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**Prof. Ing. Jan Flusser, DrSc.**

**Lecture 4 – Clustering**

# Unsupervised Classification

## (Cluster analysis)

**Training set is not available, No. of classes may not be *a priori* known**

# What are clusters?

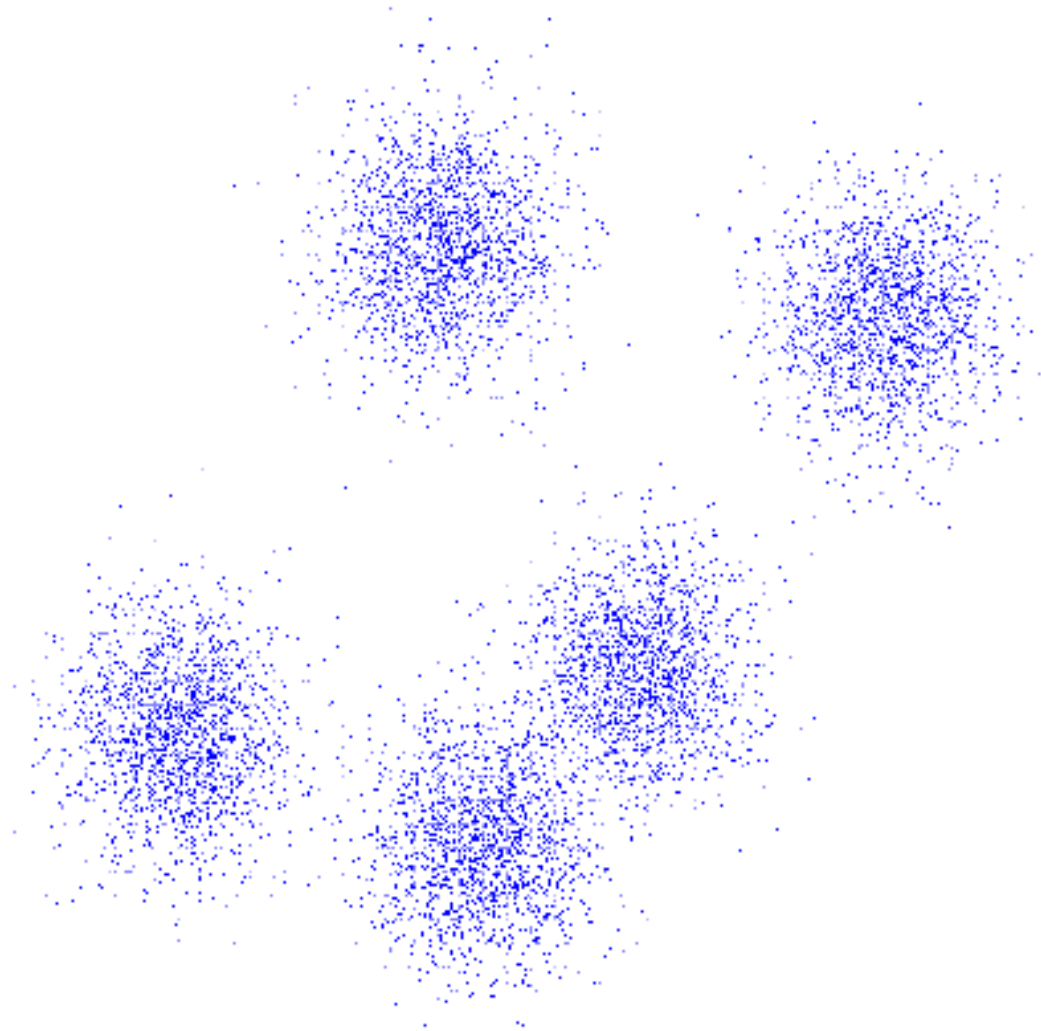
## Intuitive meaning

- compact, well-separated subsets

## Formal definition

- many attempts, mostly unsatisfactory  
( $t$ -connectivity, diameter constraint, ...)
- any partition of the data into disjoint subsets

# What are clusters?



# How to compare different clusterings? (Ward criterion)

Variance measure  $J$  should be minimized

$$J = \sum_{i=1}^N \sum_{\mathbf{x} \in C_i} \|\mathbf{x} - \mathbf{m}_i\|^2$$

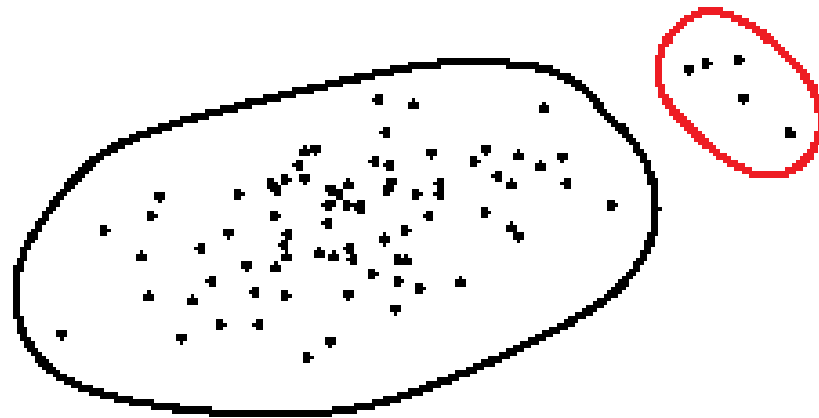
**Drawback** – only clusterings with the same  $N$  can be compared. Global minimum  $J = 0$  is reached in the degenerated case.

# Minimization of $J$

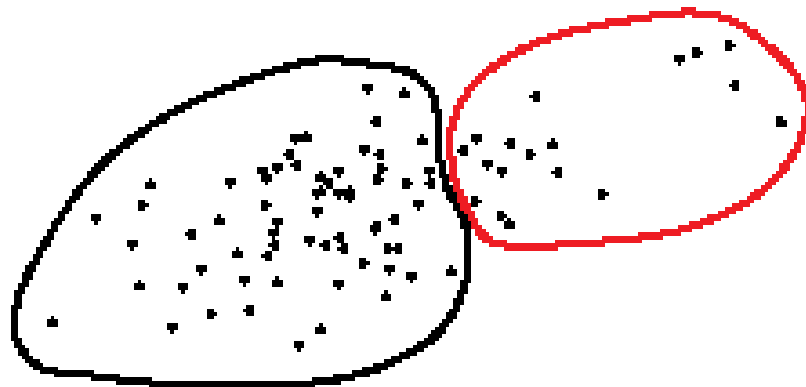
## Drawbacks

- The results are sometimes “intuitively wrong” because  $J$  prefers clusters with approx the same size

# An example of a “wrong” result

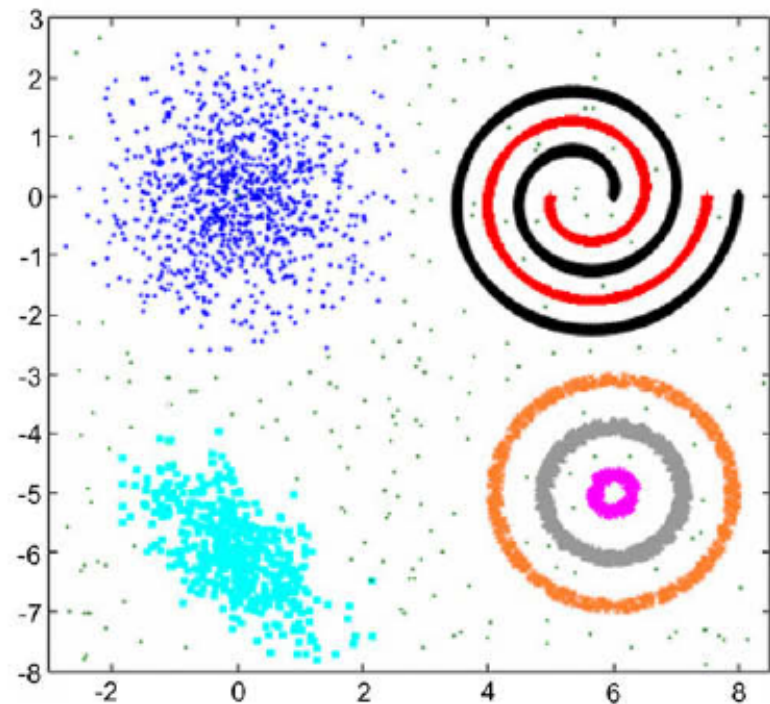
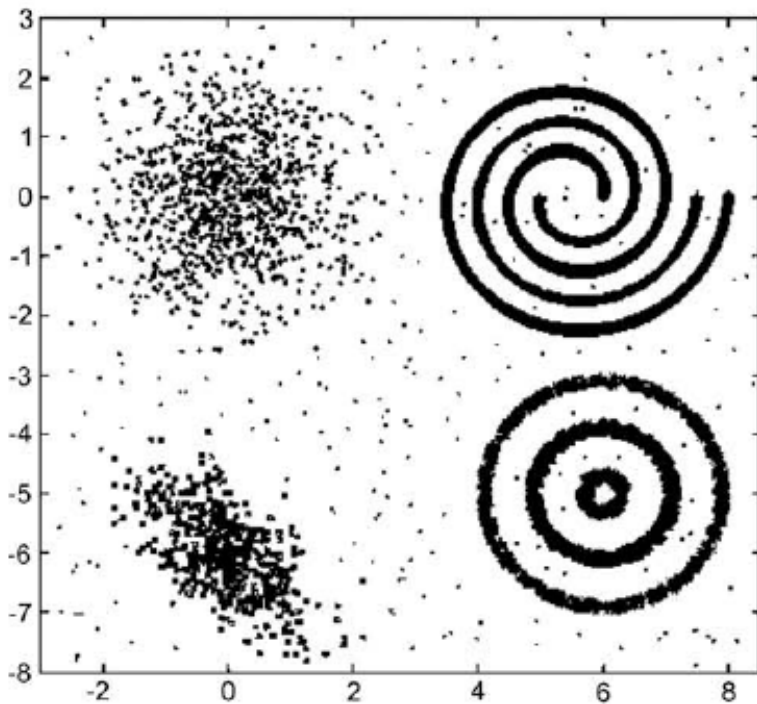


$J_e = large$



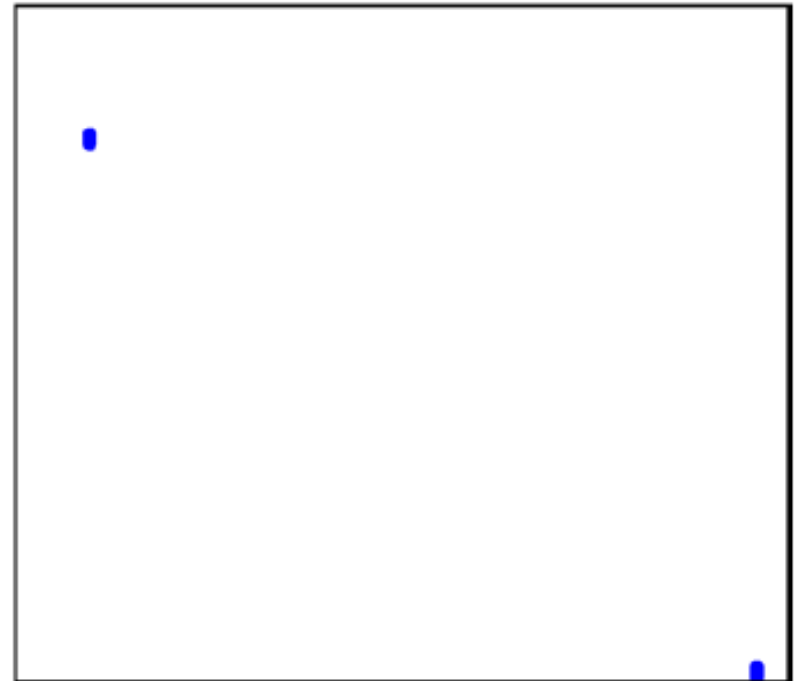
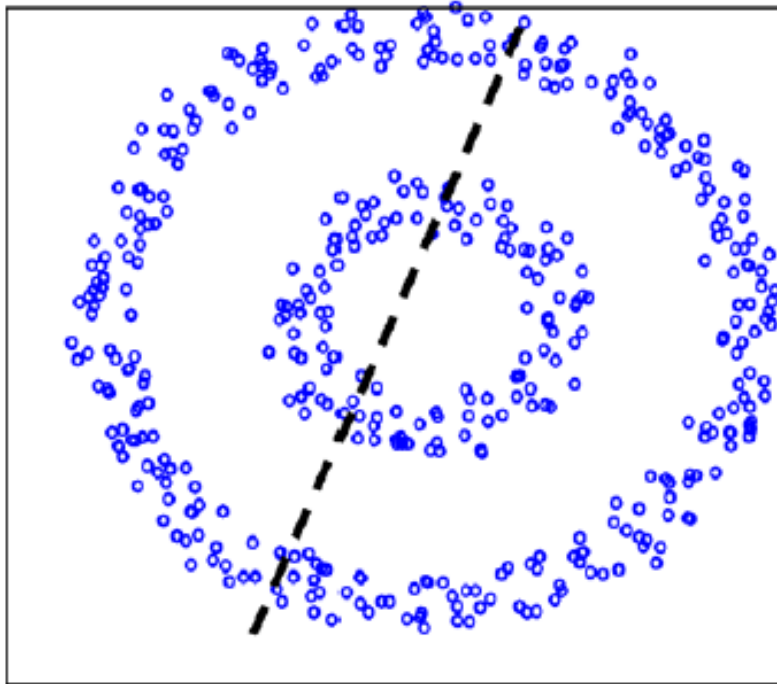
$J_e = small$

**Drawback** – assumes uncorrelated features  
and “convex” clusters





# Solution – proper transform of the features



# Clustering techniques

- **Iterative methods**
  - typically if  $N$  is given
- **Hierarchical methods**
  - typically if  $N$  is unknown
- **Other methods**
  - sequential, graph-based, branch & bound, fuzzy, genetic, model-based, etc.

# Sequential clustering

- $N$  may be unknown
- Very fast but not very good
- Each point is considered only once

**Idea:** a new point is either added to an existing cluster or it forms a new cluster. The decision is based on the **user-defined distance threshold**.

# Sequential clustering

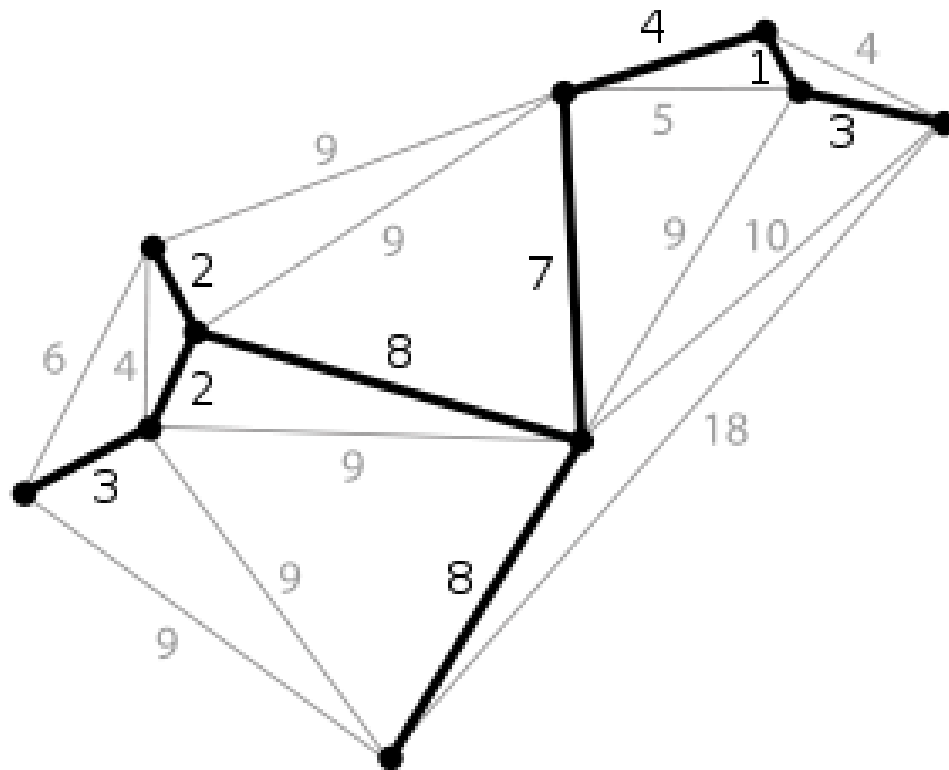
## Drawbacks:

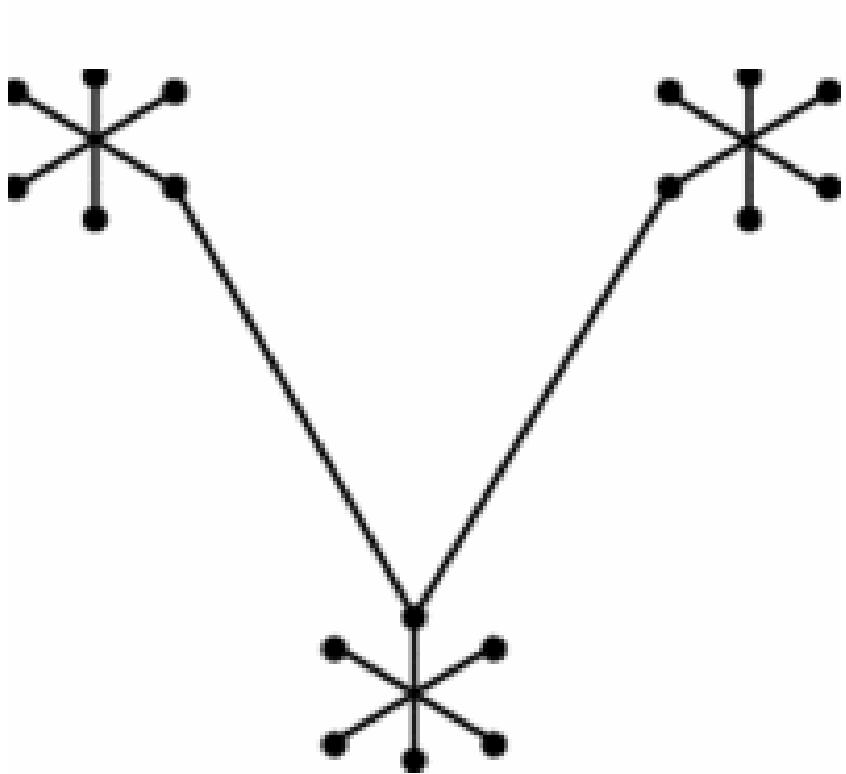
- Dependence on the distance threshold
- Dependence on the order of data points

# Graph-based clustering

**Idea:** Construct a shortest spanning tree and then divide it into clusters by removing some edges.

- Naive approach: remove  $N-1$  longest edges
- Better: remove  $N-1$  long and **inconsistent** edges





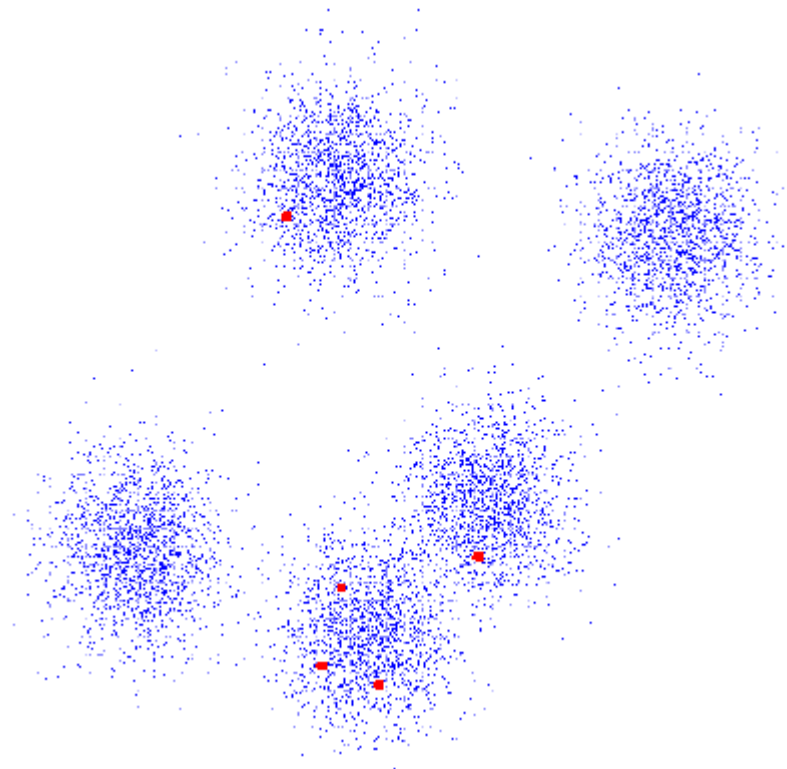
# Iterative clustering methods

- $N$ -means clustering
- Iterative minimization of  $J$
- ISODATA  
Iterative Self-Organizing DATa Analysis



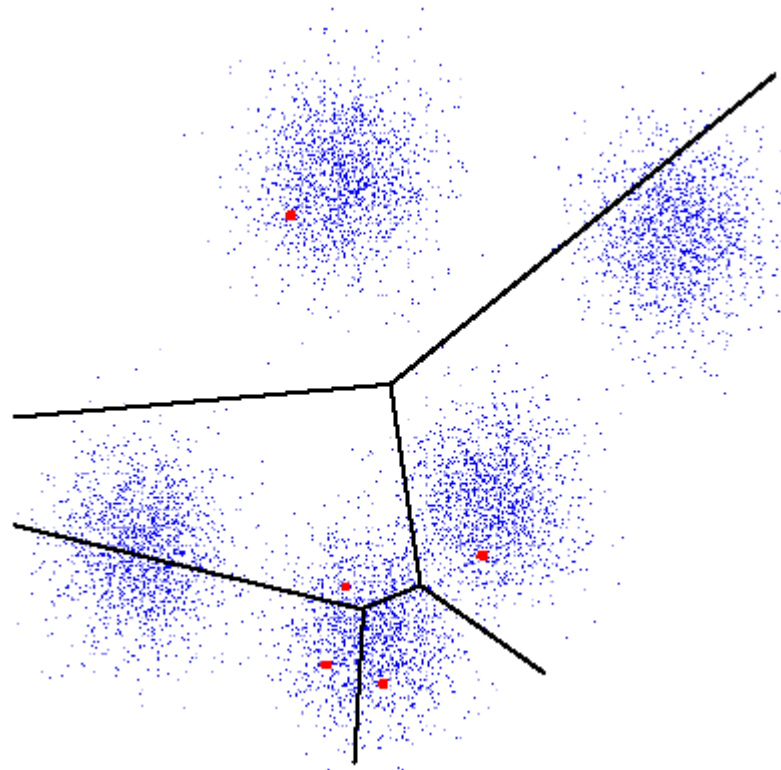
# $N$ -means clustering

1. Select  $N$  initial cluster centroids.



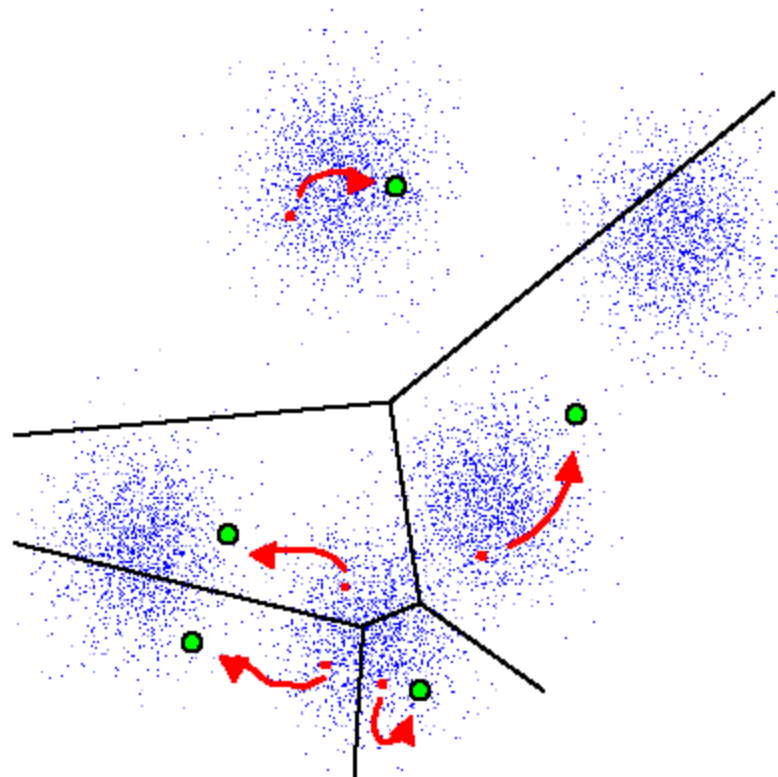
# $K$ -means clustering

2. Classify every point  $x$  according to minimum distance.



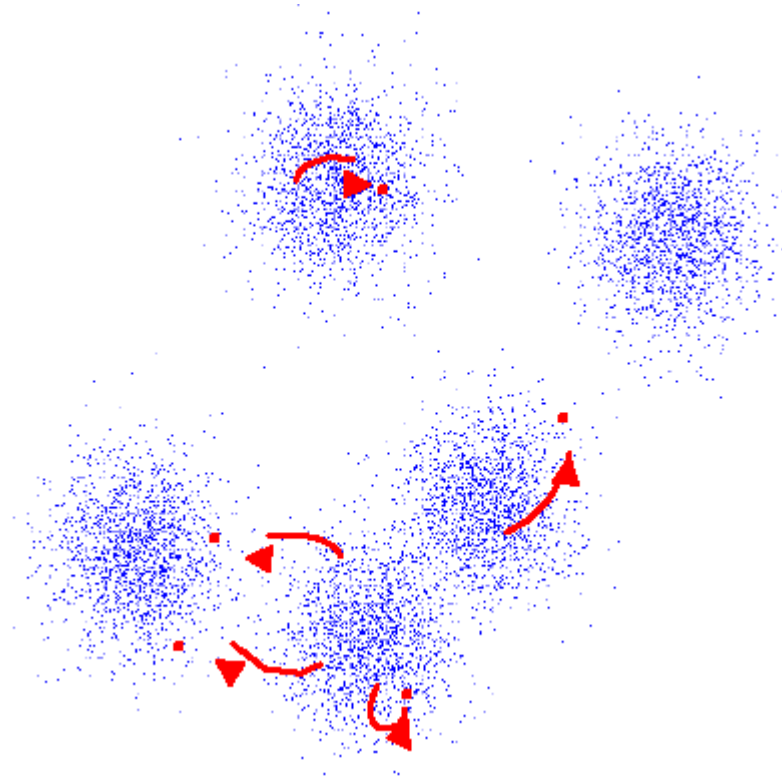
# $N$ -means clustering

3. Recalculate the cluster centroids.



# $k$ -means clustering

4. If the centroids did not change then STOP  
else GOTO 2.



# $K$ -means clustering

## Drawbacks

- The result depends on the initialization.
- $J$  is not minimized
- The results are sometimes “intuitively wrong”.

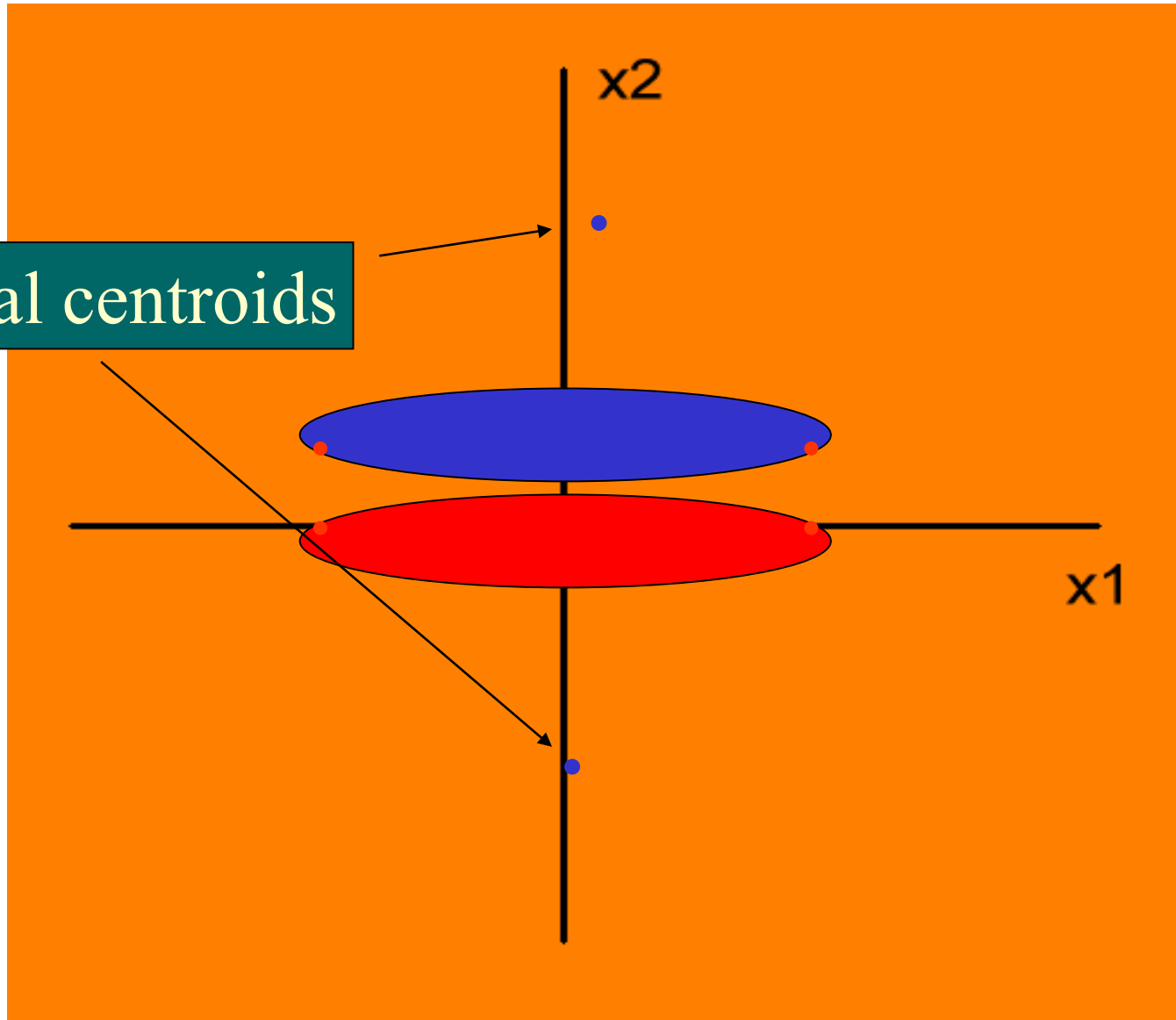
# $N$ -means clustering – An example

Two features, four points, two clusters ( $N = 2$ )

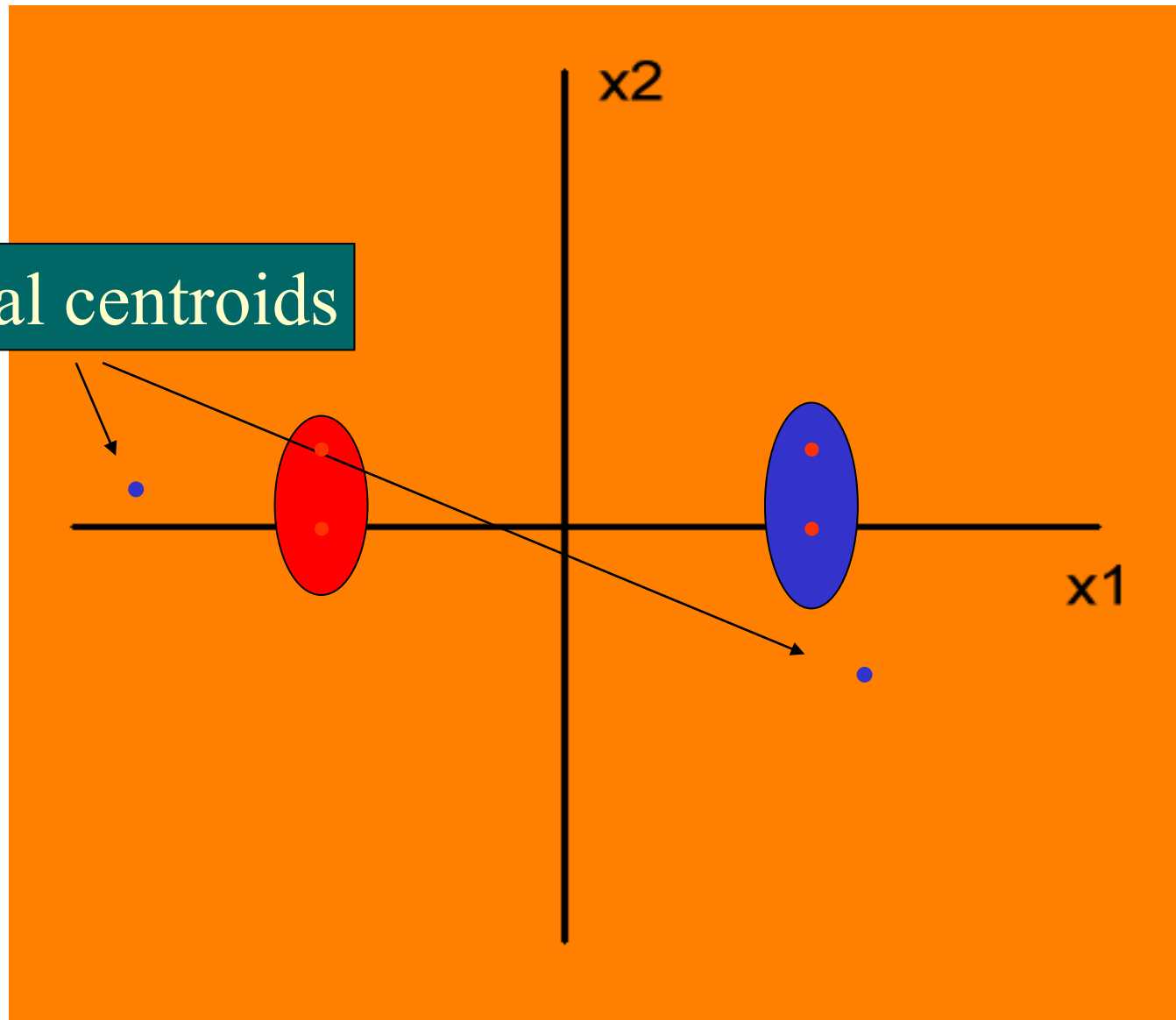
Different initializations  $\rightarrow$  different clusterings

# $K$ -means clustering – An example

Initial centroids



# $K$ -means clustering – An example





# Iterative minimization of $J$

1. Let's have an initial clustering (by  $K$ -means)
2. For every point  $x$  do the following:  
Move  $x$  from its current cluster to another cluster, such that the decrease of  $J$  is maximized.
3. If all data points do not move, then STOP.

# Iterative minimization of $J$

## Drawbacks

- The algorithm is optimal in each step but in general global minimum of  $J$  is not reached.

# ISODATA

Iterative clustering,  $N$  may vary.

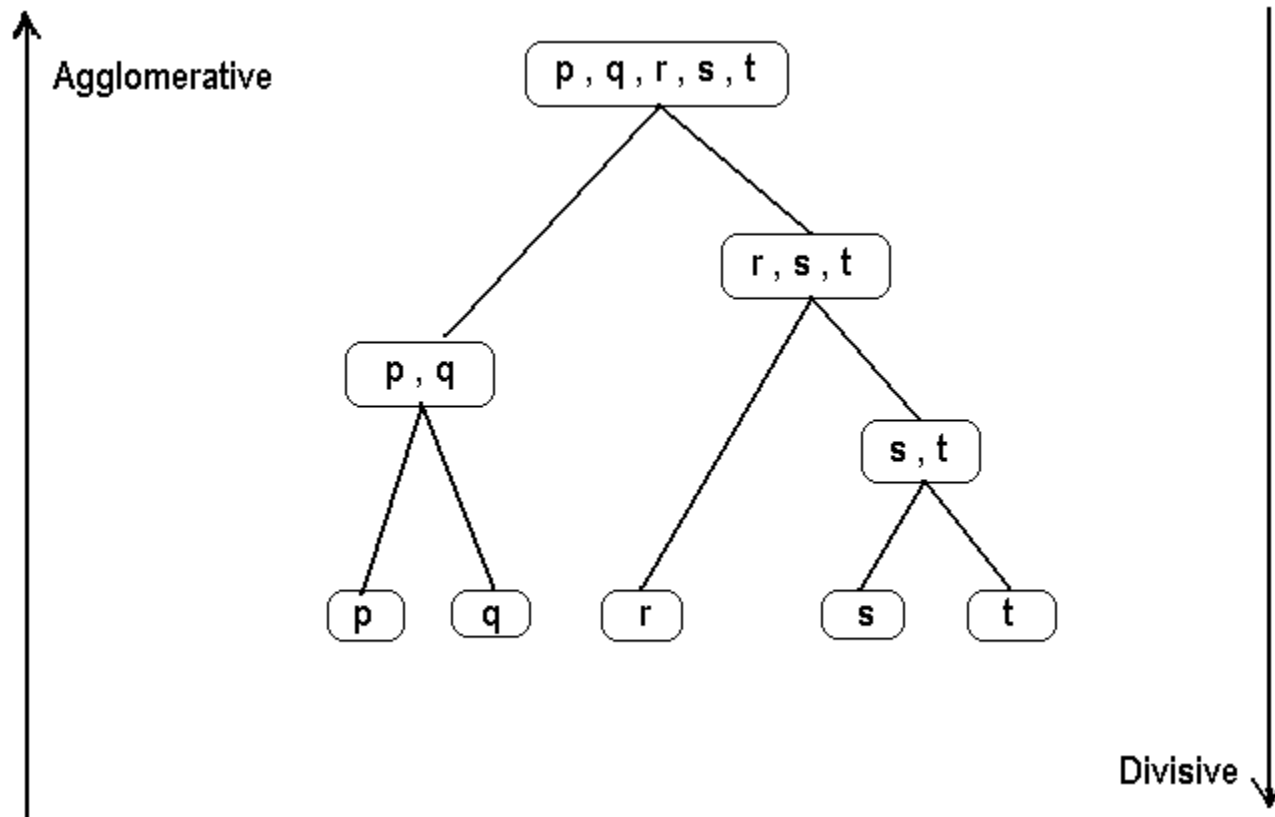
Sophisticated method, a part of many statistical software systems.

Postprocessing after each iteration

- Clusters with few elements are cancelled
- Clusters with big variance are divided
- Other merging and splitting strategies can be implemented

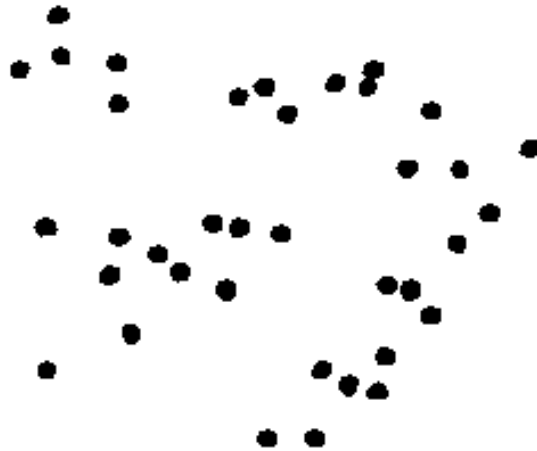
# Hierarchical clustering methods

- Agglomerative clustering
- Divisive clustering



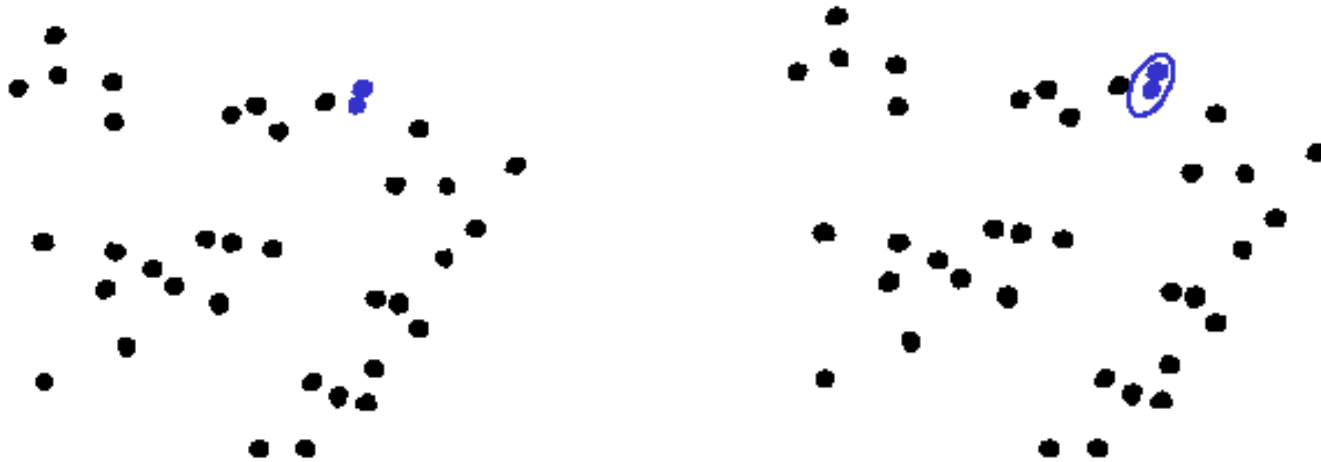
# Basic agglomerative clustering

1. Each point = one cluster



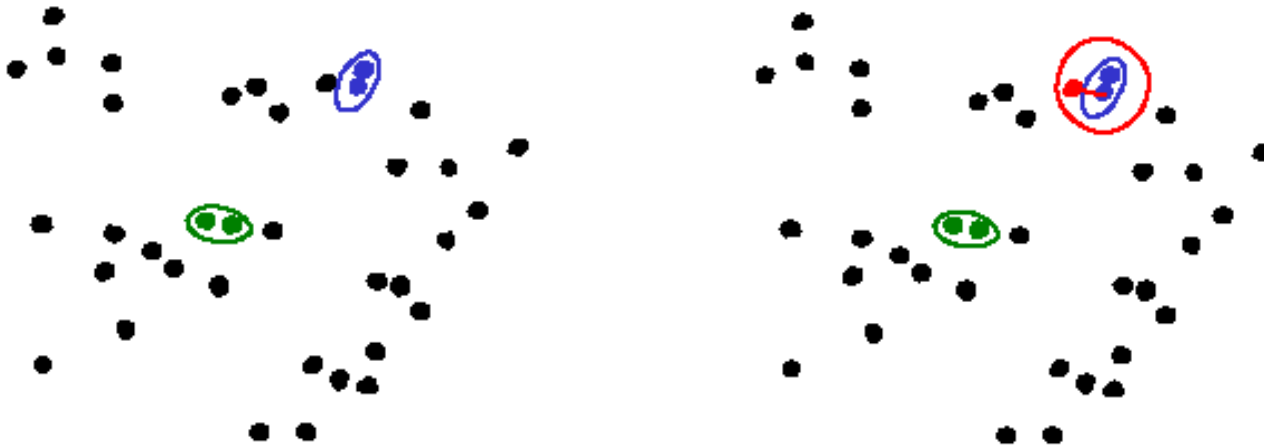
# Basic agglomerative clustering

1. Each point = one cluster
2. Find two “nearest” or “most similar” clusters and merge them together



# Basic agglomerative clustering

1. Each point = one cluster
2. Find two “nearest” or “most similar” clusters and merge them together
3. Repeat 2 until the stop constraint is reached



# Basic agglomerative clustering

Particular implementations of this method differ from each other by

- The STOP constraints
- The distance/similarity measures used

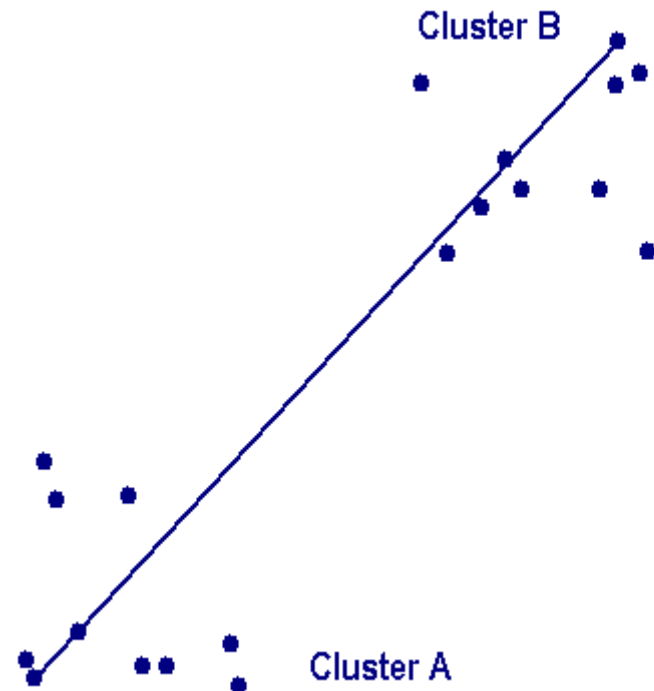
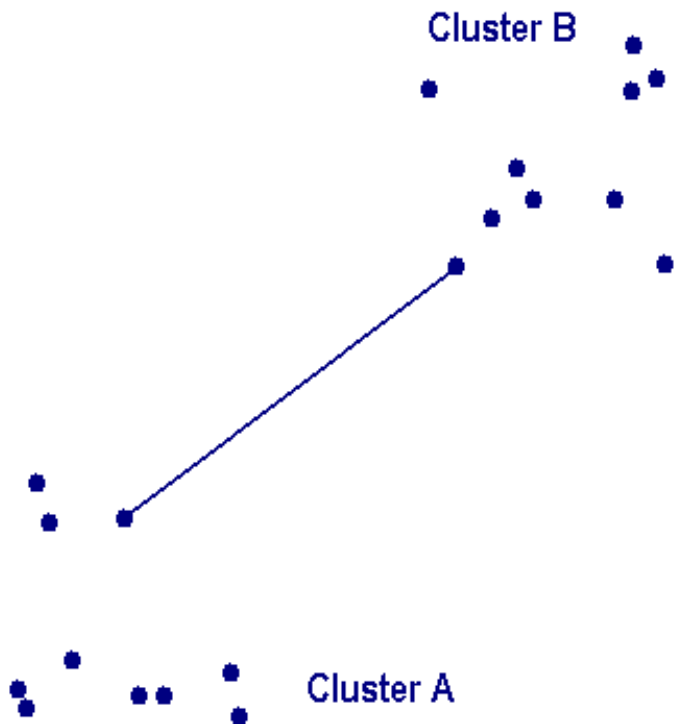


# Simple between-cluster distance measures

$$d(A,B) = d(m_1, m_2)$$

$$d(A,B) = \min d(a,b)$$

$$d(A,B) = \max d(a,b)$$



# Other between-cluster distance measures

$d(A, B) =$  Hausdorff distance  $H(A, B)$

$$d(A, B) = J(A \cup B) - J(A, B)$$

# Efficient implementation of hierarchical clustering

At each level, the distances are calculated using the distances from the previous level.

The upgrade is much faster than a complete calculation.

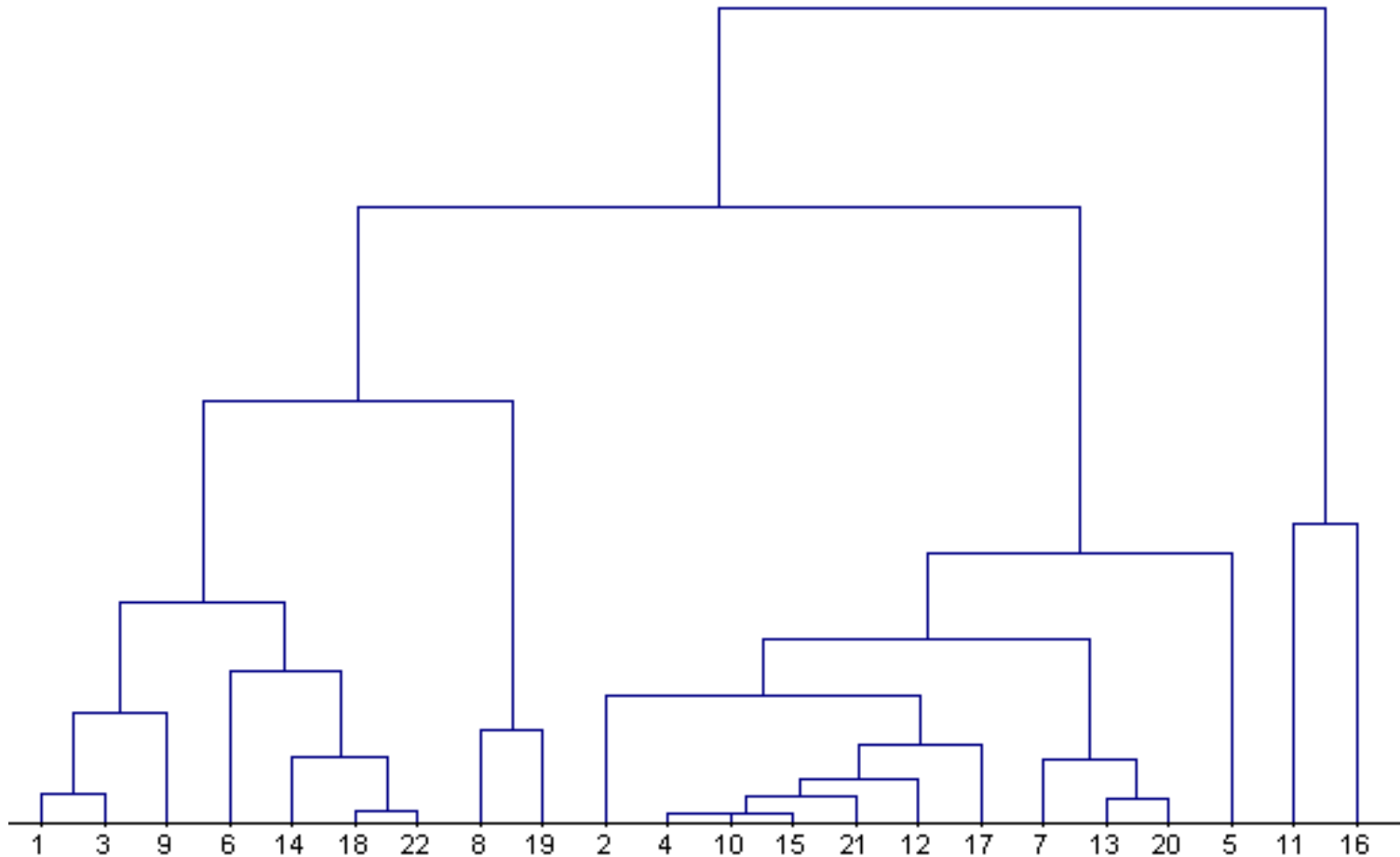
# Definite agglomerative clustering

Basic algorithm – at a certain level, there may be multiple candidates of merging.

Random selection leads to ambiguities.

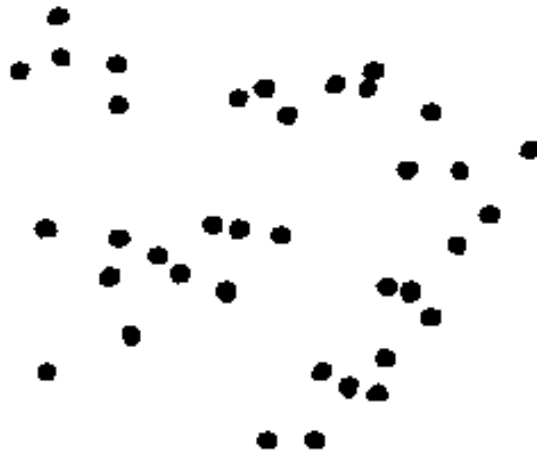
Definite algorithm – **all** such candidates are merged at that level → the number of new clusters emerging at one level may be greater than 1.

# Agglomerative clustering – representation by a clustering tree (dendrogram)



# Basic divisive clustering

1. All points = one cluster



# Basic divisive clustering

1. All points = one cluster
2. Divide the cluster into two parts  $A, B$  such that  $d(A, B)$  is maximized
3. Select a new cluster to split
4. Apply 2 to the selected cluster
5. Repeat 3-4 until the stop constraint is reached

Full search in Step 2 is very expensive –  $O(2^n)$

## Suboptimal Step 2 of divisive clustering

1. Find point  $p$  such that the mean of  $d(p,x)$  is maximized
2. Divide  $C$  into  $A \cup B$ , where  $B = \{p\}$  ( $p$  is a seed point of a new cluster  $B$ )
3. For  $x$  from  $A$  calculate the mean dist  $d(x,A)$  and  $d(x,B)$ . If  $d(x,A) > d(x,B)$  move  $x$  into  $B$ .
4. Repeat 3 for each  $x$  from  $A$ .



## Suboptimal Step 2 of divisive clustering

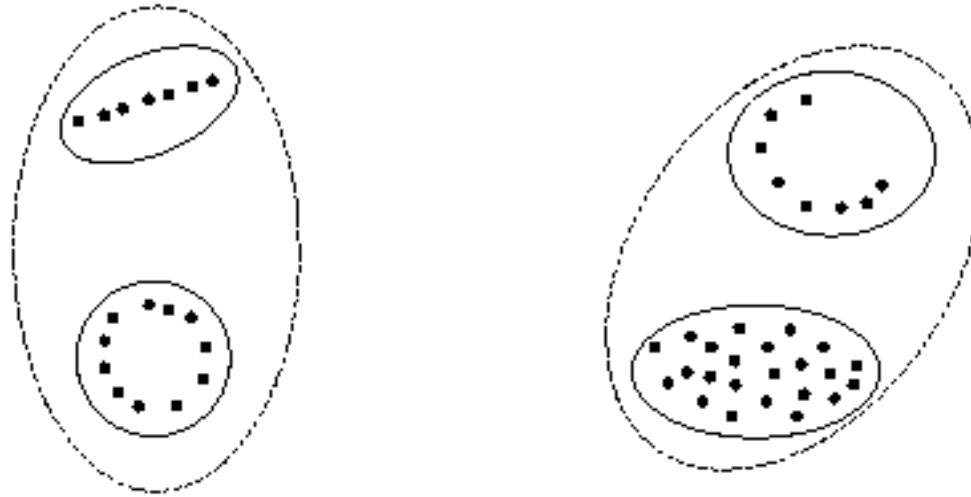
Alternatively, any iterative algorithm for  $N=2$   
can be used ( $N$ -means,  $J$  - minimization, ...)

## Step 3 of divisive clustering

Selecting the next cluster to split

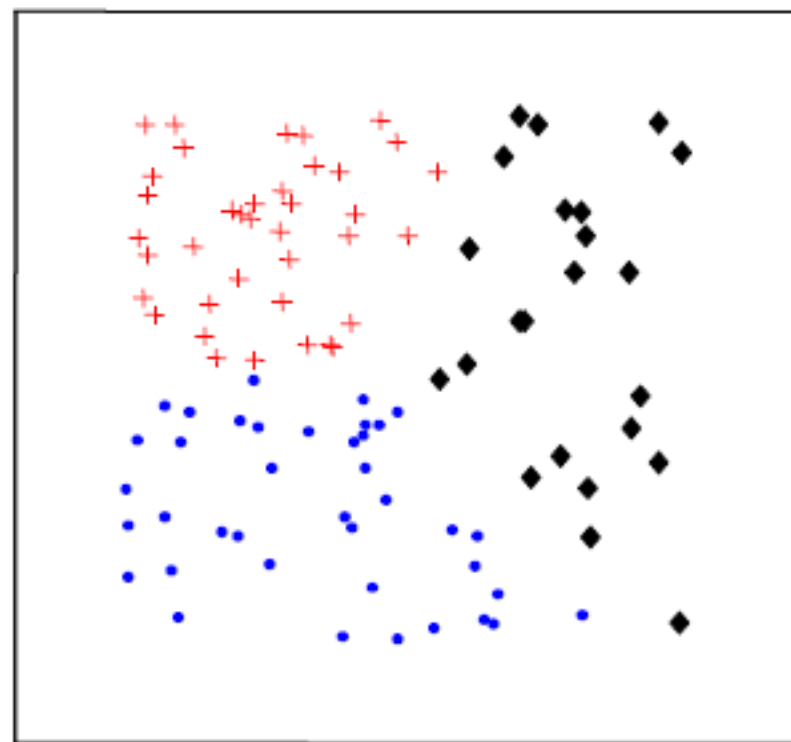
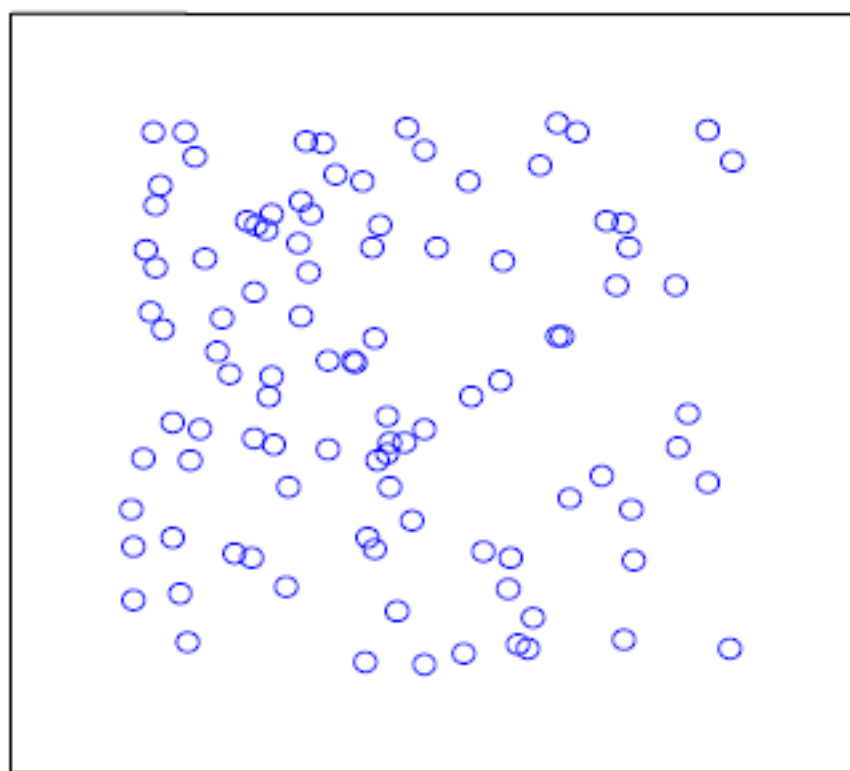
- randomly (when performing complete decomposition)
- by maximum diameter
- by maximum variance
- by maximum  $J$

How many clusters are there?



2 or 4 ?

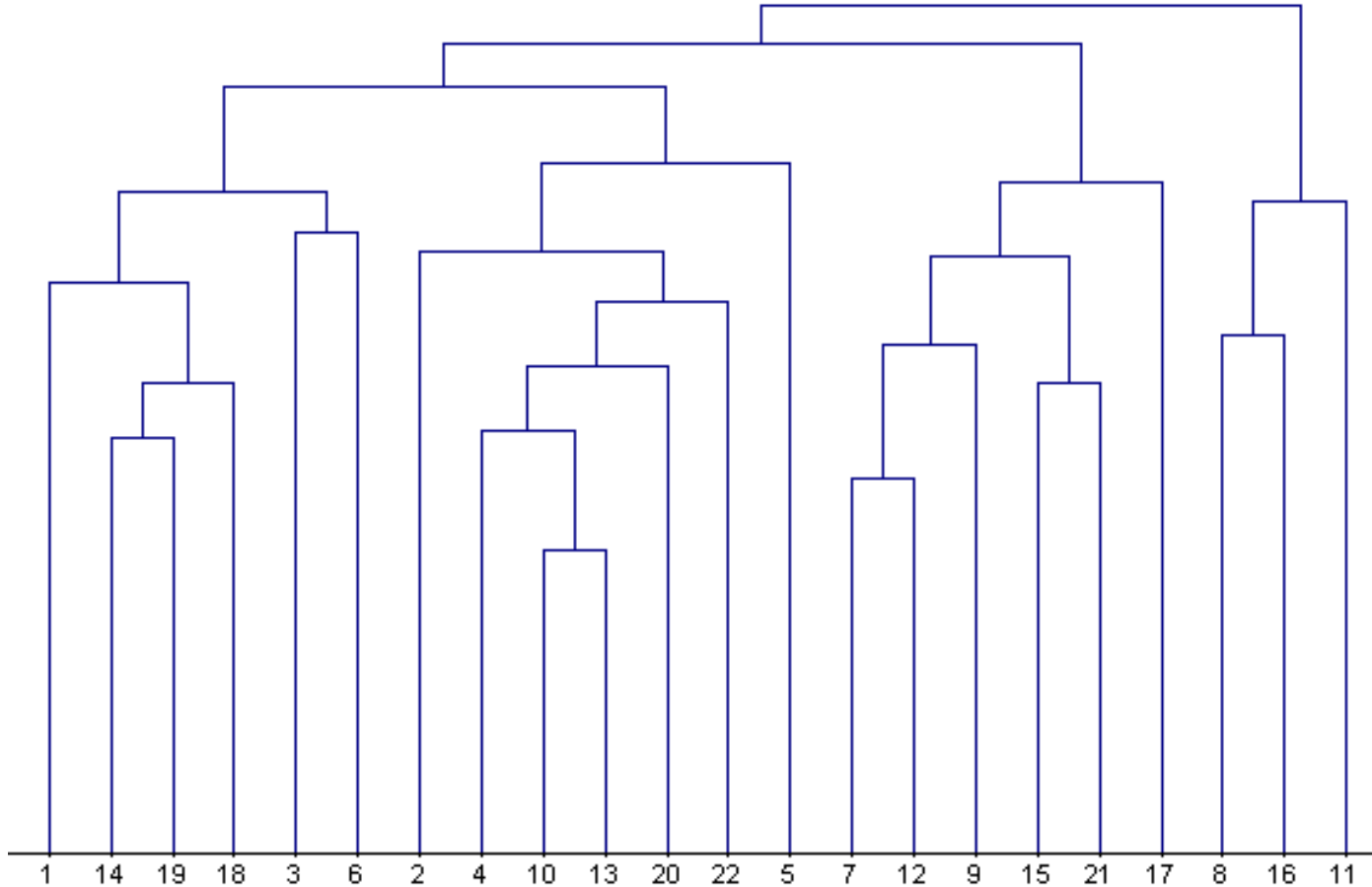
Clustering is a very subjective task



# How many clusters are there?

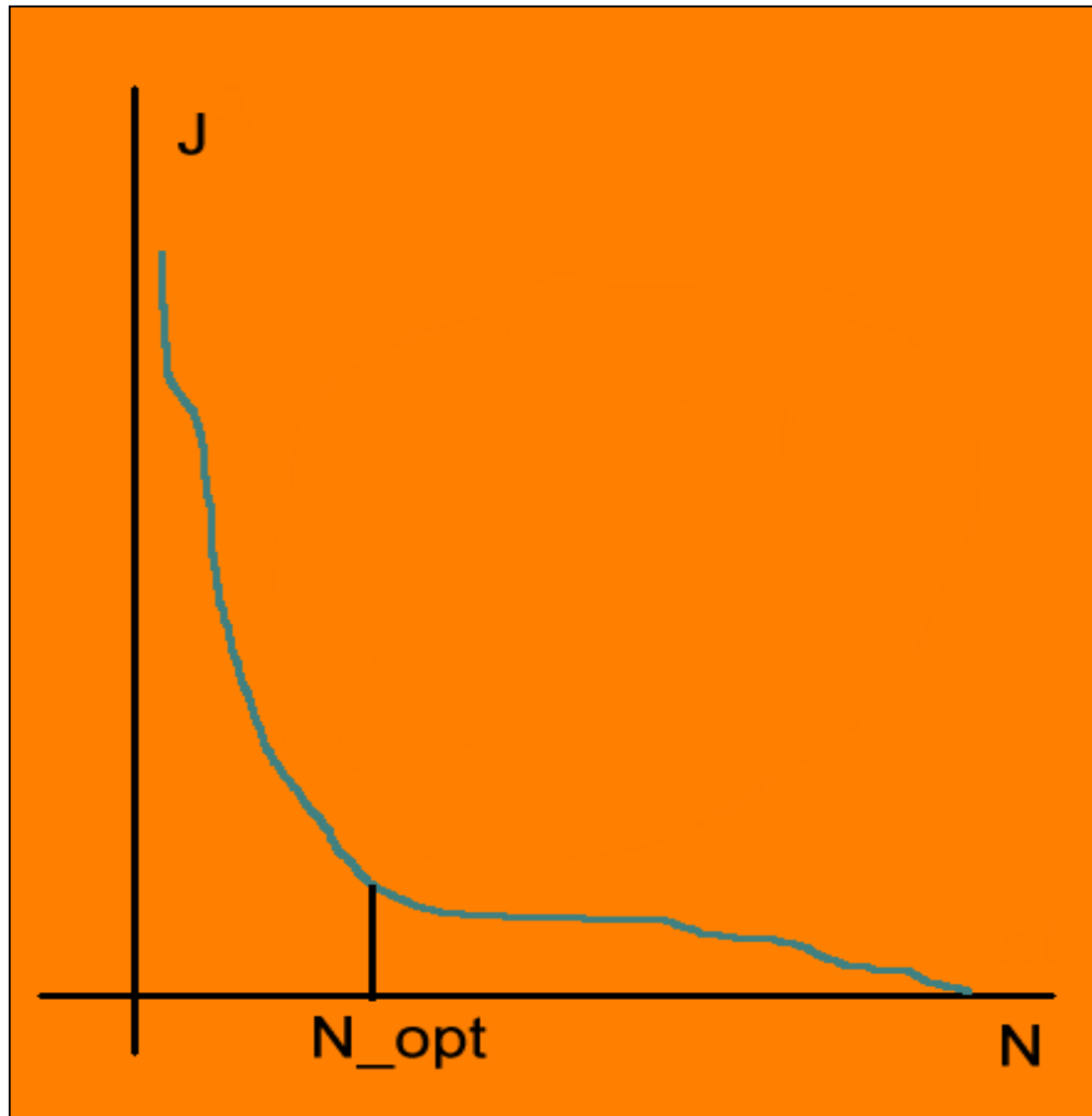
- Difficult to answer even for humans
- “Clustering tendency”, “cluster validity”
- Hierarchical methods –  $N$  can be estimated from the complete dendrogram
- The methods minimizing a cost function –  $N$  can be estimated from the “knees” in  $J$ - $N$  graph

# Life time of the clusters

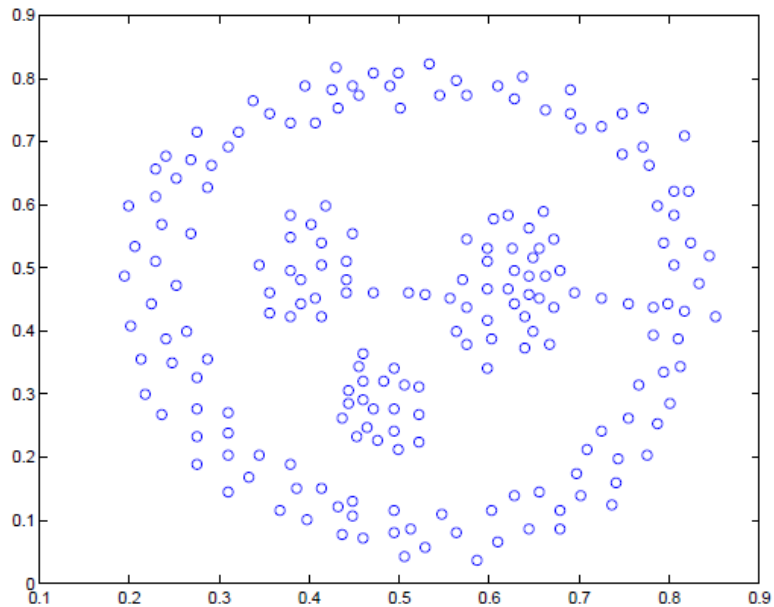


**Optimal number of clusters = 4**

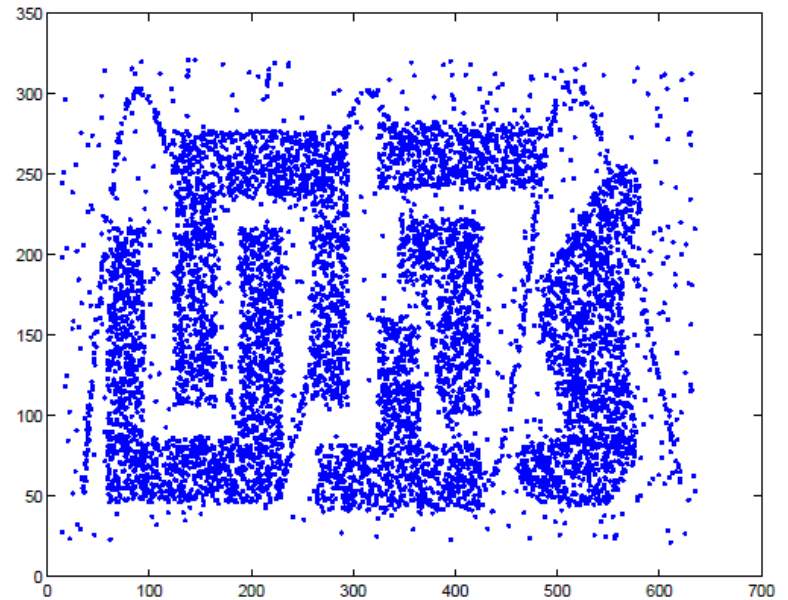
# Optimal number of clusters



# Hybrid clustering with the choice of N



(a)



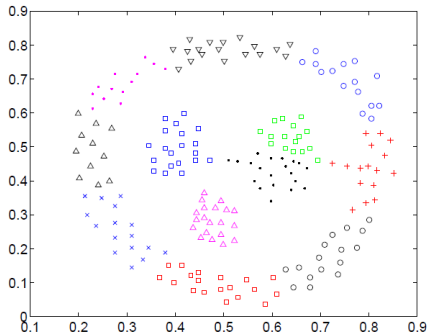
(b)



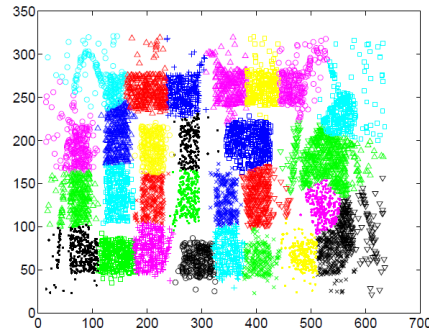
# Hybrid clustering with the choice of N

Iterative

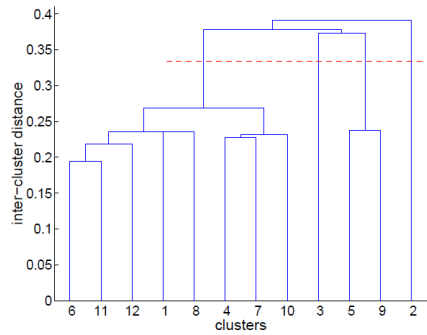
Hierarchical merging



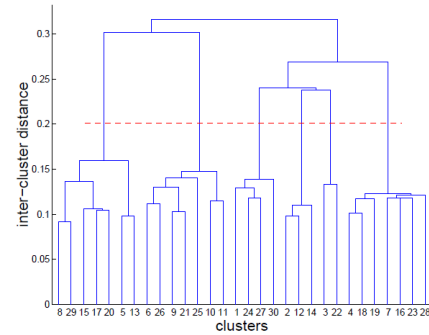
(a)



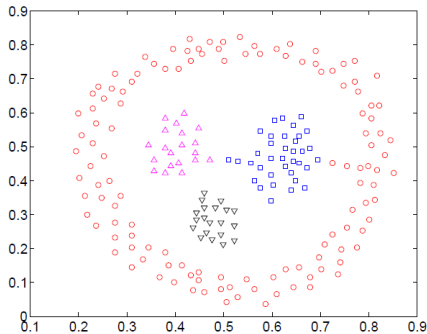
(b)



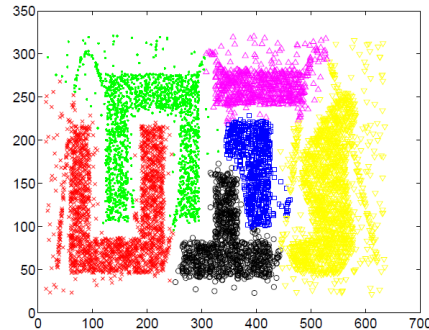
(c)



(d)



(e)



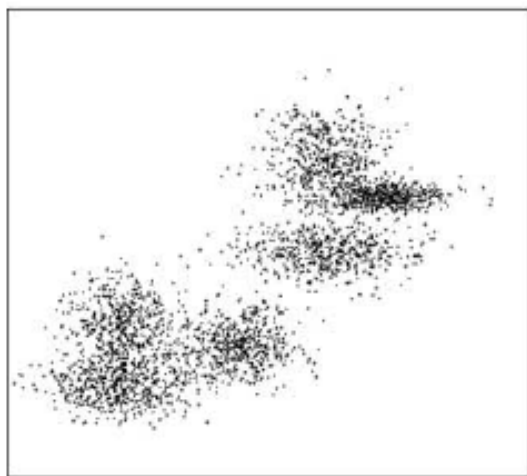
(f)

# Model-based (parametric) clustering

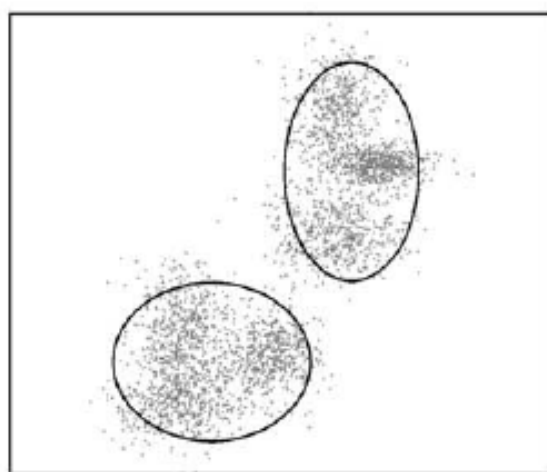
Fitting a Gaussian mixture to the data

$$f(\mathbf{x}) = \sum_{g=1}^G \pi_g \phi(\mathbf{x} | \boldsymbol{\mu}_g, \boldsymbol{\Sigma}_g)$$

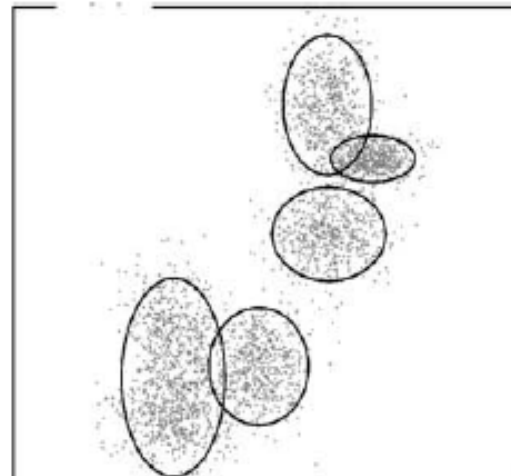
**Problem:** the number of components



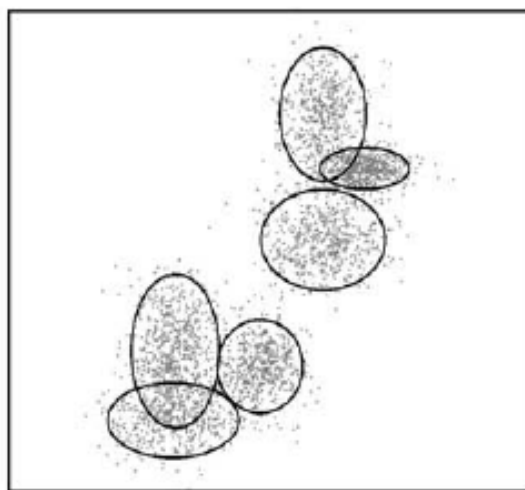
(a) Input data



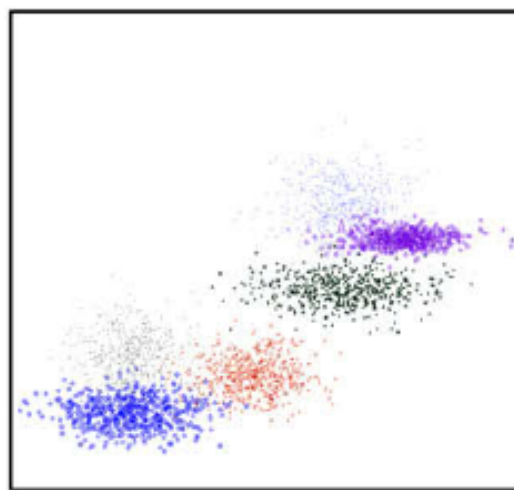
(b) GMM (K=2)



(c) GMM (K=5)



(d) GMM (K=6)

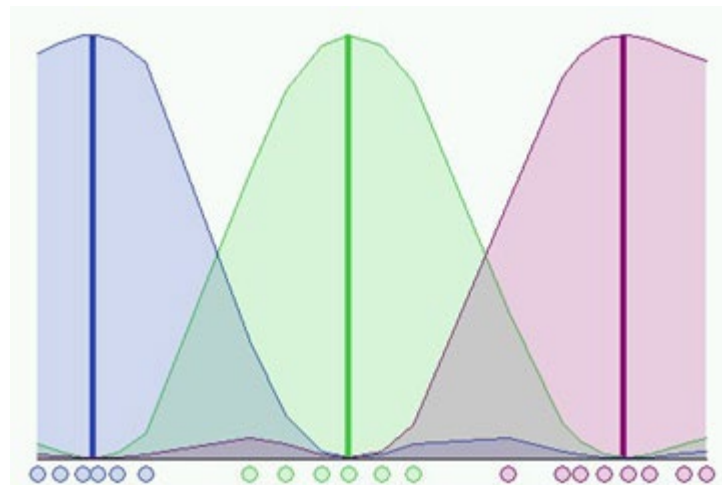


(e) True labels, K = 6

# Fuzzy clustering

Clusters = Fuzzy sets (Set, Mem. f)

Fuzzy C-means



# Other clustering criteria

## Scatter matrices

- between cluster matrix

$$B = \sum_{i=1}^N n_i (\mathbf{m}_i - \mathbf{m})(\mathbf{m}_i - \mathbf{m})^t$$

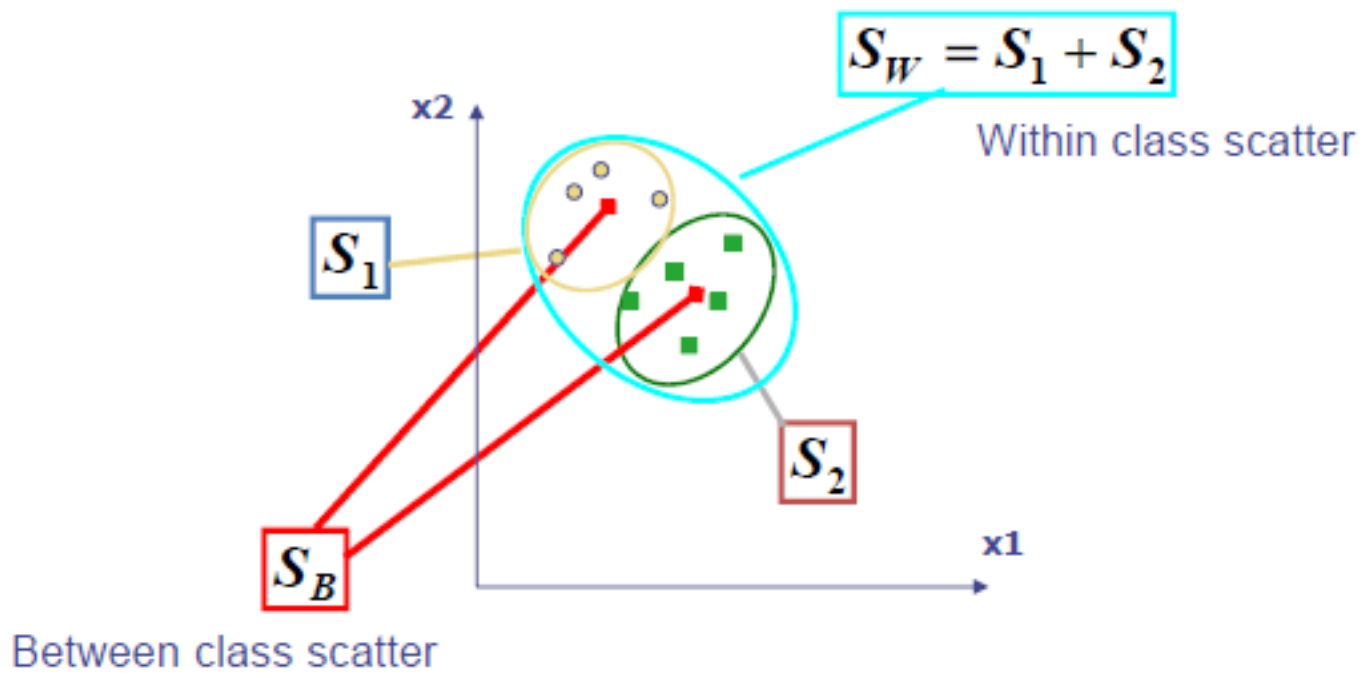
- within cluster matrix

$$W = \sum_{i=1}^N W_i$$

$$W_i = \sum_{\mathbf{x} \in C_i} (\mathbf{x} - \mathbf{m}_i)(\mathbf{x} - \mathbf{m}_i)^t$$

- total scatter matrix

$$T = \sum_{\mathbf{x}} (\mathbf{x} - \mathbf{m})(\mathbf{x} - \mathbf{m})^t$$



# Other clustering criteria

$$\min \operatorname{tr}(W) = \sum_{i=1}^N \operatorname{tr}(W_i) = \sum_{i=1}^N \sum_{\mathbf{x} \in C_i} \|\mathbf{x} - \mathbf{m}_i\|^2 = J$$

$$\min \det(W)$$

$$T = W + B$$

$$\max \operatorname{tr}(W^{-1}B)$$

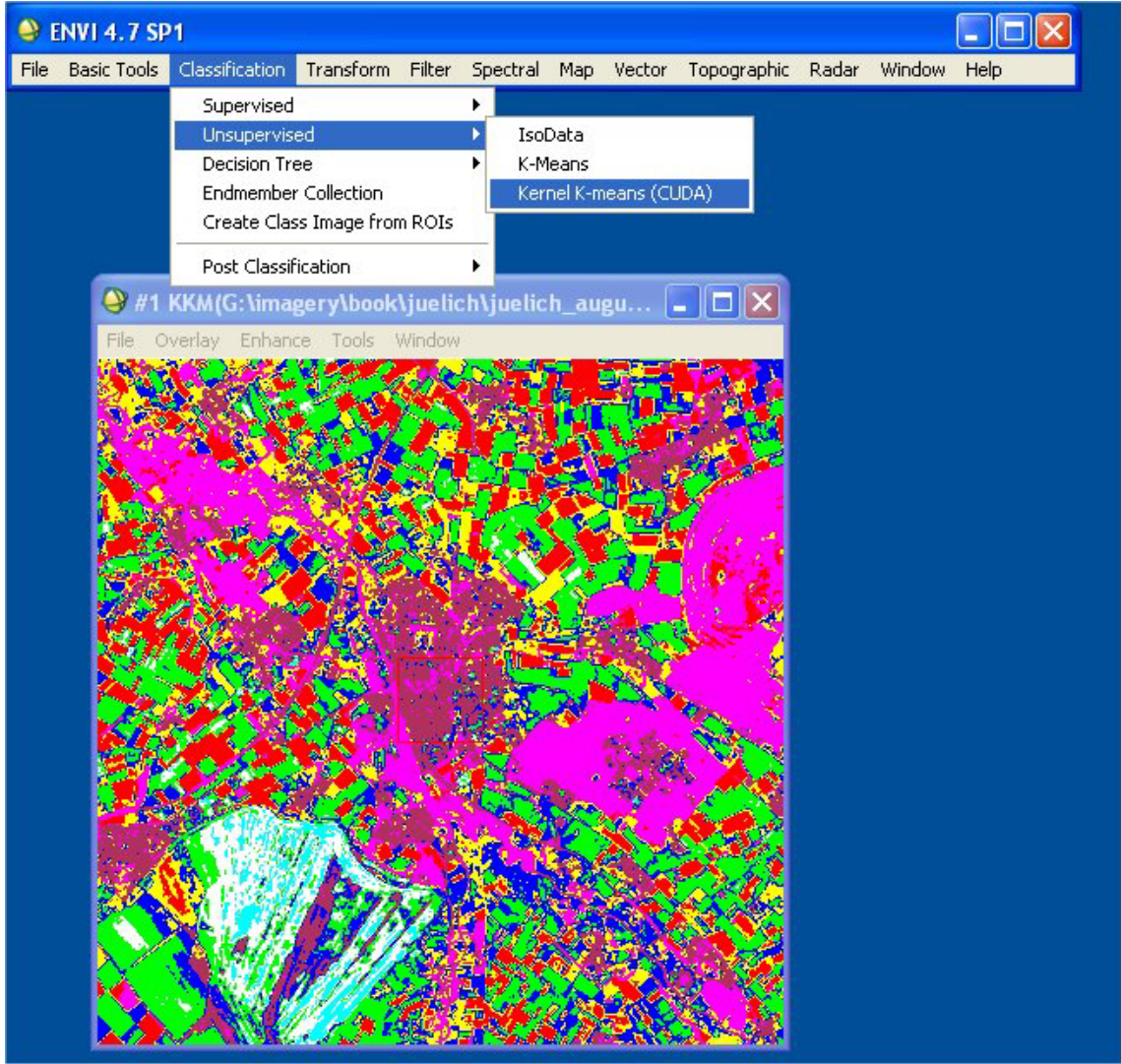
# Applications of clustering in image proc.

- **Segmentation – clustering in color space**
- **Preliminary classification of multispectral images**
- **Clustering in parametric space – RANSAC, image registration and matching**

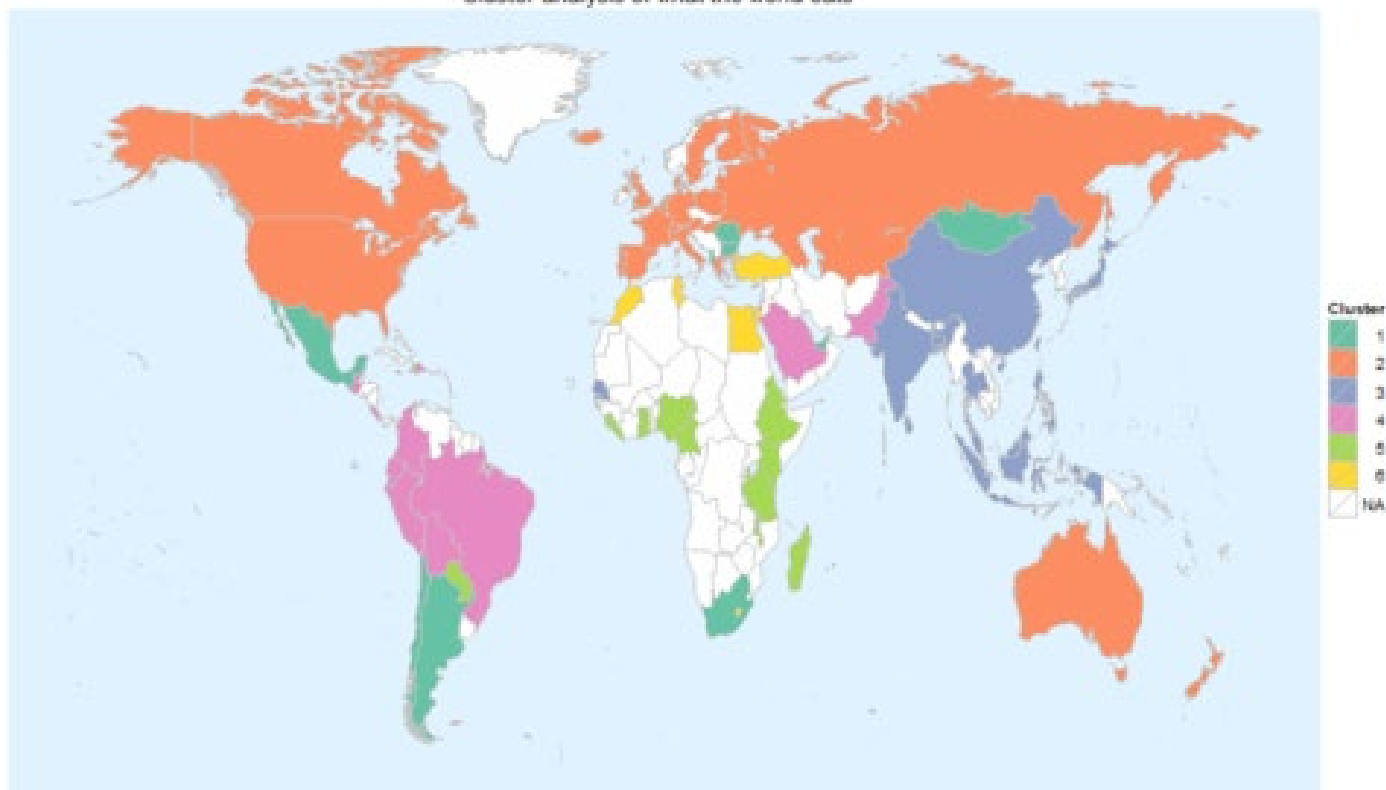
**Numerous applications are outside image processing area**







Cluster analysis of what the world eats





**Thank you !**

**Any questions ?**