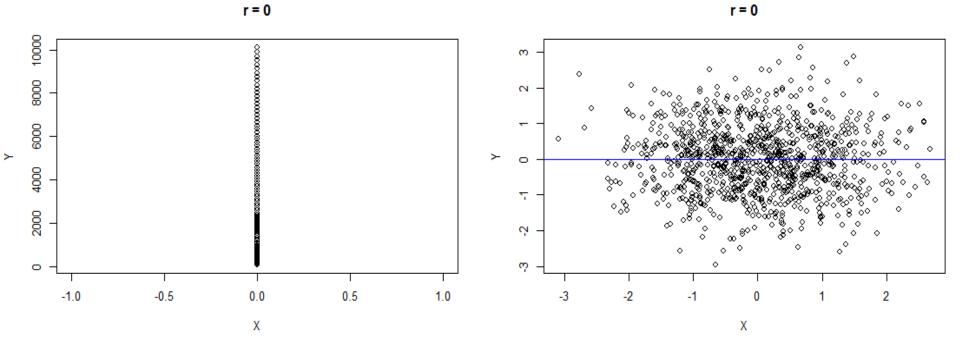




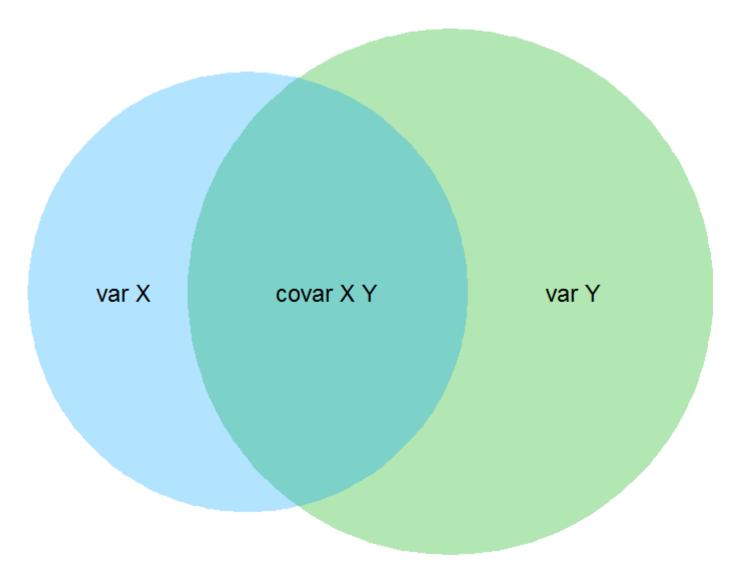


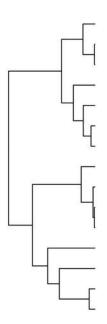
http://guessthecorrelation.com/

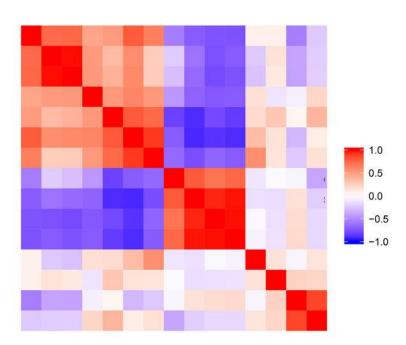
r = 0

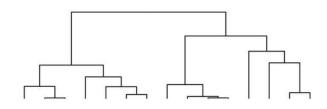


• r = covariance / combined total variance



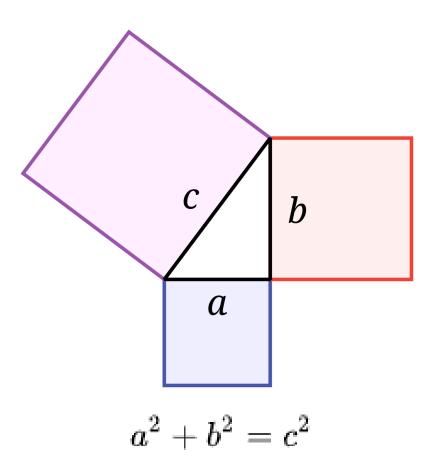






## 2.2 Euclidean distance

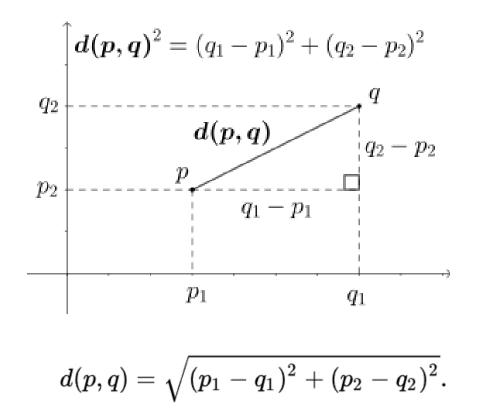
• Do you recall this?



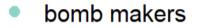
wikimedia commons

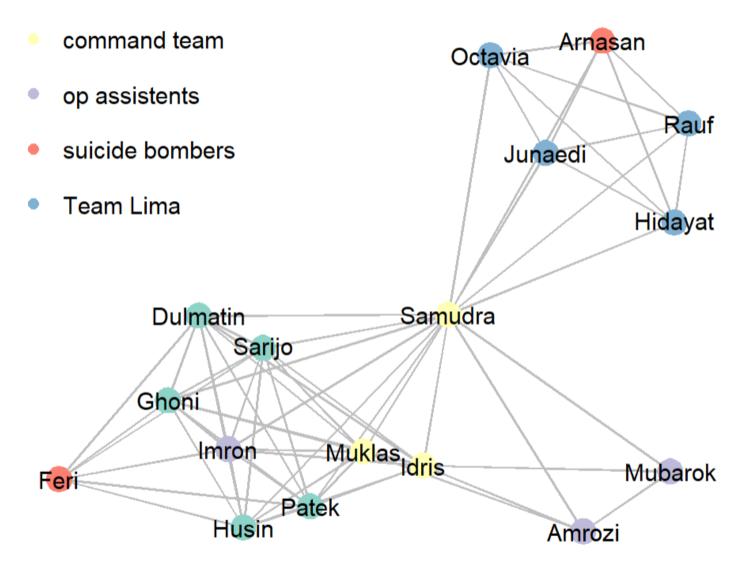
# 2.2 Euclidean distance

- Euclidean distance (d) is a generalization of the Pythagorean theorem
- d(p, q), of two objects (p, q) equals the length of the straight line between them



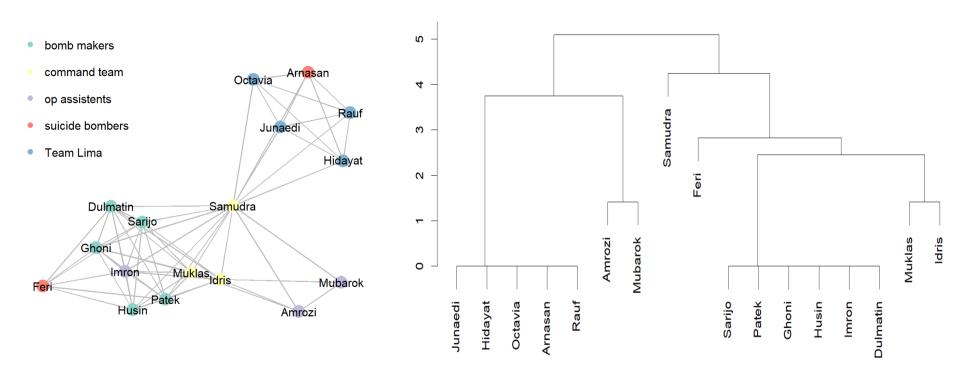
wikimedia commons





Koschade 2007

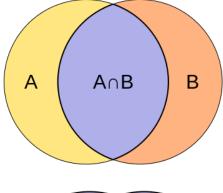
### 2.2 Euclidean distance



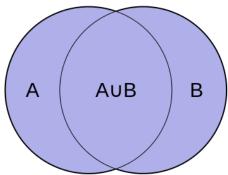
# 2.3 Jaccard's coefficient

measurement level	number of values	range	coefficient
categorical-categorical	binary	<0, 1>	Jaccard's

		sample B	
		present	absent
sample A	present	A ∩ B	A – B
	absent	B – A	∉A∪B

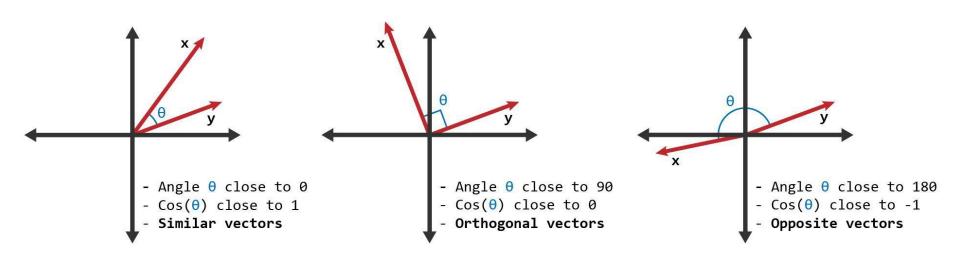


- J = the size of the intersection (A ∩ B) by the size of the union (A + B = A ∪ B) of the samples
- $J = A \cap B / (A \cup B)$
- Does not account for observations missing in both samples (∉ A ∪ B)



wikimedia commons

#### 2.4 Cosine distance



Karabiber 2024

# 2.4 Cosine distance (CD)

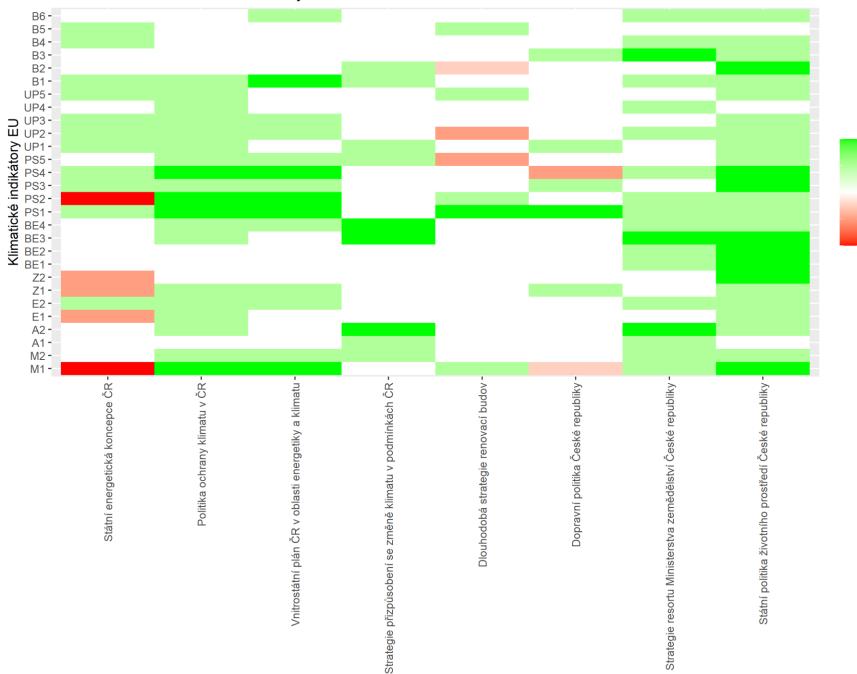
	doc_A	doc_B
geopolitics	4	0
climate change	0	1
Brexit	0	12
Euro	3	9
sovereignty	5	2

$$cosine \ similarity = |A||B|cos\theta = \frac{A \cdot B}{|A||B|} = \frac{\sum_{i}^{n} A_{i}B_{i}}{\sqrt{\sum_{i}^{n} A_{i}^{2}} \sqrt{\sum_{i}^{n} B_{i}^{2}}}$$

• CS = 
$$\frac{(4*0+0*1+0*12+3*9+5*2)}{\sqrt{(16+0+0+9+2)}*\sqrt{(0+1+144+81+4)}} = 0.345$$

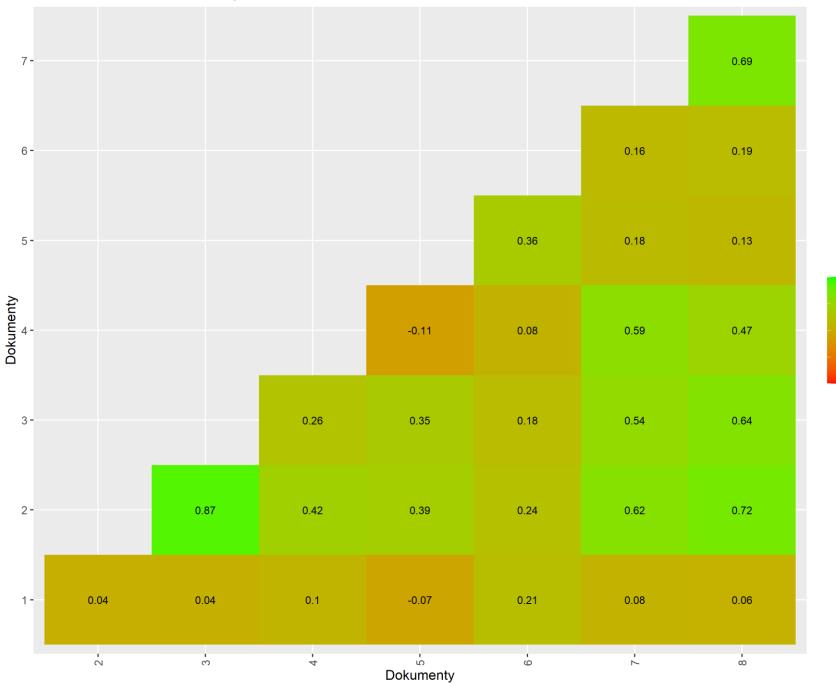
• CD = 1 - 0.345 = 0.655

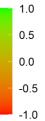
Koherence dokumentů s indikátory EU



2 1 0 -1 -2

Kosinová vzdálenost koncepčních dokumentů





# 2. Similarity measures: summary

- Cosine and Euclidean distances appropriate for higherdimensional data
- Cosine distance used for text-based data
- Euclidean distances and Jaccard coefficient used for network data
- Pearson coefficient appropriate for continuous data and linear relationships
- There are many more measures of similarity

### Cluster analysis: process

- 1. Sampling
- 2. Similarity measure
- 3. Clustering method
- 4. Cluster solution interpretation
- 5. Cluster solution diagnosis

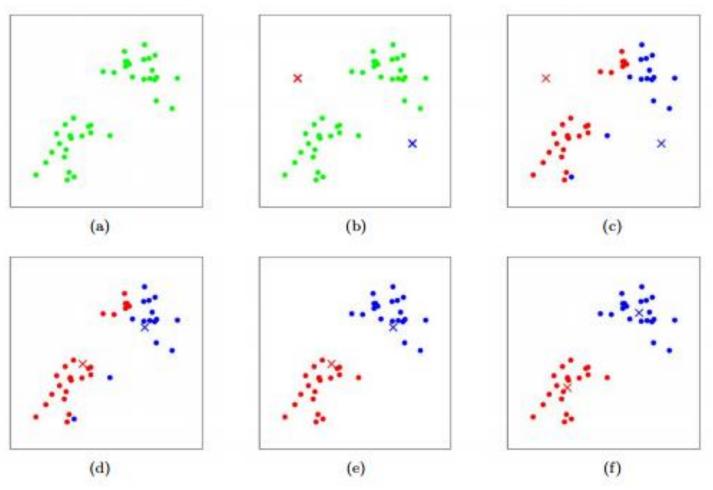
# 3. Clustering methods

- After the (dis)similarities between objects are calculated, we need to select a particular clustering method that partitions (clusters) the data according specific rules
- There are several clustering approaches, k-means clustering and hierarchical clustering belong to the most common

# 3.1 k-means clustering

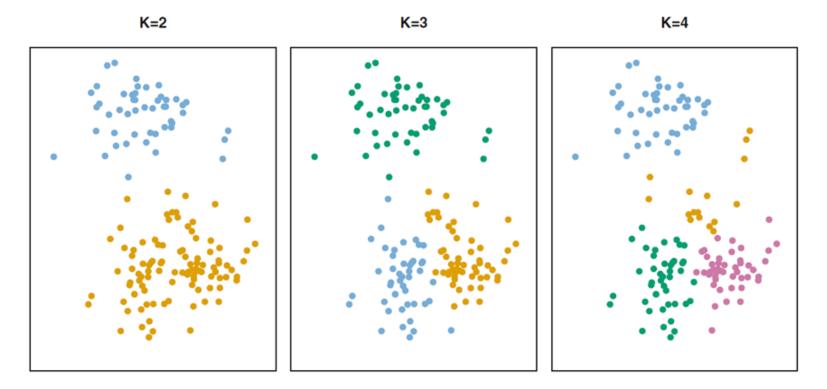
- **k-means clustering** partitions the data into *k* clusters based on the mean distances in each cluster
- 1. The number of clusters *k* needs to be **pre-selected**
- 2. The algorithm starts randomly assign cluster centers (centroids)
- 3. Each object is assigned to the nearest centroid based on a particular similarity measure
- 4. Within the clusters, **the centroids are updated** based on the mean similarity of the objects classified to the respective cluster
- 5. The steps 3-4 repeat until the centroids no longer change  $\rightarrow$  solution

#### 2-cluster solution



Jordan 2012

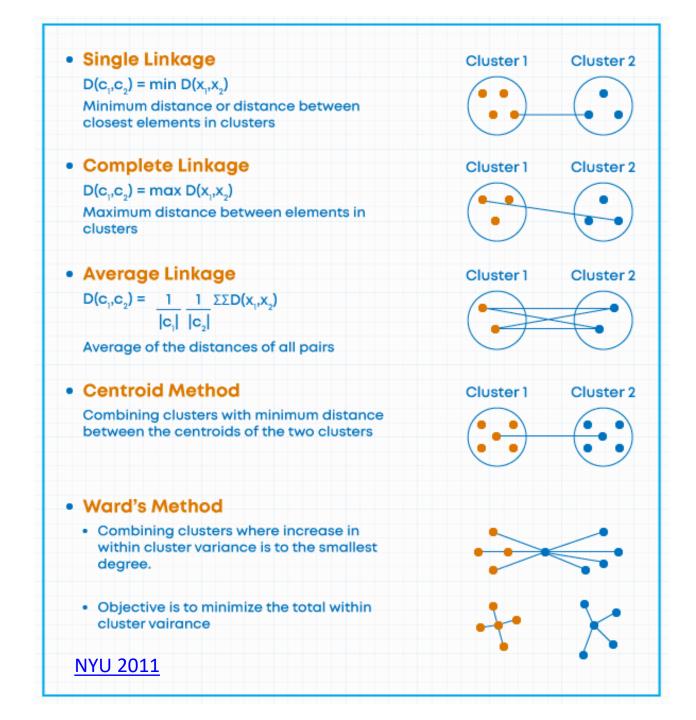
### Determining k

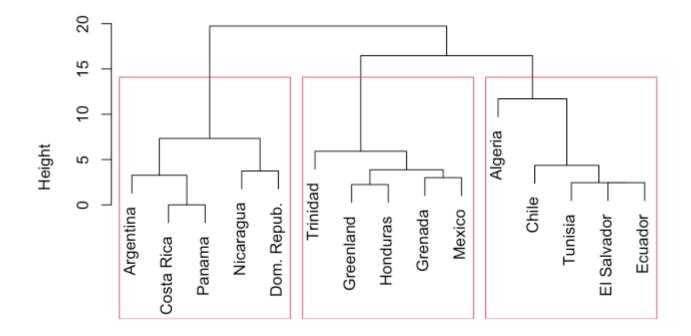


A simulated data set with 150 observations in two-dimensional space. Each figure show the results of applying K-means clustering with different values of K, the number of clusters. The color of each observation indicates the cluster to which it was assigned using the K-means clustering algorithm. Note that there is no ordering of the clusters, so the cluster coloring is arbitrary. These cluster labels were not used in clustering; instead, they are the outputs of the clustering procedure.

# 3.1 Hierarchical clustering

- Hierarchical clustering can be performed in agglomerative and divisive sequence
- The number of clusters is **not pre-selected**
- The linkage method (how are clusters merged/split) needs to be defined
- 1. Treat each objects as a separate cluster (**agglomerative** sequence)
- 2. The average distances between the clusters are calculated (average linkage)
- 3. The clusters with the lowest average distance are merged
- 4. The steps 2-3 are repeated until there is only a single cluster
- 5. The process is represented by **dendrogram**
- 6. Considering substantive insights, the *k*-cluster solution is identified





<u>NYU 2011</u>

#### Cluster analysis: process

- 1. Sampling
- 2. Similarity measure
- 3. Clustering method
- 4. Cluster solution interpretation
- 5. Cluster solution diagnosis

# 4. Cluster solution interpretation

- Do the clusters have **prima facie** validity (**eyeballing** test)?
- **Substantive** and **theoretical insights** are vital to what extent the solution aligns with our expectations?
- Size of the clusters do clusters markedly differ in their size?
- **Outliers** are there any?
- Do clusters **reduce our data well**? → cluster **solution diagnostics**

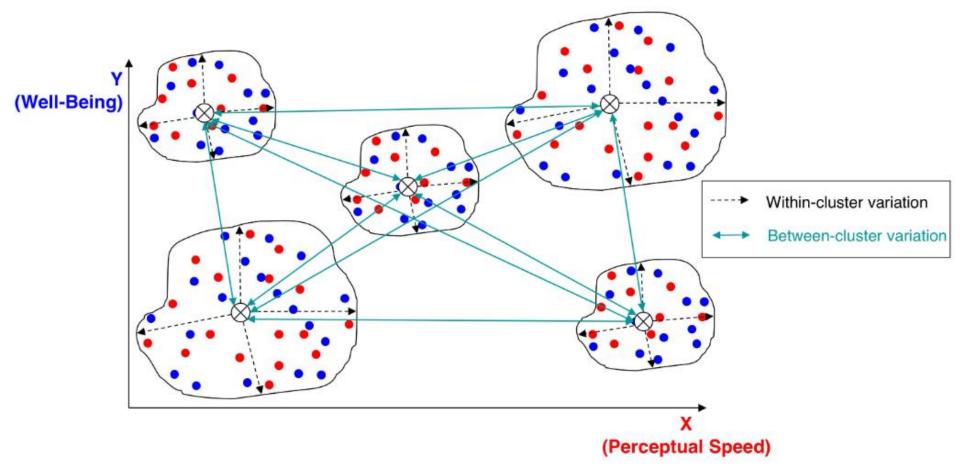
# 5. Cluster solution diagnostics

- Important to assess the **quality of our solution**
- Are the clusters internally **cohesive**?
- Are the clusters well **separated**?
- What is the optimal **number of clusters**?

# 5.1 Within/between sum of squares

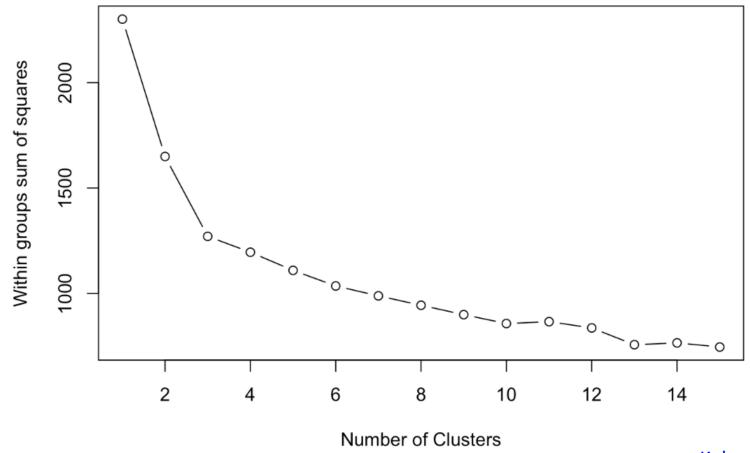
- The within-cluster sum of squares (WCSS) calculate the sum of squared distances between each object and its cluster centroid (k-means clustering approach)
- The **between-cluster sum of squares** (**BCSS**) measure the sum of squared distance between centroid of each cluster and the overall centroid of all objects
- The WCSS capture the cohesiveness of clusters, while BCSS measure the separation between clusters

#### 5.1 WCSS and BCSS



SSRI 2020

## **Elbow graph**



<u>Kulma 2017</u>

## 5.2 Silhouette score

• The **Silhouette score** ranges <-1, 1>; where high positive values indicate a good fit of the object within its own cluster, zero indicates a borderline position, and negative values indicate that the object is misclassified

• 
$$s = \frac{b-a}{\max(b-a)}$$
; *a* = average within-cluster d, *b* = average between-cluster d

- The Silhouette is calculated for each object and then average is taken to evaluate the cluster solution
- The Silhouette score can be applied both to the k-means and hierarchical clustering
- Rule of thumb: **s > 0.5** good solution; s < 0.25 bad solution

#### Exercise

## Exercise

- Open Jamovi and install **snowCluster** extension
- Load into Jamovi file "state.csv"
- Check variables relig\_prot, urban, and relig\_high in the codebook
- Describe the variables
- Apply (1) k-means clustering and (2) hierarchical clustering
- Interpret the results