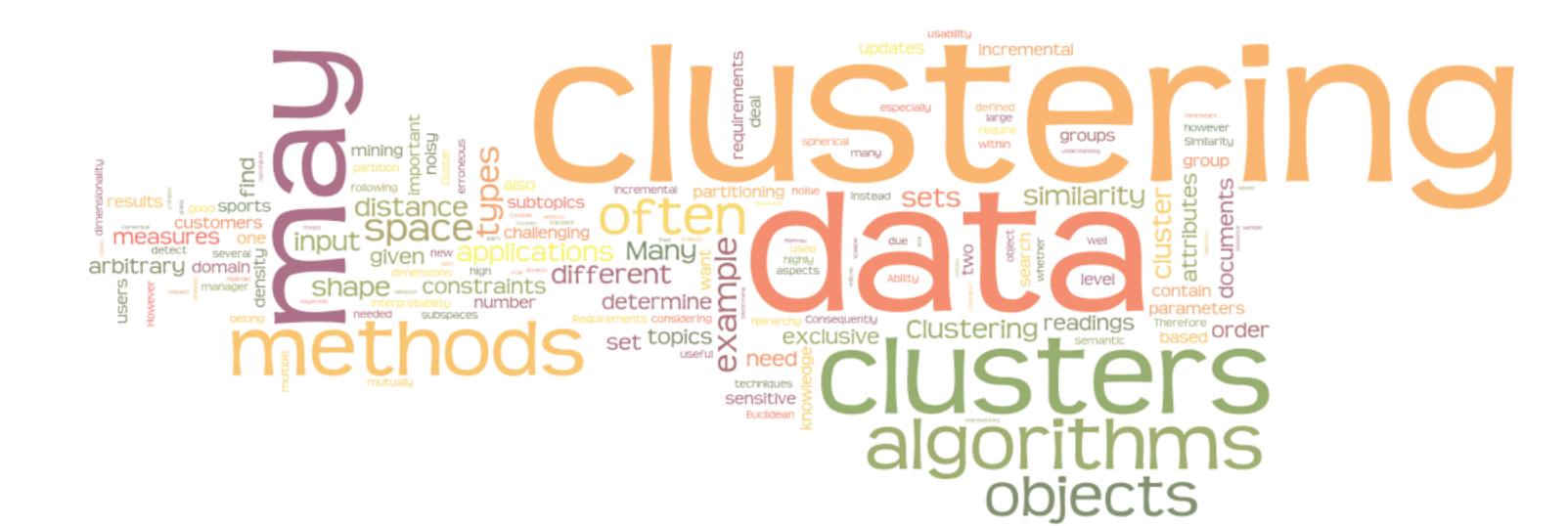
Clustering

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Motivation

Suppose we have large amount of unlabelled texts, based on which we are able to:

- Create a useful content-recommendation service
 («It looks like you love to read about battle-rap. Do you want more? »)
- make conclusions on topics represented in textual data -- or other data structure features (trending topics, emerging trends, blogosphere analysis)
- find duplicate content ("the same piece of news, can be skipped", "plagiarism!")

And so on, and so on... Another ideas?

Clustering

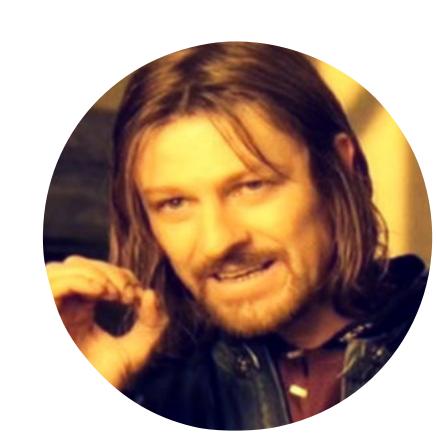
Problems?

There are:

- Objects (documents) space X
- Set of objects (documents) X^I

Our task:

 split document collection into groups (clusters) so that the documents in one group were similar to each other, whereas documents from different groups should be dissimilar



Clustering

There are:

- Objects (documents) space X
- Set of objects (documents) X¹

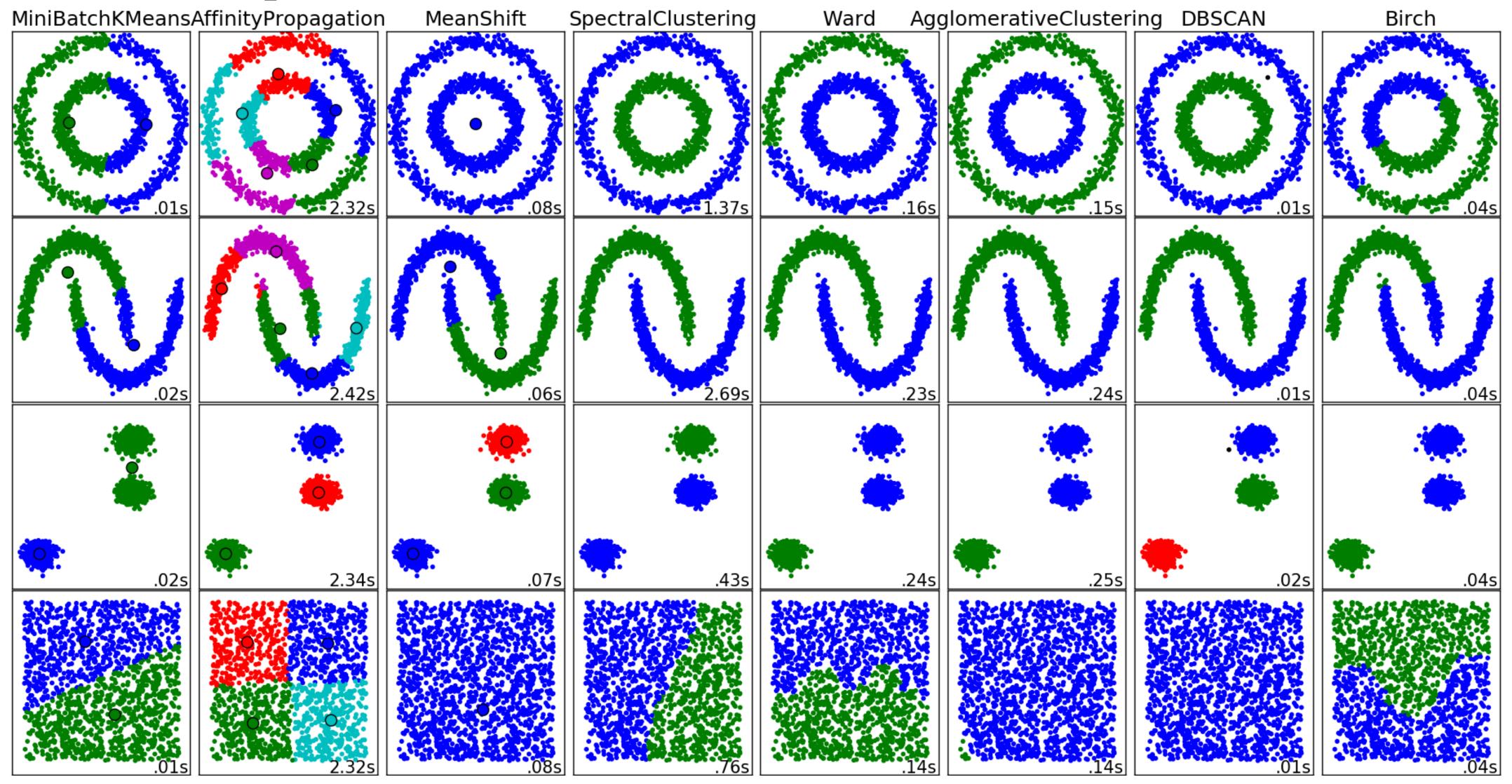
Our task:

 split document collection into groups (clusters) so that the documents in one group were similar to each other, whereas documents from different groups should be dissimilar

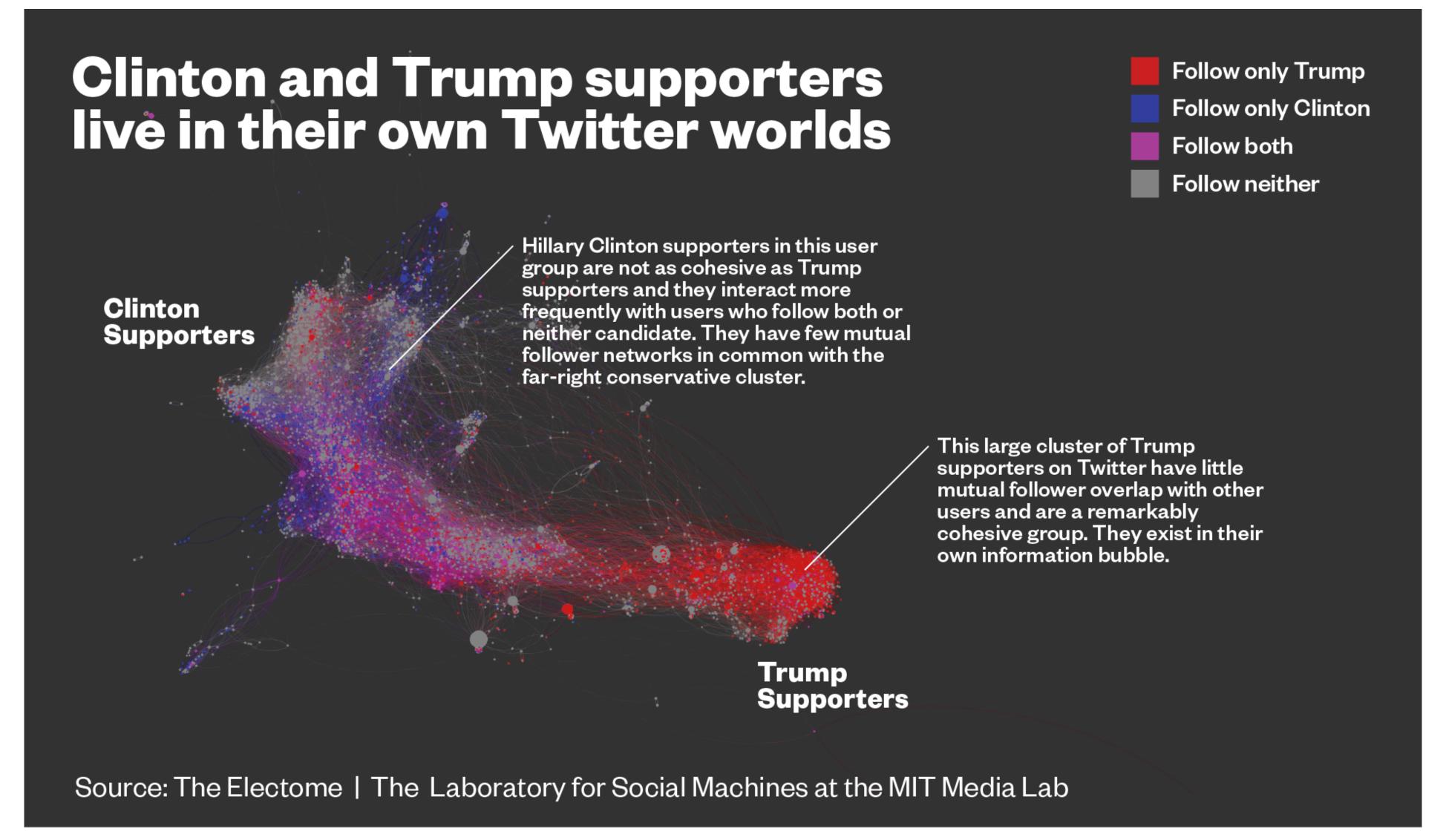
Problems?

- This task implicitly introduce some similarity measure on documents space
- We know nothing about number of clusters K (usually this is the main parameter for the algorithm)
- It's hard to introduce and estimate clustering quality
- There are some problems with huge number of dimensions (<u>curse of</u> <u>dimensionality</u>)

Examples



Examples



Ideas:

- Annotate and check by hand
- Apply to already labeled dataset
- extrinsic evaluation: estimate the 'usefulness' increase for some application
- intrinsic evaluation: estimate some clustering 'quality index'

Can you broke any of this methods?



Ideas:

- Annotate and check by hand
 - and spend all your life labeling wikipedia
- Apply to already labeled dataset
 - so why should we cluster and not train classification model?
- extrinsic evaluation: estimate the 'usefulness' increase for some application
 - but this way we don't look at clusters quality
- intrinsic evaluation: estimate some clustering 'quality index'
 - we look at one index when we optimize, then we look at a 'better' one... why not use the better one for optimization?



...with labeled corpora;

Option 1:

Annotate each pair of objects in the test set: with 1 if they are in the same cluster or 0 if they are in different ones; Then we do the same with our predictions

Thus we can evaluate quality the same way as we can do with classification:

- 1) we can compute **Accuracy** (how many pairs are correctly/incorrectly put into the same cluster)
- 2) or we can compute Precision, Recall, F-measure

...with labeled corpora;

Option 2:

'How pure is each cluster': max share of some true cluster in each of the predicted ones:

 $\frac{1}{N} \sum_{m \in M} \max |d \cap m|$

D — 'true' clusters

M — predicted clusters

Pairs:

• n(n-1)/2 pairs is a lot, the size of dataset (n) can't be large due to that

Purity:

- large number of clusters delivers large purity value!
- if every element is a cluster, purity = 1.0

A lot of clustering evaluation indices were invented, each of them is ugly in its own way:)

For a good start one may take a look at the Wikipedia article

Sorts of sholustering

We can compare clustering algorithms in terms of:

- computational complexity
- do they build flat or hierarchical clustering?
- can the shape of clustering be arbitrary?
 - if not is it symmetrical?
 - can clusters be of dierent size?
 - clusters vary in density of contained objects?
- robustness to outliers
- Hard or soft cluster assignment?

We can compare clustering algorithms in terms of:

- Distance matrix and/or measure function?
- Can we scale it?
- Which intuition these algorithms are based on?
 - graph-based? distribution?
 - density? hierarchy?
- Can we interpret our result?

Clustering algorithms:

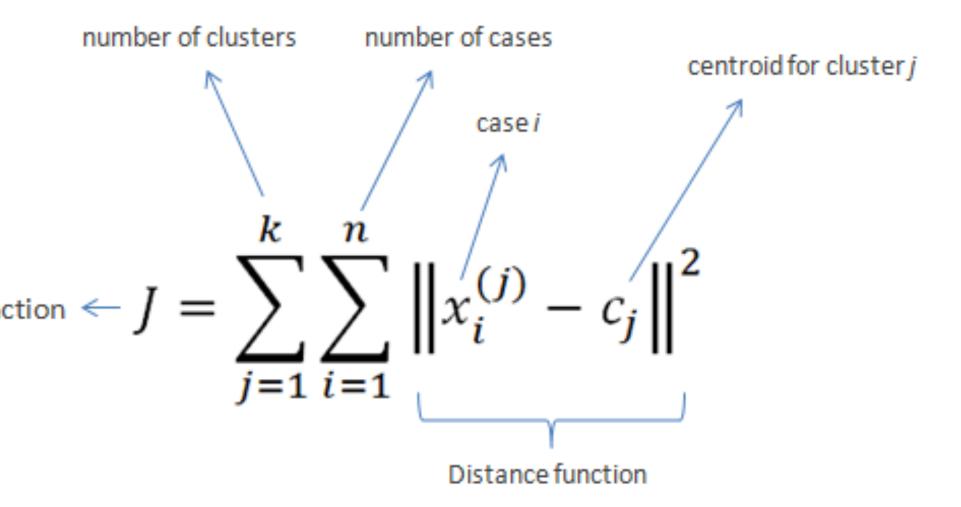
- Representative-based clustering
- Probabilistic clustering
- Hierarchical clustering
- Density-based clistering

Note 1: We already know how to represent text as a vector; all methods we will discuss are of course applicable in other domains and for other data types

Note 2: NB! The algorithms we are going to discuss have numerous modifications and implementations can differ greatly. Take care when carrying out experiments and training models for production environment!

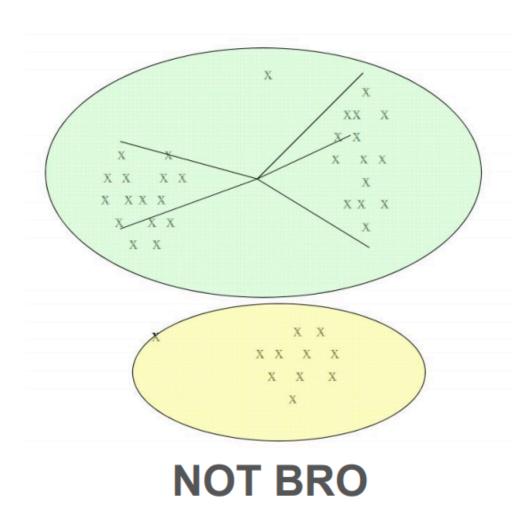
K-means:

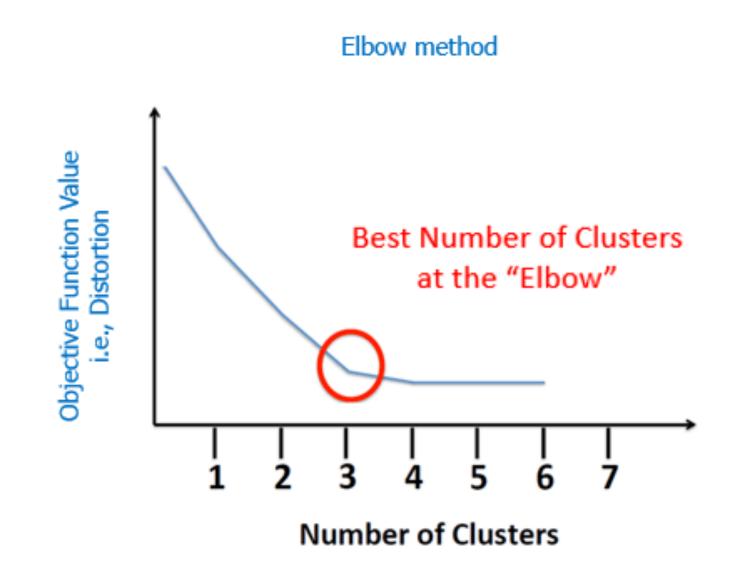
- 1. Applicability: vectors
 - Goal: minimize the sum of squares of distances from centroid of each cluster to each element of the cluster
- 1. Set the number of clusters.
- 2. Choose documents at random -- clusters centroids.
- 3. Include the remaining documents into the closest cluster.
- 4. Compute new cluster centroids as a mean vector in the cluster.
- 5. Repeat steps 3-4, until
 - 1. Centroids stop to change?
 - 2. The partition of the dataset stops to change?
 - 3. Reach num. iteration threshold

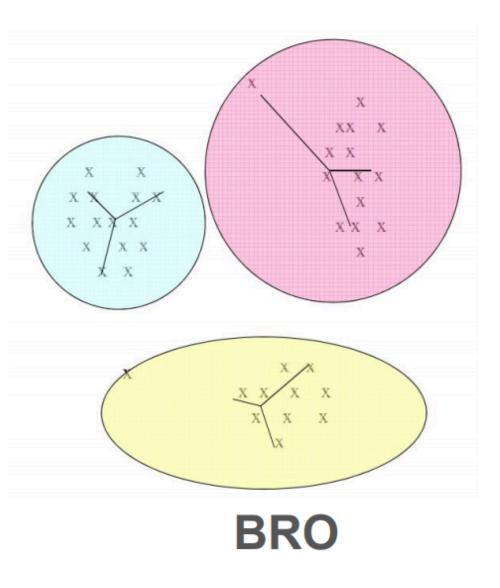


K-means:

- 1. Gradually increase **K**
- 2. Look at the average distance to centroid
- 3. At some value of K it will stop to drop fast; this is the recommended K value







K-means:

Restrictions:

- Sometimes it's hard to find «nearest neighbours»
- can't apply to the domain where there is no such thing as an 'average object'
- is prone to ball-like clusters detection
- always finds K clusters
- sometimes heavily depends on initial centroids candidates choice
- However, there are quite a few modifications useful in real life

Definitions:

- Our space X: $X = \mathbb{R}^n$, our objects $x_i = (f_1(x_i), f_2(x_i) \dots f_k(x_i)) \in X$
- The are **center points** for each cluster: $\mu_y = (\mu_{y1} \dots \mu_{yn})$
- Let $|\omega_y| \sum_{y \in Y} |\omega_y| = 1$, prior probability for each cluster
- Assume that X[|] i.i.d from mixture of distributions: $p(x) = \sum_{y \in Y} \omega_y p(x)$
- Assume that all distributions are Gaussian;

$$f_{\mathbf{X}}(x_1,\ldots,x_k) = rac{\exp\left(-rac{1}{2}(\mathbf{x}-oldsymbol{\mu})^{\mathrm{T}}oldsymbol{\Sigma}^{-1}(\mathbf{x}-oldsymbol{\mu})
ight)}{\sqrt{(2\pi)^k|oldsymbol{\Sigma}|}}$$

Definitions:

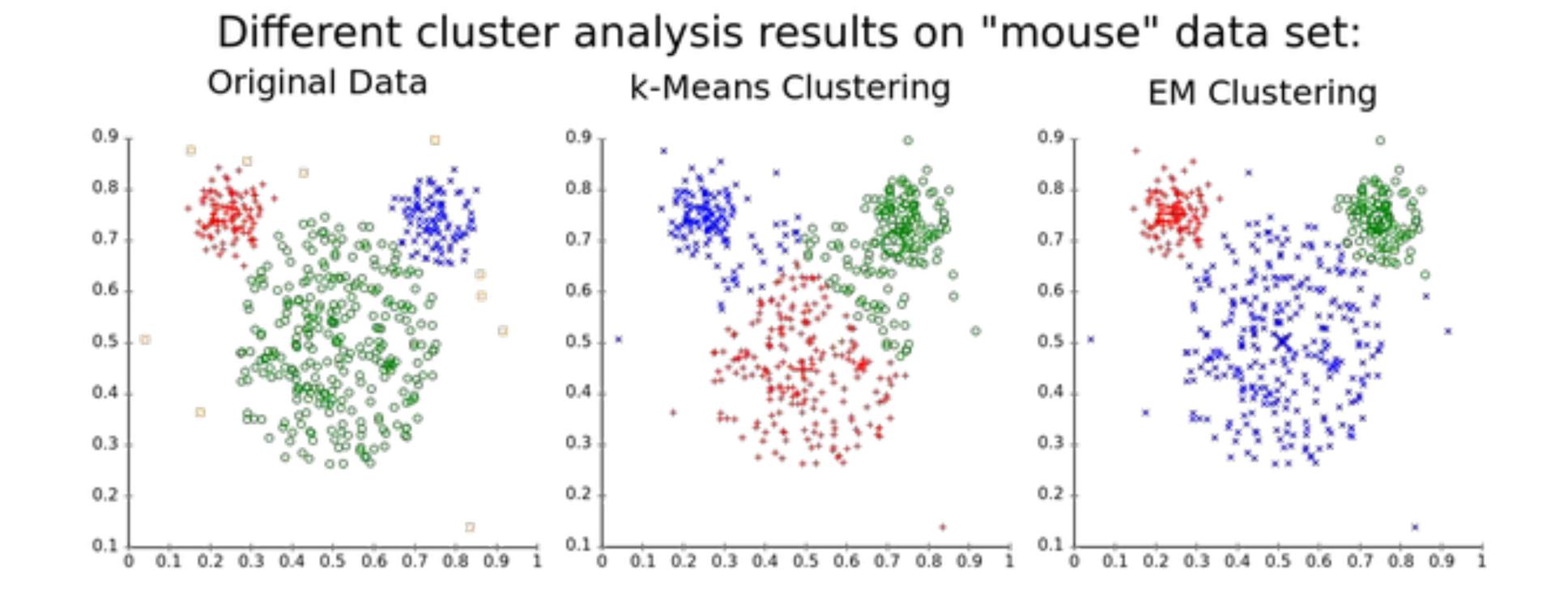
- 1. There are start points for $\omega_{y}, \mu_{y}, \Sigma_{y}, \forall y \in Y$
- 2. E(expectation)-step for y, i=1... $g_{iy} \leftarrow P(y \mid x_i) = \frac{\omega_i p_y(x_i)}{\sum_{z \in \mathcal{Z}} \omega_z p_z(x)}$
- 3. M(maximisation)-step

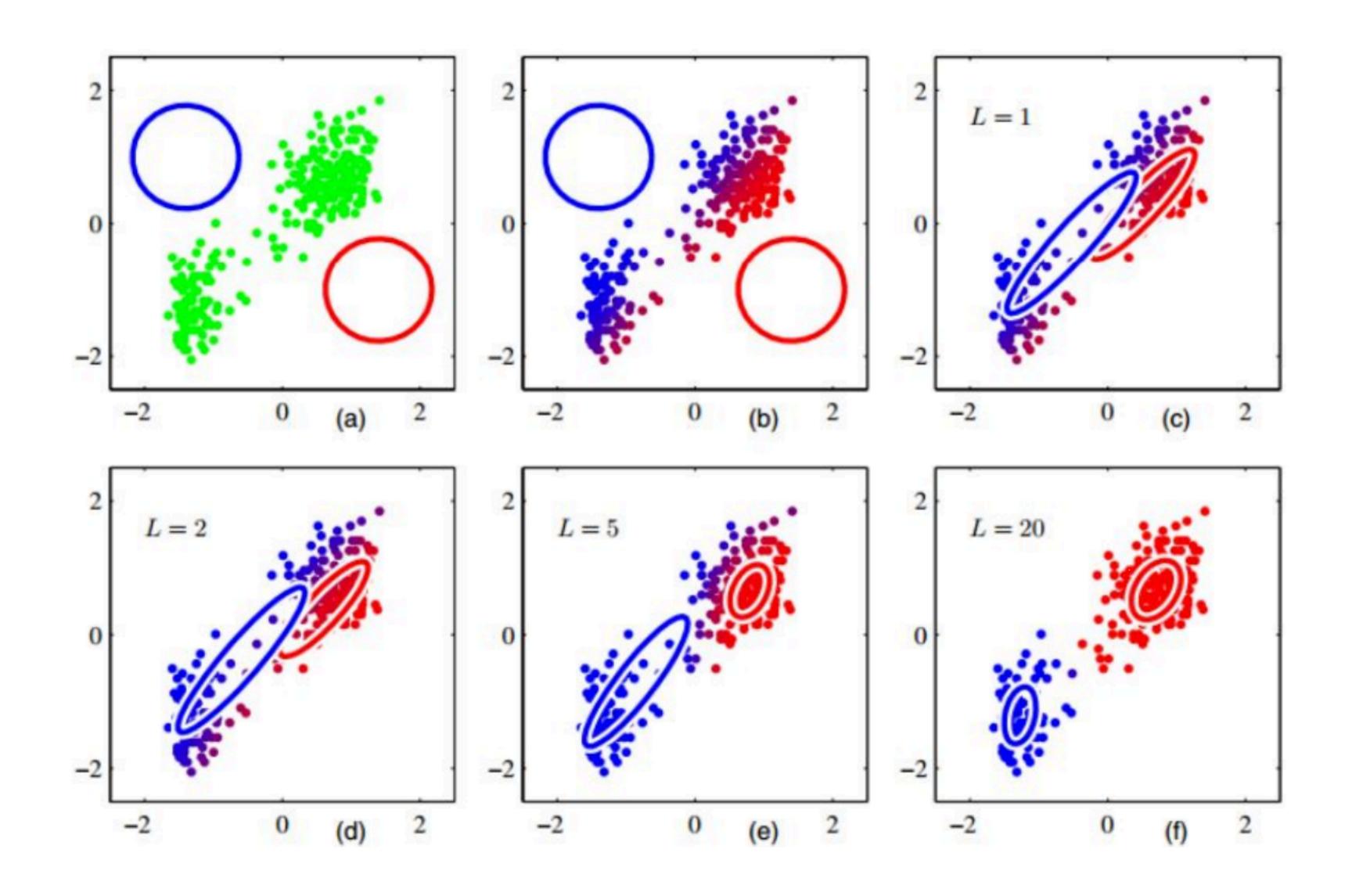
$$\omega_{y} \leftarrow \frac{1}{l} \sum_{i=1}^{l} g_{iy}$$
 $\mu_{yi} = \frac{1}{l * \omega_{y}} \sum_{i=1}^{l} g_{iy} f_{j}(x_{i})$
 $\sigma_{yj} \leftarrow \frac{1}{l * w_{y}} \sum_{i=1}^{l} g_{iy} (f_{j}(x_{i}) - \mu_{yj})^{2}$

- 4. $y_i = argmax_{y \in Y}g_{iy}$
- 5. Repeat until y_i wouldn't fix;

This way we'll have

- 1. 'fuzzy clustering' (soft clustering): clusters probabilities for each document,
- 2. possibility to restrict/give some hint on the possible shapes (distribution family) of the cluster





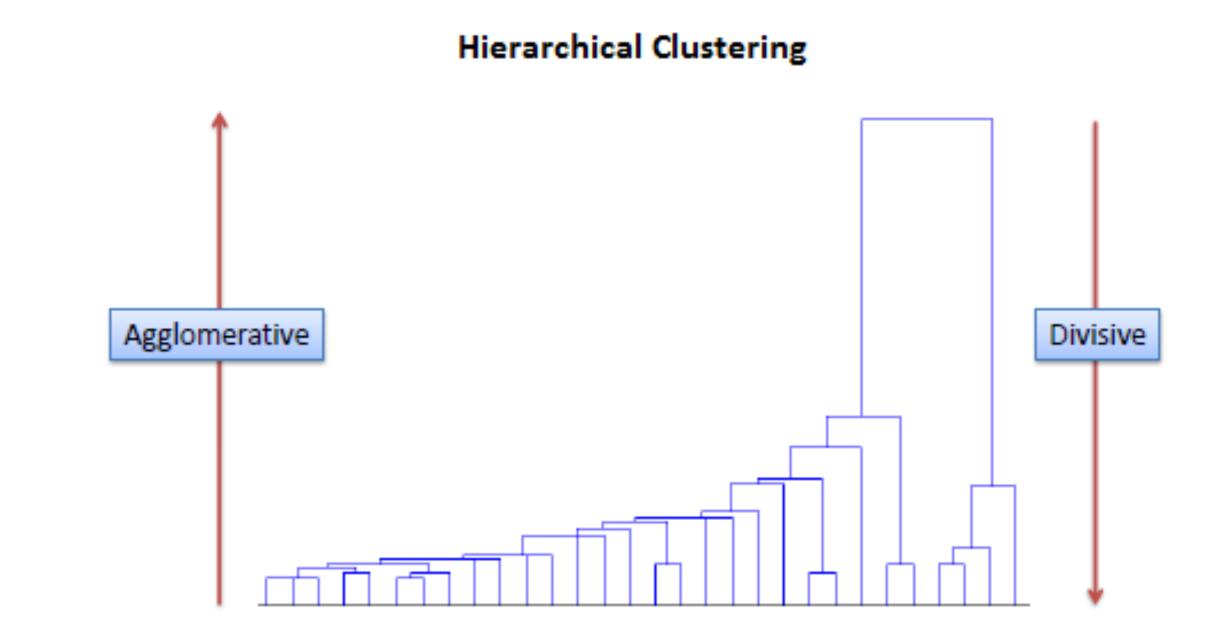
Hierarchical clustering

Two methods types

- 1. agglomerative
- 2. divisive

Each hierarchical method builds a **dendrogram** for further pruning = clusters selection.

Dendrogram shows measures of closeness between objects and sets of objects



Divisive approach:

Example: let's choose some flat clustering method A (e.g. KMeans)

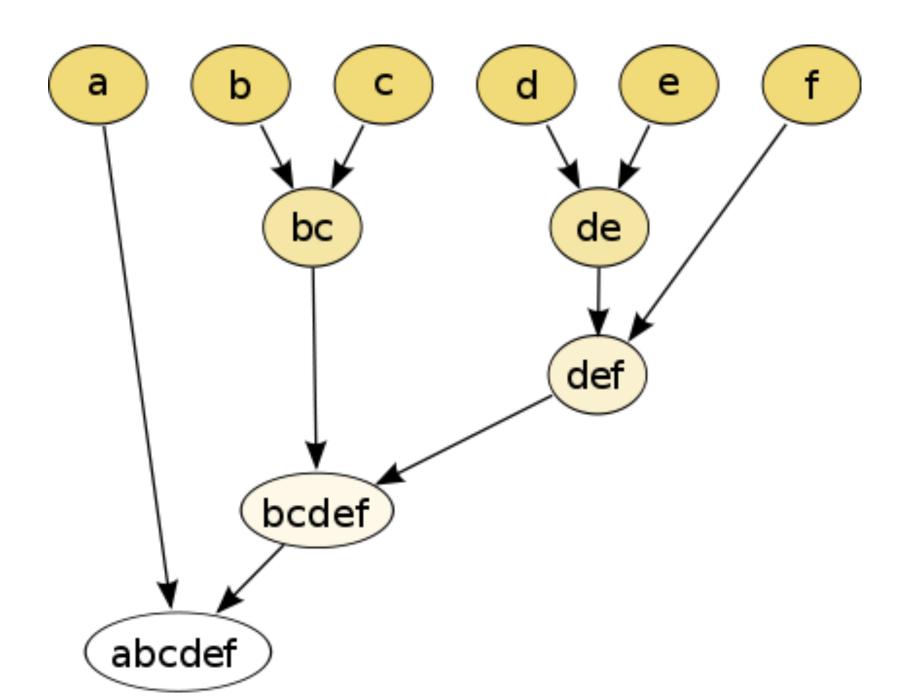
- 1. Initially just 1 cluster containing all elements, root of the tree
- 2. Apply method A to the leaf of the tree (chosen by some rule).
- 3. Add resulting clusters as leaves (x being their 'parent').
- 4. Repeat 2-3, until the cardinality of each 'leaf' is equal to 1.

Divisive approach:

A more popular and intuitive approach

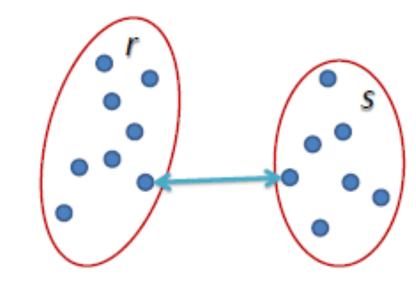
- 1. Initially each element is a cluster of size 1
- 2. Using a certain rule, we choose two closest clusters and merge them into one

3. Problems?

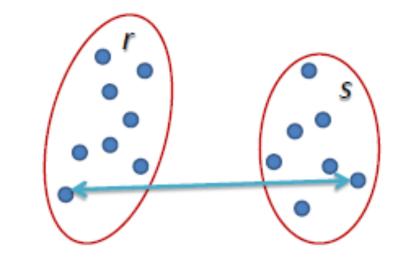


How to compute distance:

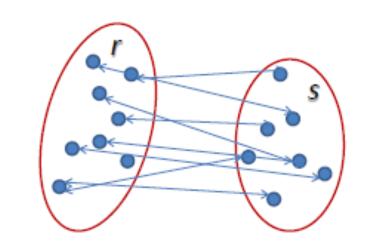
- Single linkage: the distance between two clusters is defined as the *shortest* distance between two points in each cluster.
- Complete Linkage: the distance between two clusters is defined as the *longest* distance between two points in each cluster
- Average Linkage: the distance between two clusters is defined as the average distance between each point in one cluster to every point in the other cluster.
- Ward linkage: difference between sum(sqr(distances)) inside the possible clusters union and sum(sqr(distances)) inside each of the two clusters separately



$$L(r,s) = \min(D(x_{ri}, x_{sj}))$$



$$L(r,s) = \max(D(x_{ri},x_{sj}))$$



$$L(r,s) = \frac{1}{n_r n_s} \sum_{i=1}^{n_r} \sum_{j=1}^{n_s} D(x_{ri}, x_{sj})$$

Tips and tricks:

- How to choose num clusters?
 //https://stats.stackexchange.com/questions/3685/where-to-cut-a-dendrogram
- Use Ward, practically it better
- Read literature, it's interesting

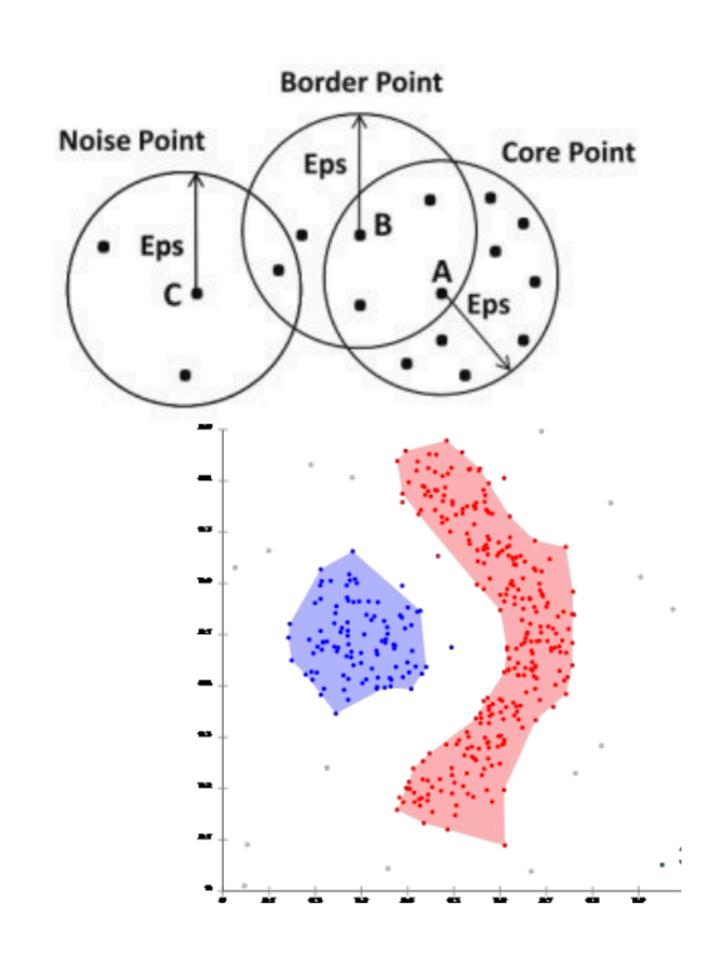
Density-based: DBSCAN

Density-Based Spatial Clustering of Applications with Noise the most cited clustering algorithm

Set ε (distance) and k (an integer)

Elements are split into 3 types:

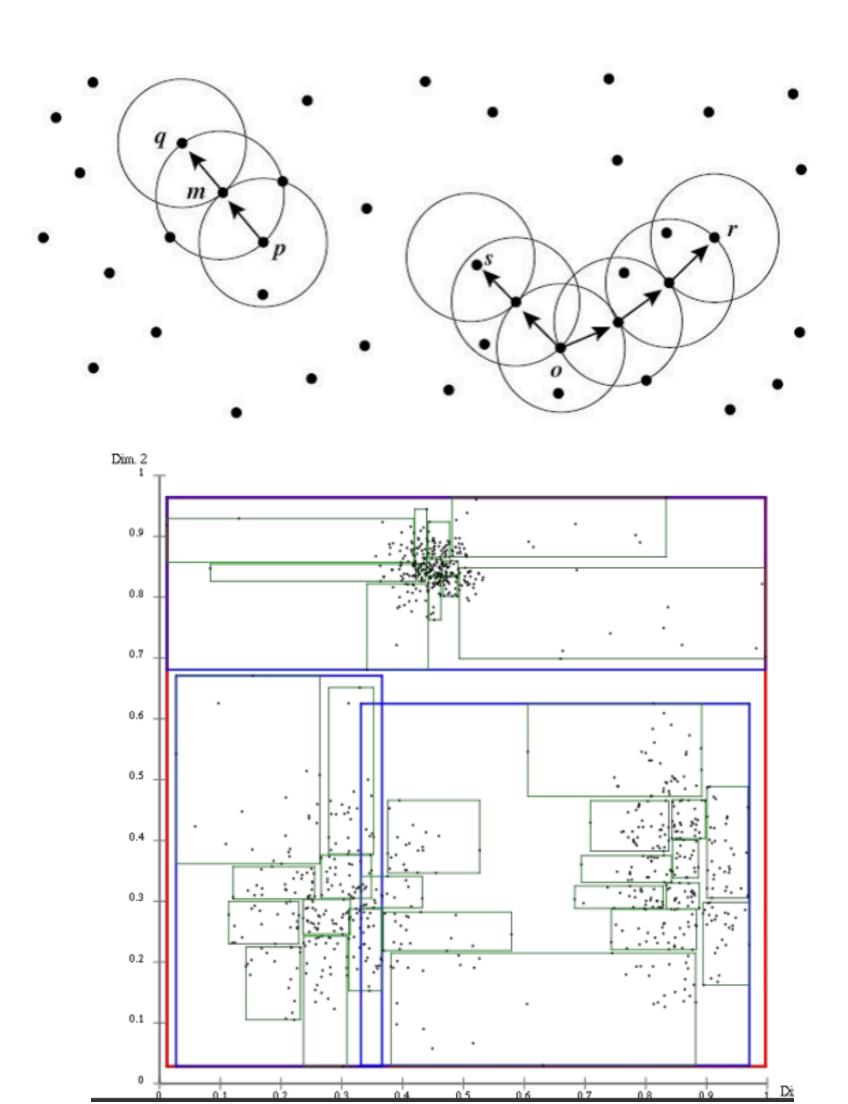
- 1. Core points: elements having at least k other elements in their ε-neighbourhood
- 2. Border points: elements having at least b element in their ε-neighbourhood
- 3. Noise points: other elements



Density-based: DBSCAN

Algorithm

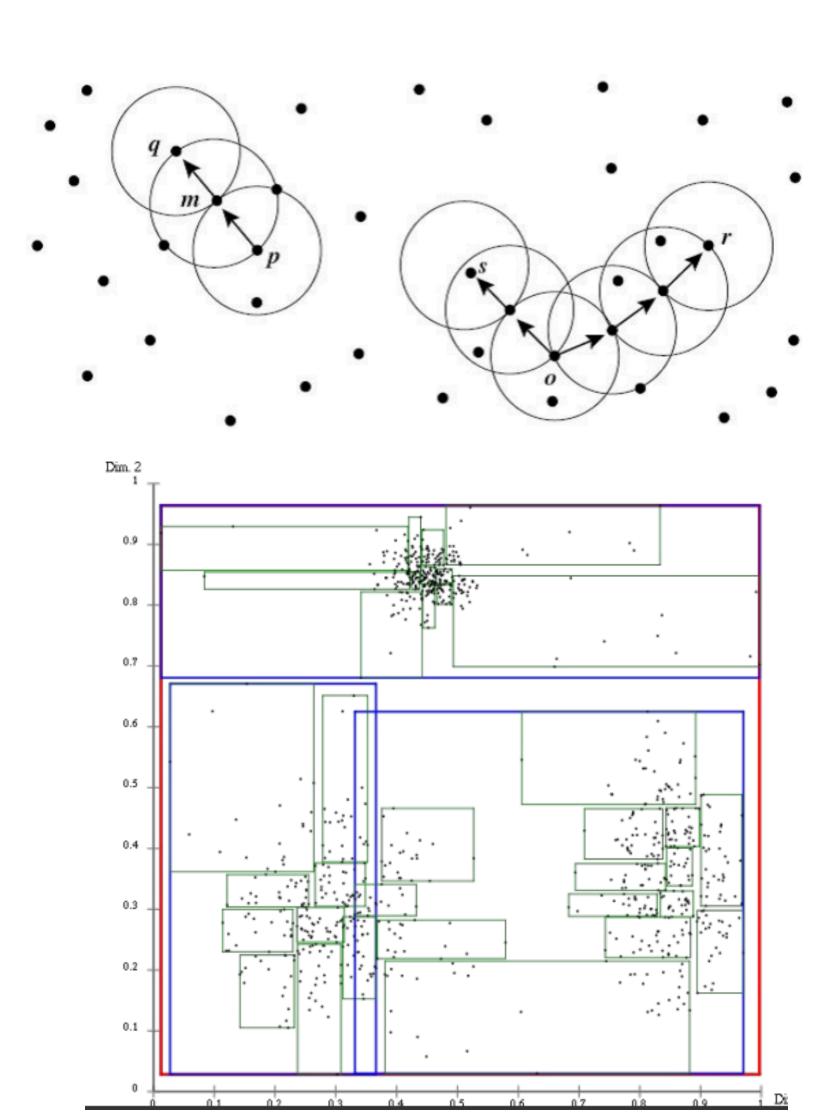
- 1) Mark elements with those three types
- 2) Build a graph, connecting core points that are no farther than ε from each other
- 3) Determine connected components
- 4) Link every border point to the closest connected component



Density-based: DBSCAN

Discussion:

- + determines the number of clusters
- + robust to outliers and noise
- + detects clusters of arbitrary shape and form
- is slow
- fails at determining clusters of different density
- tuning parameters may be a challenge



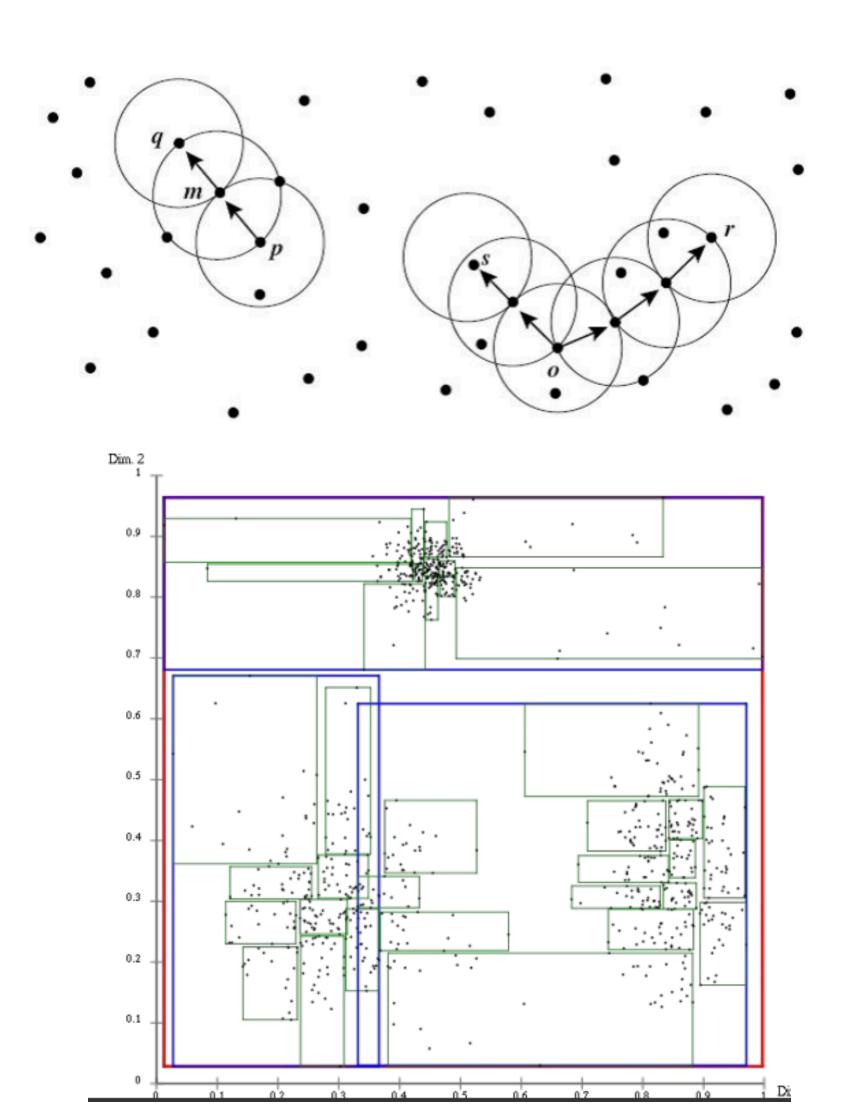
Other guys:

CURE (Clustering Using REpresentatives: hybrid of hierarchical and flat clustering; keeping several representative data points for each cluster)

BIRCH (hierarchical, designed for large datasets, we build a tree of subclusters, preserving certain constraints, SIGMOD 10y test time award)

OPTICS and other DBSCAN modifications (DBSCAN taking density into account)

Community and graphs



Tools & Refs

Mainstream instruments allowing to try different approaches

- scipy.cluster
- sklearn.cluster
- custom libraries, e.g. pyclustering
- 1.CSC 2014 course
- 2. Mining of Massive Datasets Jure Leskovec, Anand Rajaraman, Jeff Ullman
- 3. Scikit-learn docs
- 4. MSU course slides and other materials MSU course slides and other materials
- 5. <u>EM-algorithm @ ml.ru</u>
- 6. Wikipedia (English)