

# Fundamentals of AI

## Clustering

### Assessment of clustering quality\*

\*Some materials in this lecture are used from

<https://towardsdatascience.com/unsupervised-machine-learning-clustering-analysis-d40f2b34ae7e>

<https://gdcoder.com/silhouette-analysis-vs-elbow-method-vs-davies-bouldin-index-selecting-the-optimal-number-of-clusters-for-kmeans-clustering/>

# Problem with clustering

- Any algorithm will deliver some clusters
- Some data are not naturally ‘clusterable’
- No clustering algorithm exists without parameters affecting the number of detected clusters
- We would like clusters *not to be* the results of statistical fluctuations in the point density
- We should always test how far the discovered structure in the data is from ‘random’ data
- Using built-in optimization criteria is not a good idea

# Basic questions

- Did we capture any *non-random* structure in the data?
- Did we choose the parameters (e.g., number of clusters) in the most optimal way?
- Do our clusters make sense in the application domain?

# Standard protocol for testing cluster validity

1. Identify the clustering structure and validation type
2. Define a validation index
3. Define a null hypothesis of no structure (for example, *all the locations of data points in the specific regions are equally likely*)
4. Establish the baseline distribution under the null hypothesis (for example, by bootstrapping, resampling or permutations)
5. Calculate the index
6. Test the hypothesis of no structure

# ‘Supervised’ vs ‘Unsupervised’ clustering validation

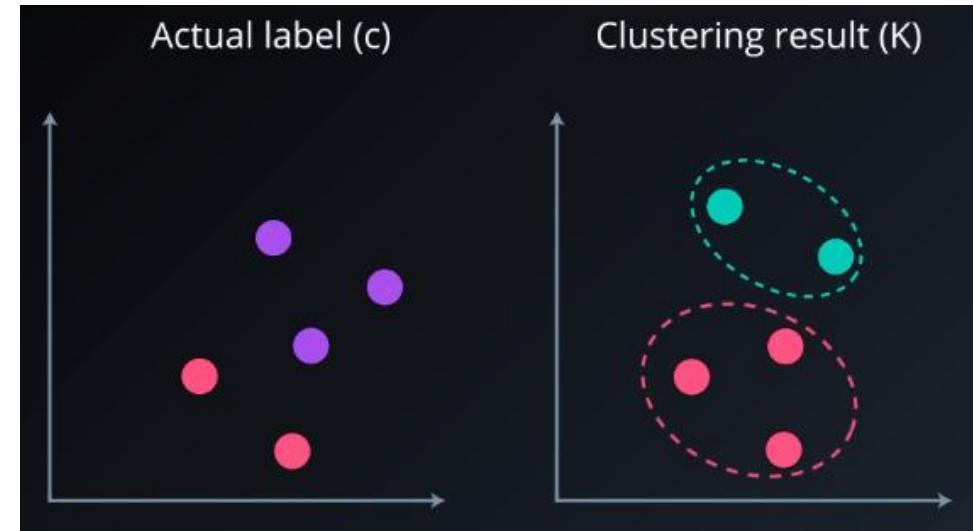
- Supervised: How similar is partitioning of the data into *clusters* and *classes*?
- Unsupervised: How ‘compact’ are the obtained clusters?

# Rand Index (accuracy of clustering with respect to 'ground truth')

- Rand index is the most basic measure of the percentage of correct decisions made by the algorithm
- **a**: is the number of points that are in the same cluster both in C and in K
- **b**: is the number of points that are in the different cluster both in C and in K.
- **n** : is the total number of samples

$$\text{Rand Index} = \frac{a + b}{n(n-1)/2}$$

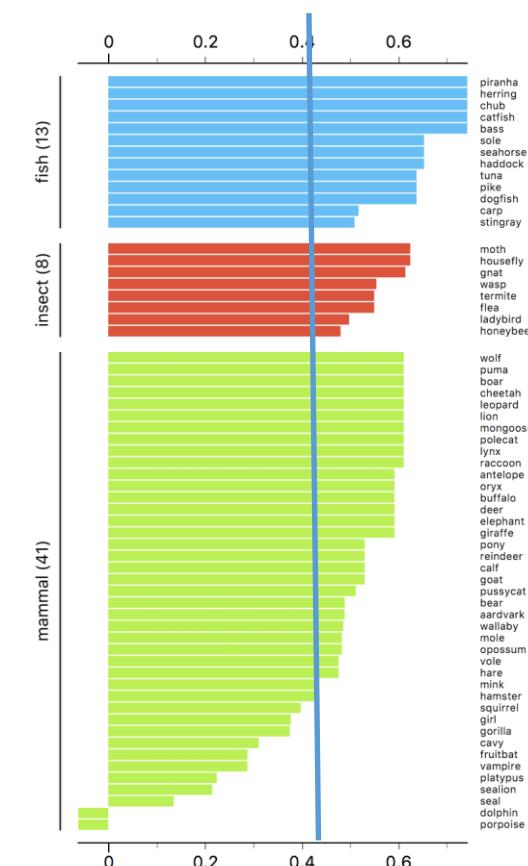
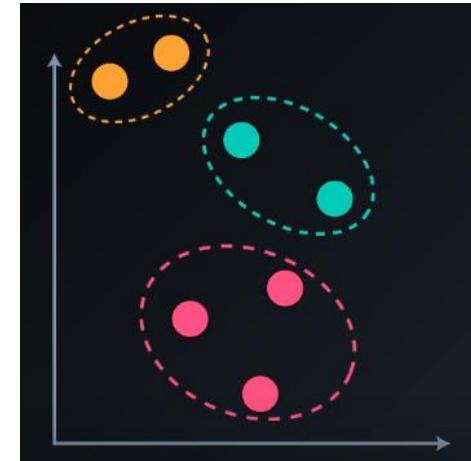
Adjusted Rand Index (ARI) corrects for the case of random label reshuffling



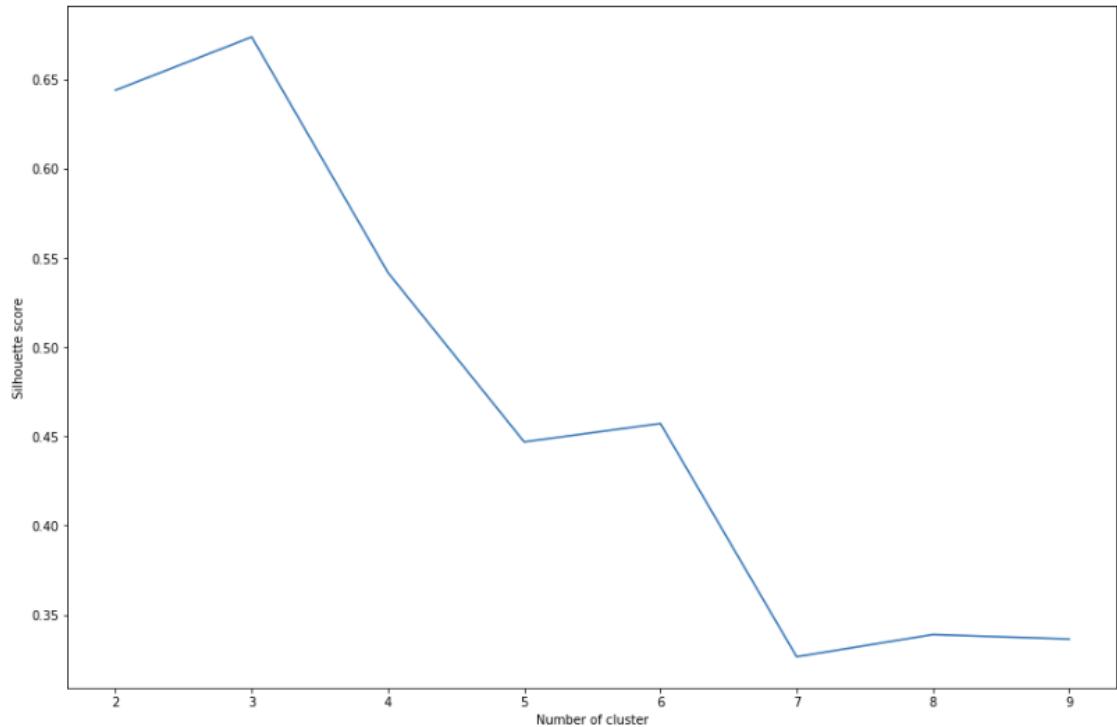
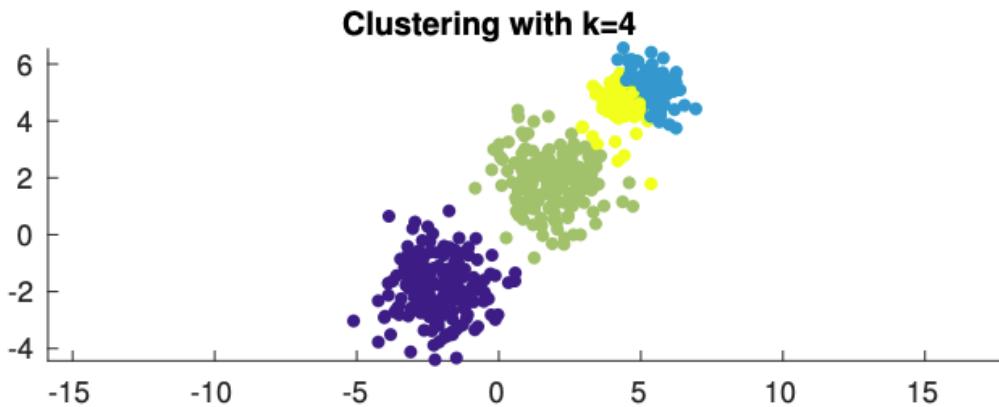
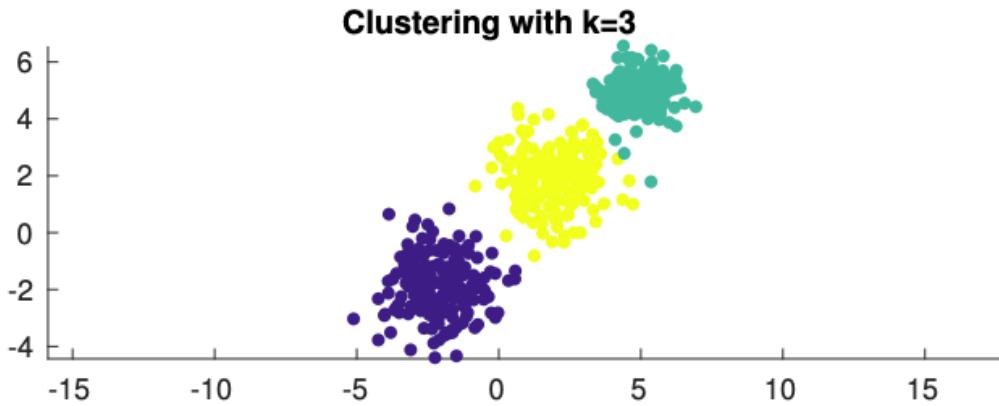
$$\text{ARI} = \frac{\text{RI} - \text{Expected Index}}{\text{Max(RI)} - \text{Expected Index}}$$

# Silhouette

- Suitable for K-means and hierarchical clustering
- $a$  = average distance to other sample in the same cluster
- $b$  = average distance to other sample in closest neighbouring cluster
- Silhouette for a point  $i$ :  $S_i = (a_i - b_i) / \max(b_i, a_i)$
- Global silhouette coefficient = average of the sum of  $S_i$  for each point



# Using silhouette to determine the right number of clusters



# Other indices (several tens)

- Calinski-Harabasz index
- Davies-Bouldin index
- Dunn index
- Density-Based Clustering Validation index

# Consensus clustering and resampling

must do for small number of points!

- How to compare two clustering results?
- Way to find ‘averaged’ clustering, if there are several clustering results available
- Typical application for cluster validation: remove k% of data points, cluster, repeat (resampling)
- Let us generate  $H$  datasets, then for each  $h = 1 \dots H$
- Connectivity matrix

$$M^h(i, j) = \begin{cases} 1, & \text{if points } i \text{ and } j \text{ belong to the same cluster} \\ 0, & \text{otherwise} \end{cases}$$

- Indicator matrix

$$I^h(i, j) = \begin{cases} 1, & \text{if } i \text{ and } j \text{ are in the same dataset} \\ 0, & \text{otherwise} \end{cases}$$

- Consensus matrix

$$C(i, j) = \left( \frac{\sum_{h=1}^H M^h(i, j)}{\sum_{h=1}^H I^h(i, j)} \right)$$

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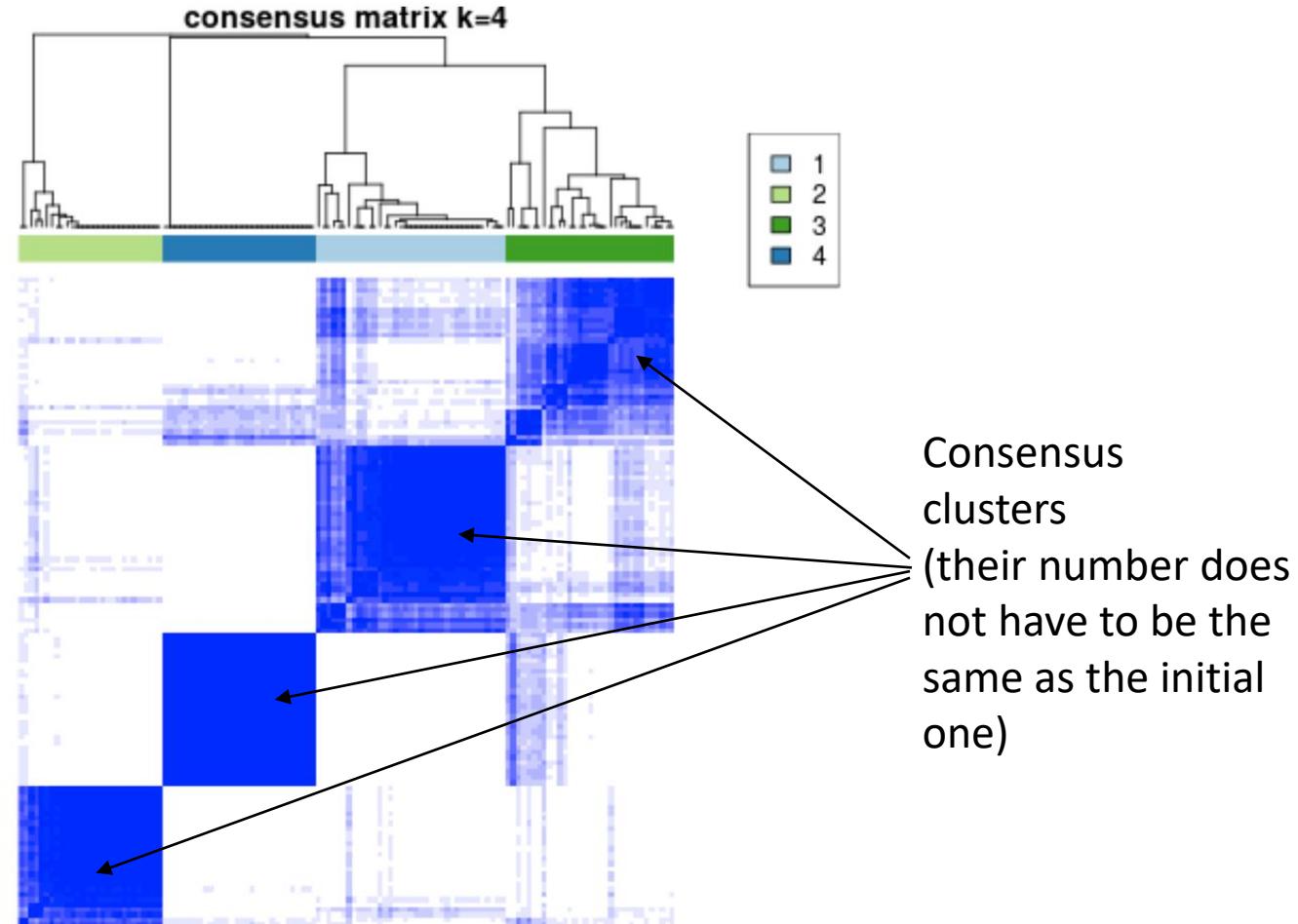
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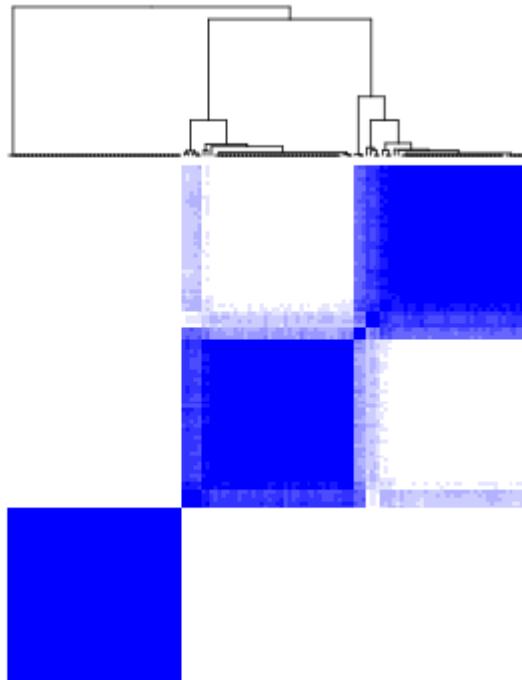
# Consensus clustering and resampling consensus clusters

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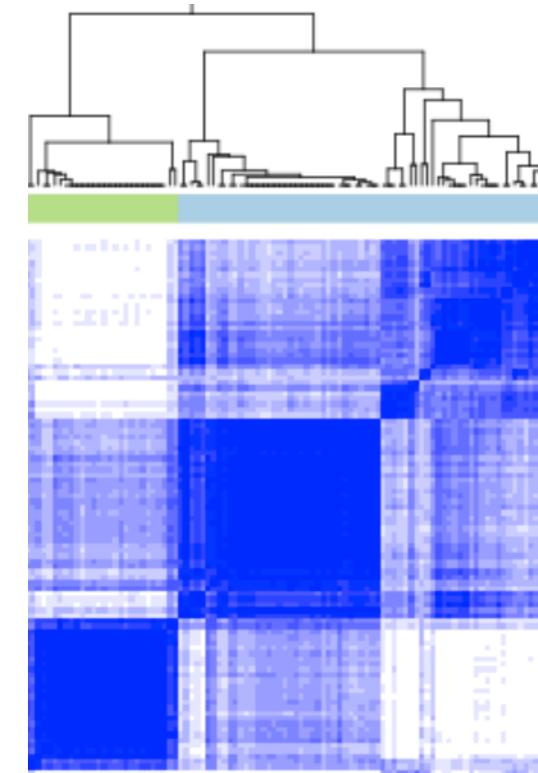
Consensus matrix can be used as a new distance matrix for clustering (e.g., hierarchical clustering)



# Consensus clustering and resampling consensus matrix



Good consensus matrix



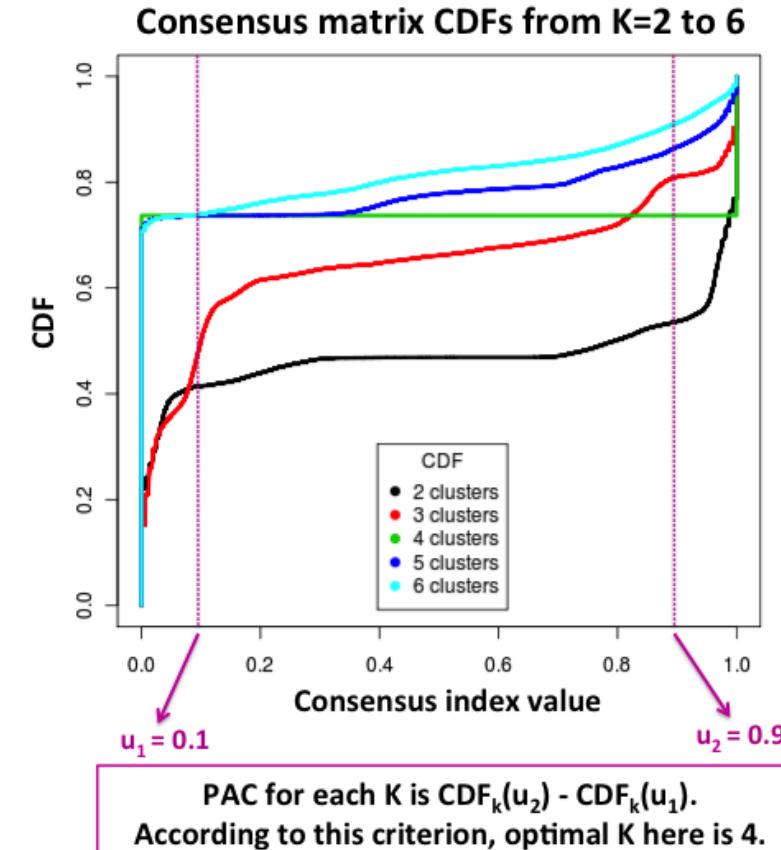
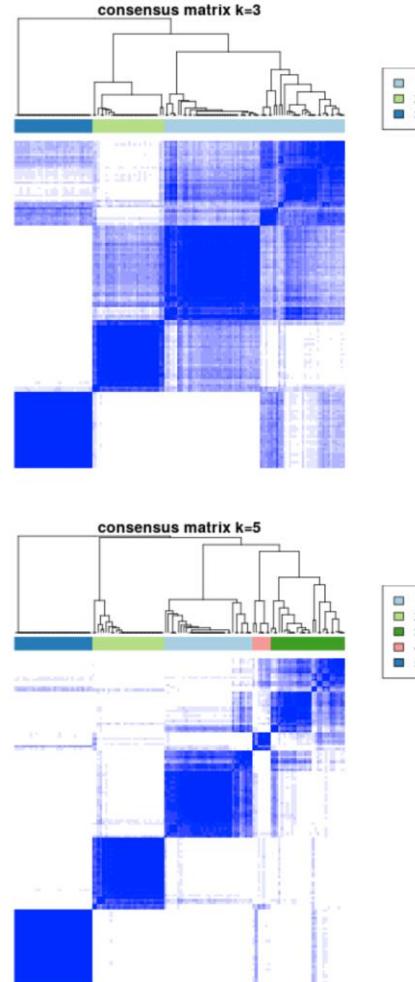
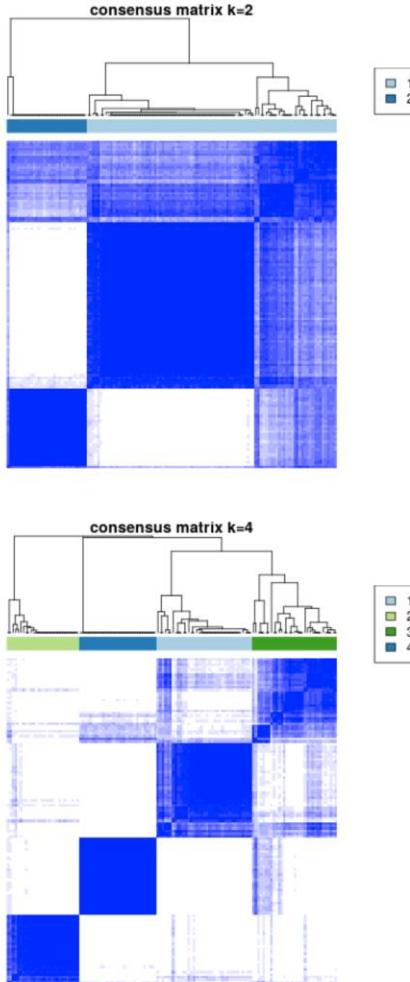
Not that good consensus matrix

How to quantify the goodness?

# Consensus clustering and resampling

Cumulative Density Function (CDF) of consensus matrix

Proportion of ambiguous clustering (PAC)

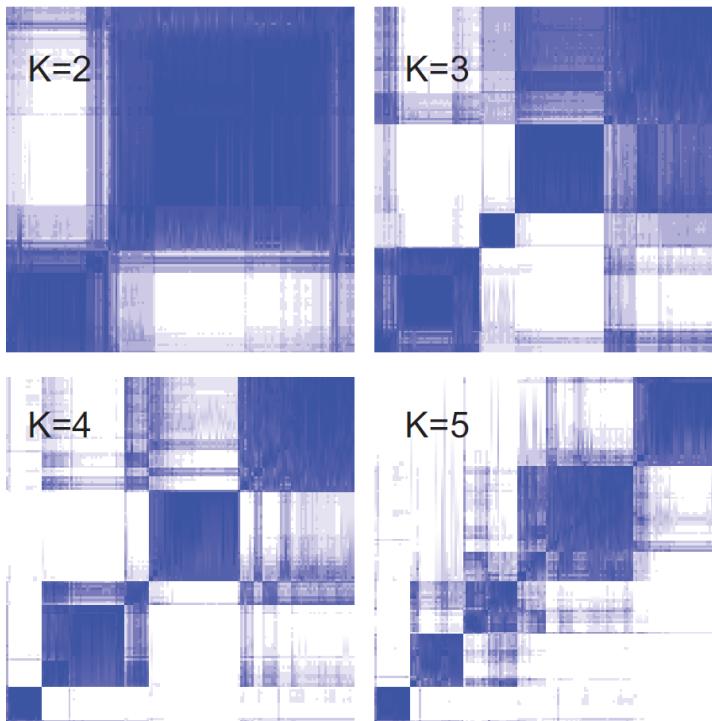


PAC measures the 'contrast' of the consensus matrix

# Consensus clustering and resampling

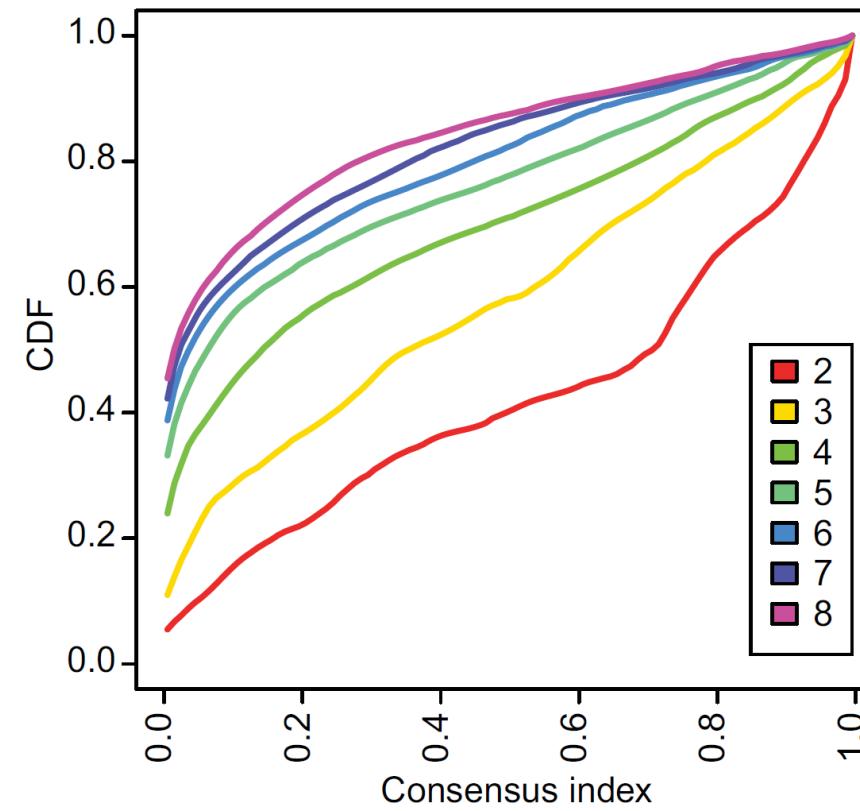
a

Consensus Clustering



b

Consensus CDF



Li et al, Scientific Reports, 2014

# What you should take with you from 'Clustering' lecture

- Distinction between clusters and classes
- Main characteristics of a clustering algorithm
- Major clustering algorithm types
- Good understanding of what k-means, hierarchical clustering and DBSCAN do
- What is the purpose of cluster validation
- Understanding Rand, silhouette index and the principle of consensus clustering