# Clustering validation

Is there any real clustering? How good is it?

- Book: Chapter 6.9
- External material: Halkidi et al. (2002): Cluster Validity Methods: Part I. ACM SIGMOD Record 31(2): 40–45. https://doi.org/10.1145/565117.565124

# Three similar problems

- 1. Clustering tendency: is there any clustering in data presented with certain features?
- 2. Determining number of clusters (or other parameters)
- 3. Evaluating goodness of clustering
  - compare different methods
  - compare against classification

### All three depend on the clustering objective!

- assumptions on clusters (e.g., compactness, shape)
- separation between clusters

# Evaluating goodness of clustering

#### 1. Internal criteria

- validity indices, similar to objective functions
- do not work, if clustering had a different objective!
- can be used to i) evaluate a single clustering or ii) compare clusterings (as relative indices)

#### 2. External criteria

- compare clustering to a predefined classification
- classes may not reflect natural clusters

#### 3. Statistical hypothesis testing

 maybe the most sound approach, but computationally demanding

# Internal validity indices

- indices assume some clustering objective → reward methods with the same objective
  - even a good clustering can get a bad score if a different objective!
  - many indices assume/favor spherical or convex clusters
- best for comparing similar algorithms and tuning parameters
- Some popular indices:
  - Average silhouette
  - Calinski-Harabasz index
  - Davies-Bouldin index

#### Silhouette index

#### Silhouette of a point x is

$$S(\mathbf{x}) = \begin{cases} 0 & \text{if } \mathbf{x} \text{ a cluster of its own} \\ \frac{b-a}{\max\{a,b\}} & \text{otherwise} \end{cases}$$

$$a = avg\{d(\mathbf{x}, \mathbf{y}) \mid \mathbf{x} \in C, \mathbf{y} \in C\}$$
  
$$b = \min_{q} avg\{d(\mathbf{x}, \mathbf{y}) \mid \mathbf{x} \in C, \mathbf{y} \in C_q, C \neq C_q\}$$

 $\approx$  how closely x matches its own cluster and how loosely the neighbouring cluster

- $S(\mathbf{x}) \in [-1, 1]$ , high values good
- Average silhouette describes goodness of entire clustering
- flexible: any distance function *d*

# Example: Silhouette of points



# What negative values mean?

$$S(\mathbf{x}) = \begin{cases} 0 & \text{if singleton} \\ \frac{b-a}{\max\{a,b\}} & \text{otherwise} \end{cases}$$

$$\begin{aligned} a &= avg\{d(\mathbf{x}, \mathbf{y}) \mid \mathbf{x} \in C, \mathbf{y} \in C\} \\ b &= \min_{q} avg\{d(\mathbf{x}, \mathbf{y}) \mid \mathbf{x} \in C, \mathbf{y} \in C\} \\ C_q, C \neq C_q \} \end{aligned}$$

image source http://www.sthda.com/
english/wiki/wiki.php?id\_contents=7952

#### Calinski-Harabasz index

$$S_{CH} = \frac{(n-K)B}{(K-1)W}$$

- between-cluster variance  $B = \sum_{i=1}^{K} |C_i| L_2^2(\mathbf{c}_i, \mathbf{m})$ , where  $\mathbf{m}$  is the mean of the whole data
- within-cluster variance  $W = \sum_{i=1}^{K} \sum_{\mathbf{x} \in C_i} L_2^2(\mathbf{x}, \mathbf{c}_i)$
- requires  $K \ge 2$
- range  $[0, \infty[$ , high values good
- When could you get value 0?

# Calinski-Harabasz index (cont'd)

$$S_{CH} = \frac{(n-K)B}{(K-1)W} = \frac{(n-K)\sum_{i=1}^{K} |C_i| L_2^2(\mathbf{c}_i, \mathbf{m})}{(K-1)\sum_{i=1}^{K} \sum_{\mathbf{x} \in C_i} L_2^2(\mathbf{x}, \mathbf{c}_i)}$$

**Note:**  $W = SSE(\mathbf{C})$ . K-means criterion minimizes  $W \Rightarrow$  maximizes B, because

$$\sum_{\mathbf{x} \in \mathcal{D}} L_2^2(\mathbf{x}, \mathbf{m}) = \sum_{i=1}^K \sum_{\mathbf{x} \in C_i} L_2^2(\mathbf{x}, \mathbf{c}_i)^2 + \sum_{i=1}^K |C_i| L_2^2(\mathbf{c}_i, \mathbf{m})$$

 $\Rightarrow$   $S_{CH}$  favours especially K-means!

**Important**: need to use  $L_2$  in clustering!

## Davies-Bouldin index

$$S_{DB} = \frac{1}{K} \sum_{i=1}^{K} \max_{j \neq i} \frac{S_i + S_j}{D_{ij}} \quad \text{, where}$$

- $S_i = \left(\frac{1}{|C_i|} \sum_{\mathbf{x} \in C_i} L_p^q(\mathbf{x}, \mathbf{c}_i)\right)^{\frac{1}{q}}$  measures dispersion of  $C_i$ 
  - usually q = 2 (stdev of distances)
  - if q = 1, average distances
- $D_{ij} = L_p(\mathbf{c}_i, \mathbf{c}_j)$  measures separation between  $C_i$  and  $C_j$
- max: for each  $C_i$ , evaluate relation to most problematic  $C_j$
- possible to take avg instead of max

**Important**: use the same  $L_p$  as the clustering algorithm!

# Davies-Bouldin index (cont'd)

$$S_{DB} = \frac{1}{K} \sum_{i=1}^{K} \max_{j \neq i} \frac{S_i + S_j}{D_{ij}}$$
, where

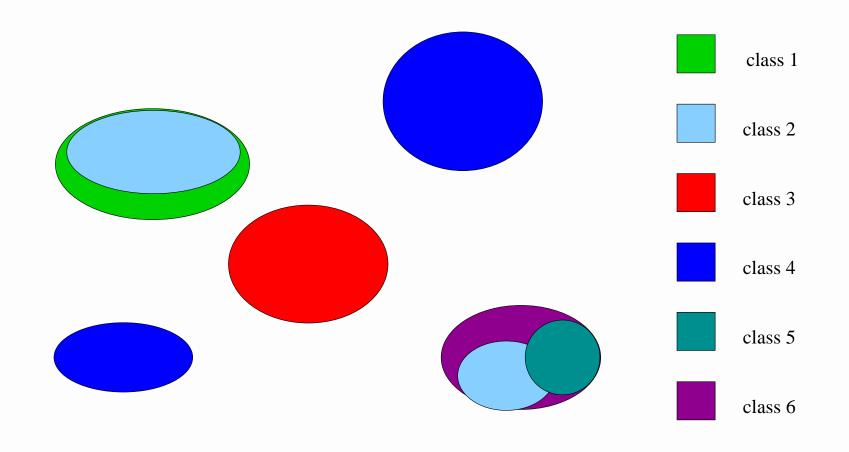
$$S_i = \left(\frac{1}{|C_i|} \sum_{\mathbf{x} \in C_i} L_p^q(\mathbf{x}, \mathbf{c}_i)\right)^{\frac{1}{q}} \text{ and } D_{ij} = L_p(\mathbf{c}_i, \mathbf{c}_j)$$

- range  $[0, \infty[$ , small values good
- When could you get value 0?

#### Possible strategies when $S_{DB}$ used to determine K:

- restrict number of singletons (e.g., 0 or a few)
- define  $S_i = a$  for some large a, when  $|C_i| = 1$

# External validation: Compare clustering against predefined classification



# A confusion matrix: clustering vs. classification

	Class 1	Class 2	Class 3	
Cluster 1	$n_{11}$	n <sub>12</sub>	n <sub>13</sub>	$m_1$
Cluster 2	n <sub>21</sub>	n <sub>22</sub>	n <sub>23</sub>	$m_2$
Cluster 3	n <sub>31</sub>	n <sub>32</sub>	n <sub>33</sub>	$m_3$
	$c_1$	c <sub>2</sub>	<i>c</i> <sub>3</sub>	n

image source Cunnigham https://slideplayer.com/slide/14318989/

#### External validation

Given clustering  $C_1, \ldots, C_K$  and classification  $D_1, \ldots, D_q$ . Many validation indices! E.g.,

purity

$$Pur(C) = \frac{1}{n} \sum_{i=1}^{K} \max_{j} |C_i \cap D_j|$$

- be careful! (increases with K)
- normalized mutual information NMI (robust, independent of K)
- Rand index

#### Normalized mutual information

Normalized mutual information by Strehl and Ghosh (2003):

$$NMI = \frac{I(C, D)}{\sqrt{H(C)H(D)}}$$

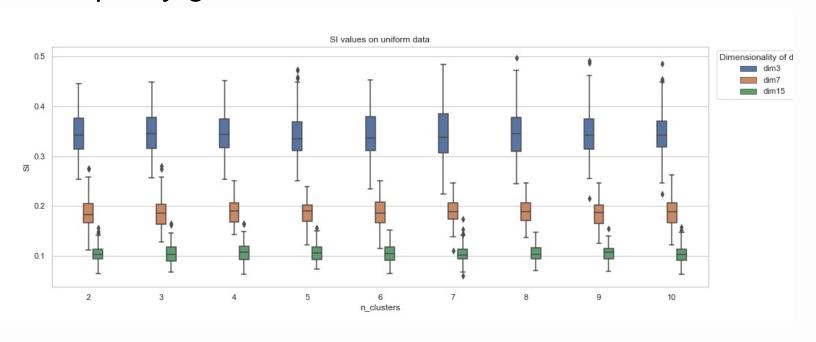
mutual information  $I = \sum_{C_i \in C} \sum_{D_j \in D} P(C_i, D_j) \log \frac{P(C_i, D_j)}{P(C_i)P(D_j)}$ entropy  $H(C) = -\sum_{C_i \in C} P(C_i) \log P(C_i)$ 

- + does not depend on the number of clusters
- many singleton clusters can cause problems

Note: Also other variants of normalized mutual information, give always equation and/or reference what you use!

# Statistical hypothesis testing: motivation

SI can be pretty good even for random data!



- each feature generated independently from uniform distribution
- 100 randomizations
- K-means repeated 100 times  $\rightarrow$  best result for each K

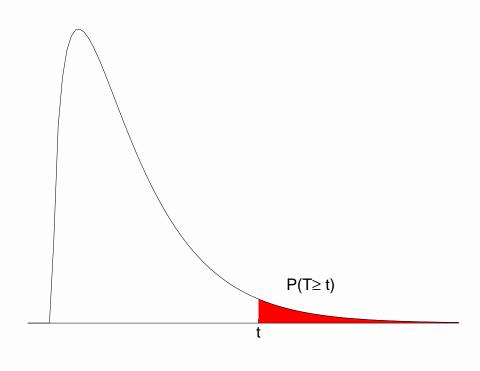
Experiment by Georgy Ananov for MDM 2023

# Statistical hypothesis testing

#### Procedure:

- 1. decide a null hypothesis  $H_0$  to test
  - describes the state where there isn't any clustering
  - e.g.,  $H_0$ : All sets of n locations in certain region are equally likely.
- 2. decide a test statistic *T* 
  - may be a validity index
- 3. What is the probability to obtain at least as good test statistic values as in data (where T = t) if  $H_0$  was true?

# Statistical hypothesis testing



Assume that large T value good

**Idea:** If  $P(T \ge t)$  very small  $\Rightarrow$  unlikely that the observed clustering had occurred by chance

•  $P(T \ge t)$  is the **p-value** that can be used as a significance measure

# Statistical hypothesis testing

Problem: How to evaluate p-value? (T's distribution seldom known!)

- often by Monte Carlo experiments (randomization tests):
  - generate random data sets fulfilling  $H_0$ , cluster them and evaluate T
  - p-value  $\approx$  proportion of random sets that obtained  $T \ge t$  (if large T good)
- computationally demanding (a lot of simulations!)
- many alternatives for  $H_0$ s and Ts

# Other evaluation: What the clustering reveals?

- Look at cluster sizes (e.g.,  $C_1$ : n-2 data points and  $C_2$ : 2 points likely outliers!)
- How do the clusters differ? (selected and external features)
  - e.g., rats clustered by body measurements (weight, tail and body length, organ weights)
  - 2 clusters: big and small rats
  - vs. 3 clusters:  $C_1$ : young or sick rats,  $C_2$ : pregnant or nursing females,  $C_3$ : other adults
- Are all clusters clear? (e.g.,  $C_1$  and  $C_3$  intermingled,  $C_2$  separate)

# Summary

- Remember validation, but be cautious!
  - even random data can produce clusterings, but they seldom pass validation
  - problem: indices biased or do not reflect the underlying clustering
  - try always more than one validation technique
- Objective, distance measure, clustering method and validation should match!

# Sources and further reading

- Halkidi et al. (2001): On clustering validation techniques, Journal of Intelligent Information Systems 17: 107–145. https://www.researchgate.net/ publication/2500099\_On\_Clustering\_Validation\_ Techniques
- Jain and Dubes (1988): Algorithms for clustering data,
   Ch 4.
- Gan, Ma, Wu (2007): Data clustering theory, algorithms, and applications, Ch 17, https://www.researchgate.net/publication/ 220694937\_Data\_Clustering\_Theory\_Algorithms\_and \_Applications

# Sources and further reading

 Vargha, Bergman, Takacs: Performing Cluster Analysis Within a Person-Oriented Context: Some Methods for Evaluating the Quality of Cluster Solutions. Journal of Person-Oriented Research, 2: 78-86, 2016.