# **CS570: Introduction to Data Mining**Basic Clustering

Reading: Chapter 10 Han, Chapter 8 Tan

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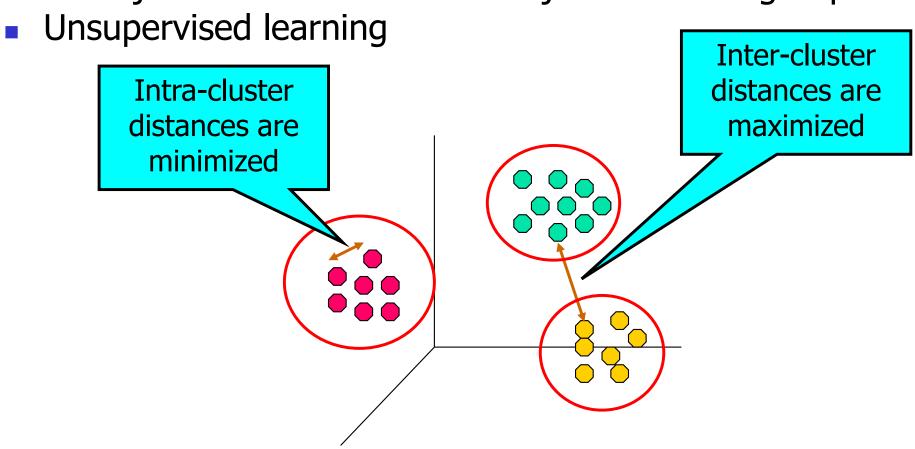
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## Cluster Analysis

- Overview
- Partitioning methods
- Hierarchical methods
- Density-based methods
- Other Methods
- Outlier analysis
- Summary

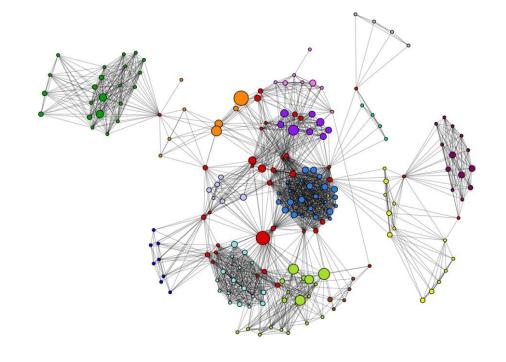
# What is Cluster Analysis?

- Finding groups of objects (clusters)
  - Objects similar to one another in the same group
  - Objects different from the objects in other groups

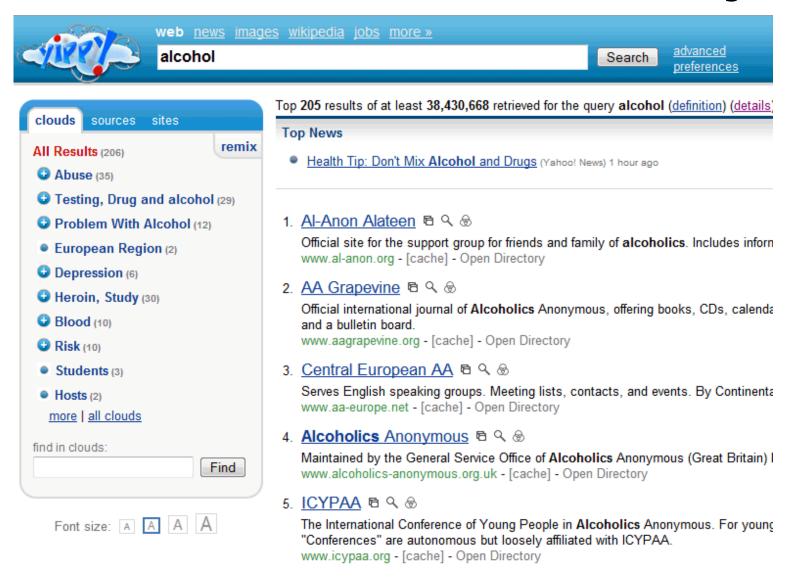


- Marketing research
- Social network analysis

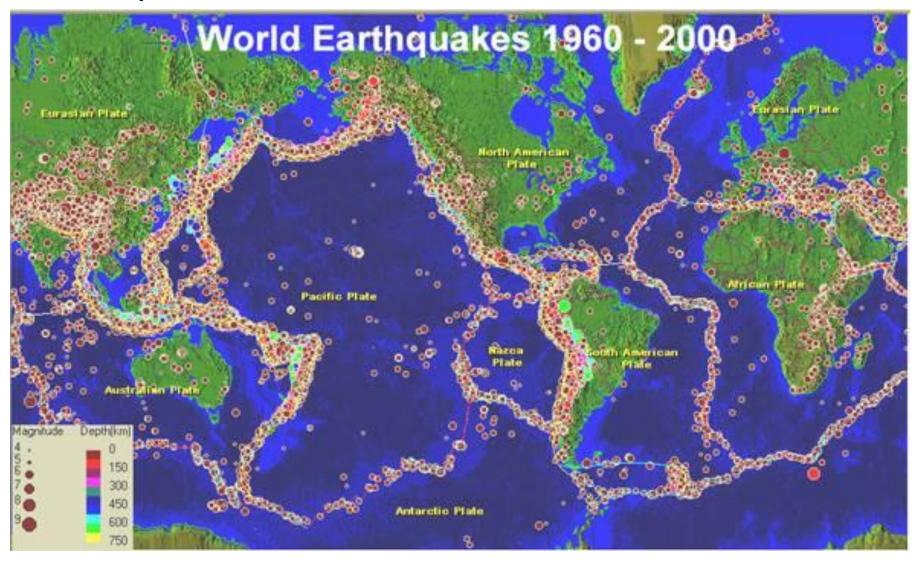




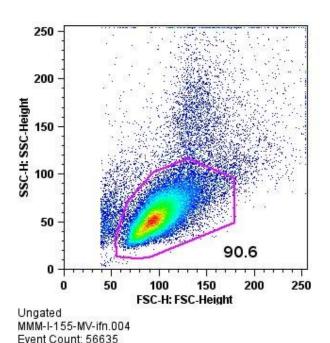
WWW: Documents and search results clustering

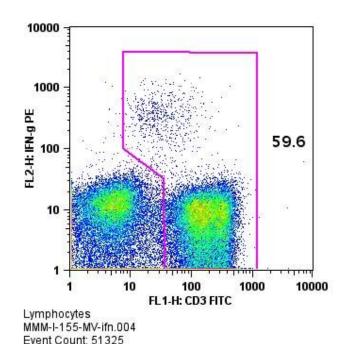


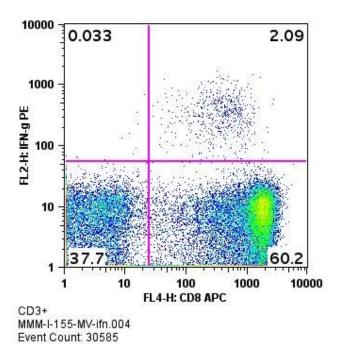
Earthquake studies



- Biology: plants and animals
- Bioinformatics: microarray data, flow cytometry data, genes and sequences





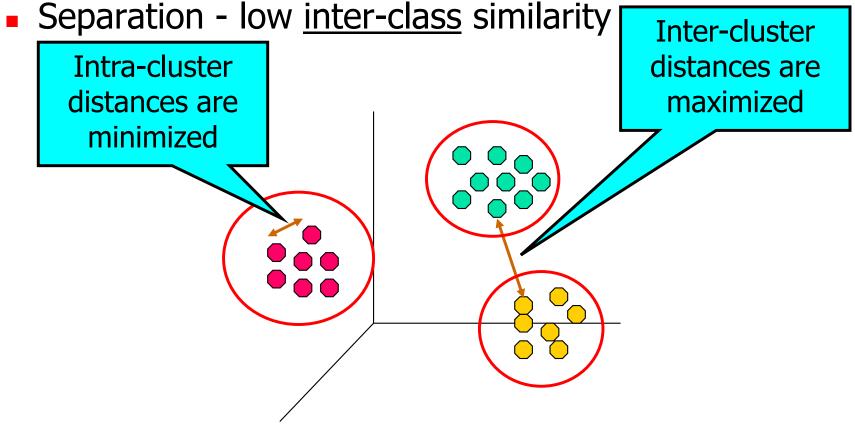


## Requirements of Clustering

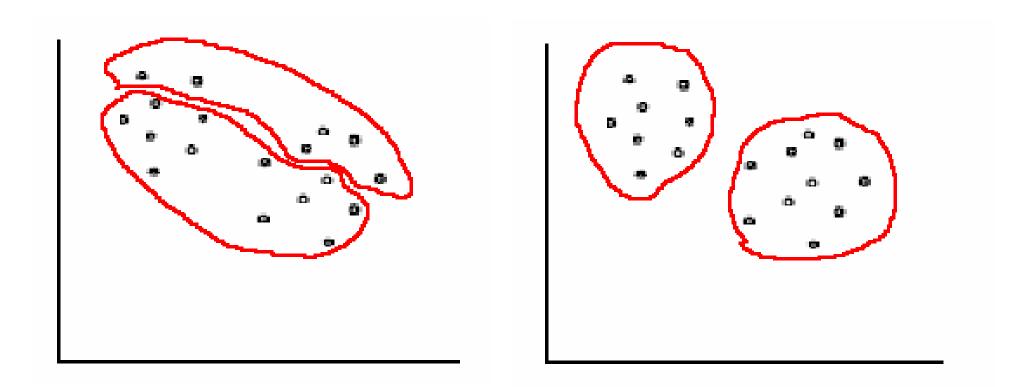
- Quality
- Scalability
- Ability to deal with different types of attributes
- Ability to handle dynamic data
- Ability to deal with noise and outliers
- Ability to deal with high dimensionality
- Minimal requirements for domain knowledge to determine input parameters
- Incorporation of user-specified constraints
- Interpretability and usability

# Quality: What Is Good Clustering?

- Agreement with "ground truth"
- A good clustering will produce high quality clusters with
  - Homogeneity high <u>intra-class</u> similarity



# Bad Clustering vs. Good Clustering



#### Similarity or Dissimilarity between Data Objects

$$\begin{bmatrix} x_{11} & \dots & x_{1f} & \dots & x_{1p} \\ \dots & \dots & \dots & \dots \\ x_{i1} & \dots & x_{if} & \dots & x_{ip} \\ \dots & \dots & \dots & \dots \\ x_{n1} & \dots & x_{nf} & \dots & x_{np} \end{bmatrix}$$
 **•** Euclidean distance

$$d(i,j) = \sqrt{(|x_i - x_j|^2 + |x_i - x_j|^2 + ... + |x_i - x_j|^2)}$$

$$d(i,j) = |x_{i_1} - x_{j_1}| + |x_{i_2} - x_{j_2}| + ... + |x_{i_p} - x_{j_p}|$$

Minkowski distance

$$d(i,j) = \sqrt{(|x_{i1} - x_{j1}|^q + |x_{i2} - x_{j2}|^q + ... + |x_{ip} - x_{jp}|^q)}$$

Chebyshev distance

$$\lim_{p \to \infty} \left( \sum_{i=1}^{n} |x_i - y_i|^p \right)^{\frac{1}{p}} = \max_{i=1}^{n} |x_i - y_i|.$$

#### Other Similarity or Dissimilarity Metrics

$$\begin{bmatrix} x_{11} & \cdots & x_{1f} & \cdots & x_{1p} \\ \cdots & \cdots & \cdots & \cdots \\ x_{i1} & \cdots & x_{if} & \cdots & x_{ip} \\ \cdots & \cdots & \cdots & \cdots \\ x_{n1} & \cdots & x_{nf} & \cdots & x_{np} \end{bmatrix}$$

- Pearson correlation  $r = \frac{\sum_{i=1}^{n} (X_i X)(Y_i Y)}{\sqrt{\sum_{i=1}^{n} (X_i \bar{X})^2} \sqrt{\sum_{i=1}^{n} (Y_i \bar{Y})^2}}.$
- Cosine measure  $\frac{X_i \bullet X_j}{\|X_i\| \cdot \|X_j\|}$
- Jaccard coefficient  $J(A,B) = \frac{|A \cap B|}{|A \cup B|}$ .
- KL divergence, Bregman divergence, ...

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## Different Attribute Types

• To compute  $\begin{vmatrix} x_i - x_i \end{vmatrix}$ 

$$|x_{if} - x_{jf}|$$

- f is numeric (interval or ratio scale)
  - Normalization if necessary

• 
$$f$$
 is ordinal  $Z_{if} = \frac{I_{if} - 1}{M_f - 1}$ 

- f is nominal
  - Mapping function

$$|x_{if} - x_{jf}| = 0$$
 if  $x_{if} = x_{jf}$ , or 1 otherwise

Hamming distance (edit distance) for strings

## Clustering Approaches

#### Partitioning approach:

- Construct various partitions and then evaluate them by some criterion,
  e.g., minimizing the sum of square errors
- Typical methods: k-means, k-medoids, CLARANS

#### Hierarchical approach:

- Create a hierarchical decomposition of the set of data (or objects) using some criterion
- Typical methods: Diana, Agnes, BIRCH, ROCK, CAMELEON

#### Density-based approach:

- Based on connectivity and density functions
- Typical methods: DBSACN, OPTICS, DenClue

#### Others

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## Partitioning Algorithms: Basic Concept

Partitioning method: Construct a partition of a database D of n objects into a set of k clusters, s.t., the sum of squared distance is minimized

$$\sum_{i=1}^k \sum_{p \in C_i} (p - m_i)^2$$

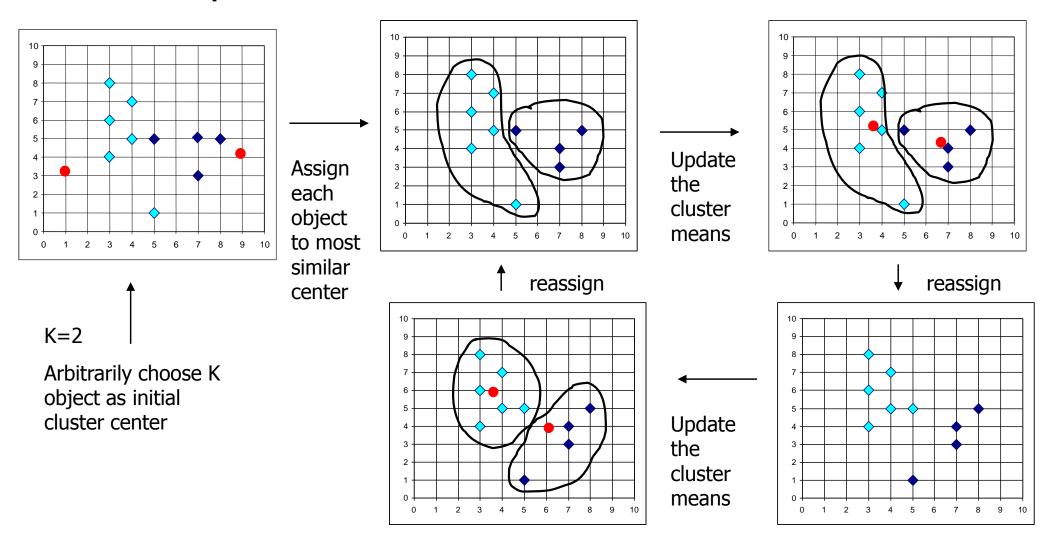
- Given a k, find a partition of k clusters that optimizes the chosen partitioning criterion
  - Global optimal: exhaustively enumerate all partitions
  - Heuristic methods: k-means and k-medoids algorithms
  - <u>k-means</u> (MacQueen'67): Each cluster is represented by the center of the cluster
  - <u>k-medoids</u> or PAM (Partition around medoids) (Kaufman & Rousseeuw'87): Each cluster is represented by one of the objects in the cluster

#### K-Means Clustering: Lloyd Algorithm

- Given k, randomly choose k initial cluster centers
- Partition objects into k nonempty subsets by assigning each object to the cluster with the nearest centroid
- Update centroid, i.e. mean point of the cluster
- Go back to Step 2, stop when no more new assignment

## The K-Means Clustering Method

#### Example



#### *K*-means Clustering – Details

- Initial centroids are often chosen randomly.
- The centroid is (typically) the mean of the points in the cluster.
- 'Closeness' is measured by Euclidean distance, cosine similarity, correlation, etc.
- Most of the convergence happens in the first few iterations.
  - Often the stopping condition is changed to 'Until relatively few points change clusters'
- Complexity is O(tkn)

*n* is # objects, *k* is # clusters, and *t* is # iterations.

#### Comments on the *K-Means* Method

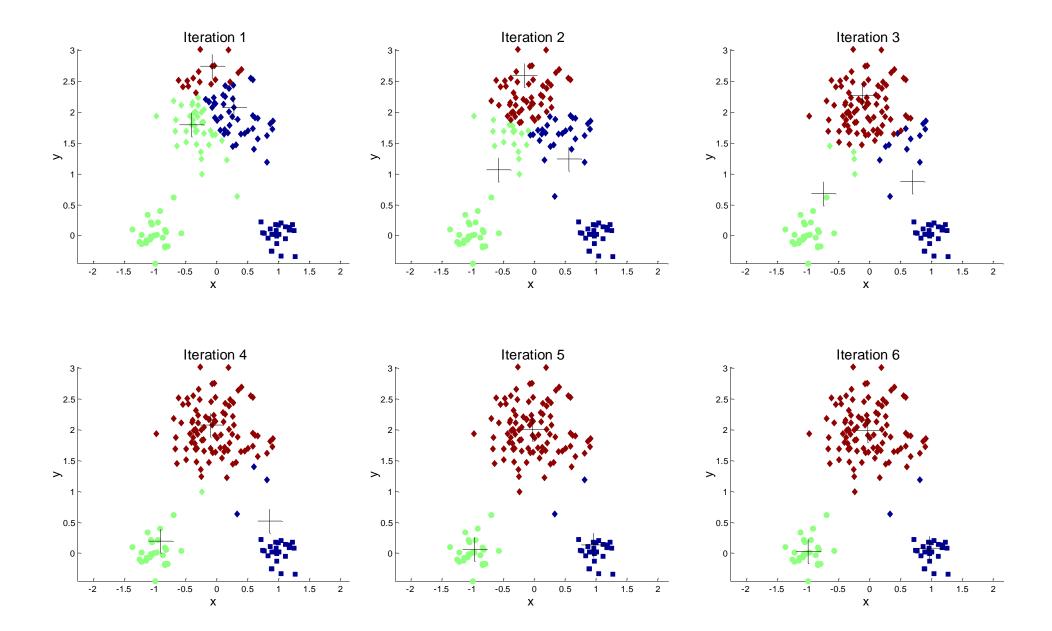
#### Strength

- Simple and works well for "regular" (spherical shape) disjoint clusters
- Relatively efficient and scalable (normally, k, t << n)
- Effective for small to medium size data sets

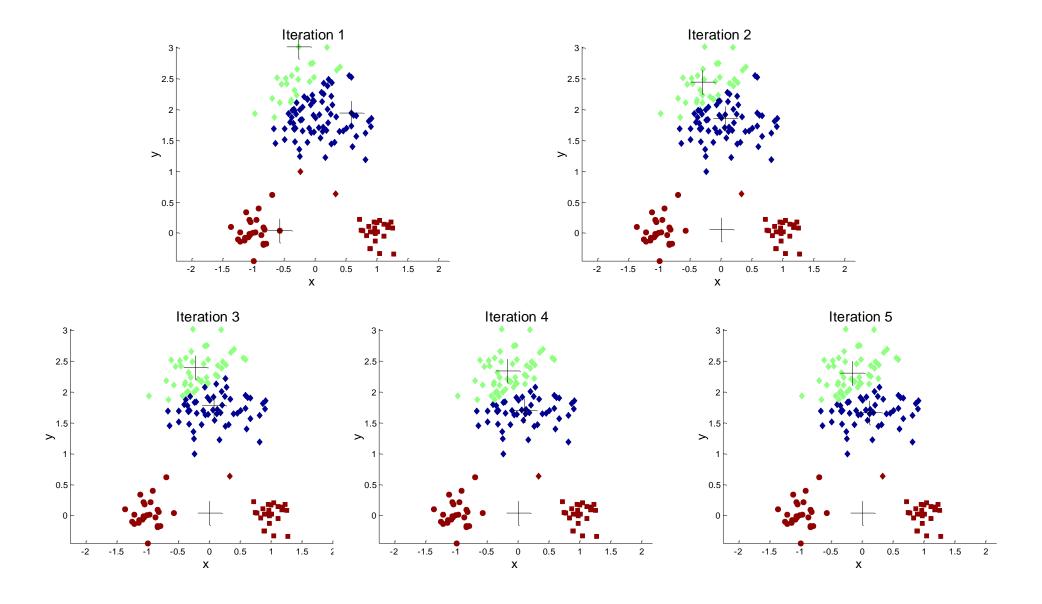
#### Weakness

- Need to specify k, the number of clusters, in advance
- Depending on initial centroids, may terminate at a local optimum
  - Potential solutions
- Unable to handle noisy data and outliers
- Not suitable for clusters of
  - Different sizes
  - Non-convex shapes

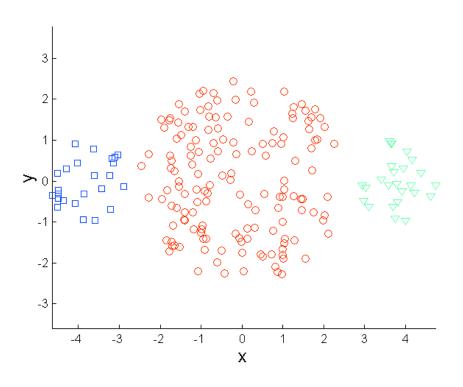
#### Importance of Choosing Initial Centroids – Case 1

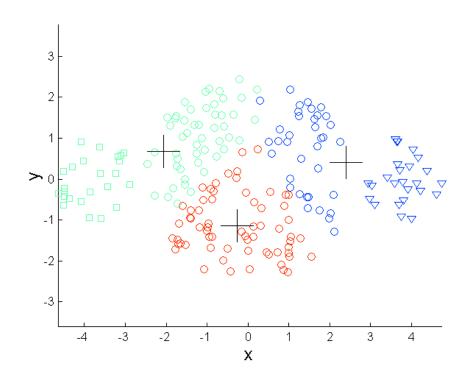


#### Importance of Choosing Initial Centroids – Case 2



#### Limitations of K-means: Differing Sizes

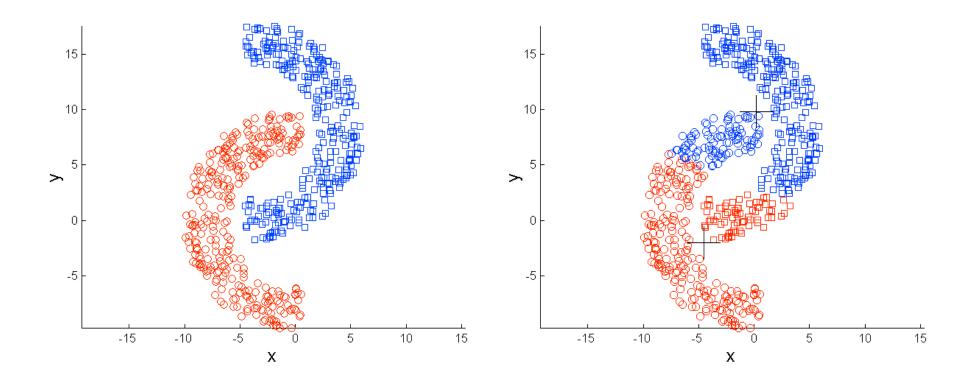




**Original Points** 

K-means (3 Clusters)

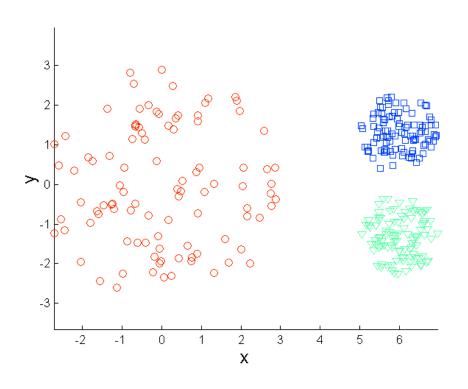
#### Limitations of K-means: Non-convex Shapes

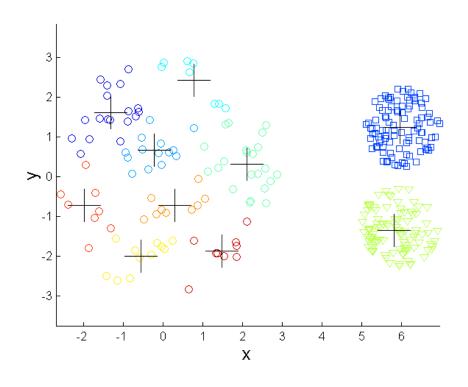


**Original Points** 

K-means (2 Clusters)

## Overcoming K-means Limitations

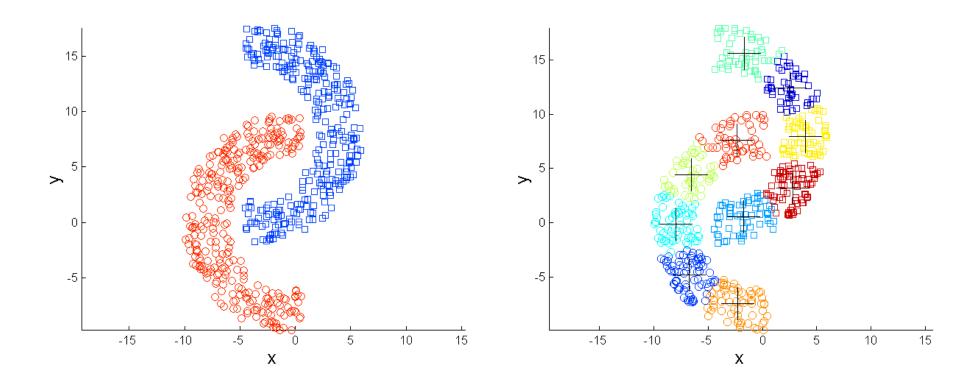




**Original Points** 

K-means Clusters

#### Overcoming K-means Limitations



**Original Points** 

K-means Clusters

#### Variations of the *K-Means* Method

- A few variants of the k-means which differ in
  - Selection of the initial k means
  - Dissimilarity calculations
  - Strategies to calculate cluster means
- Handling categorical data: k-modes (Huang'98)
  - Replacing means of clusters with <u>modes</u>
  - Using new dissimilarity measures to deal with categorical objects
  - Using a <u>frequency</u>-based method to update modes of clusters
  - A mixture of categorical and numerical data: k-prototype method

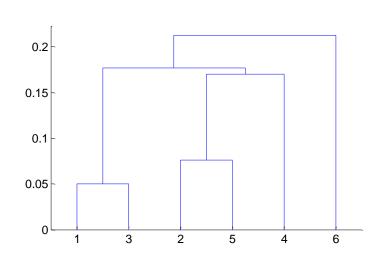
## Cluster Analysis

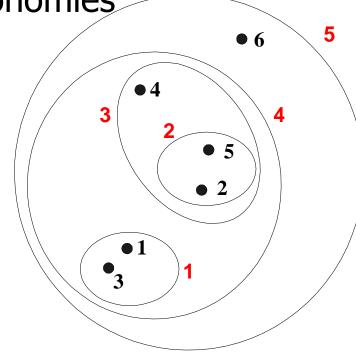
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# **Hierarchical Clustering**

- Produces a set of nested clusters organized as a hierarchical tree
- Can be visualized as a dendrogram, a tree like diagram
  - Clustering obtained by cutting at desired level
- Do not have to assume any particular number of clusters

May correspond to meaningful taxonomies





## **Hierarchical Clustering**

- Two main types of hierarchical clustering
  - Agglomerative:
    - Start with the points as individual clusters
    - At each step, merge the closest pair of clusters until only one cluster (or k clusters) left

#### Divisive:

- Start with one, all-inclusive cluster
- At each step, split a cluster until each cluster contains a point (or there are k clusters)

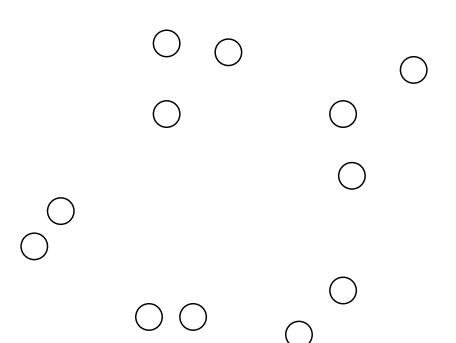
## **Agglomerative Clustering Algorithm**

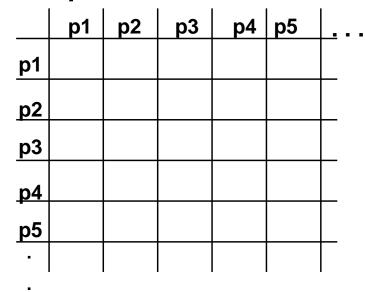
- 1. Compute the proximity matrix
- 2. Let each data point be a cluster
- 3. Repeat
- 4. Merge the two closest clusters
- 5. Update the proximity matrix
- 6. **Until** only a single cluster remains

## **Starting Situation**

Start with clusters of individual points and a

proximity matrix

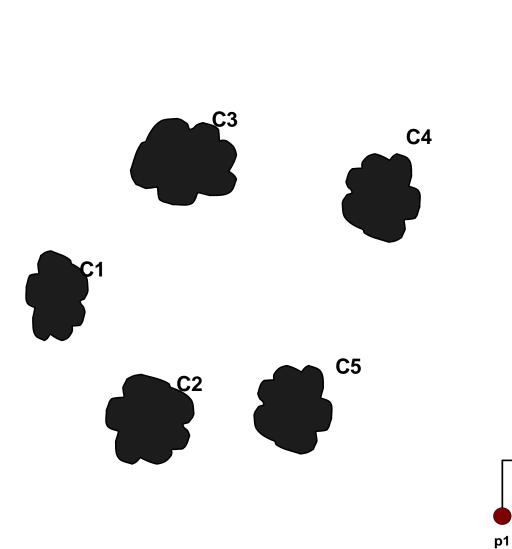


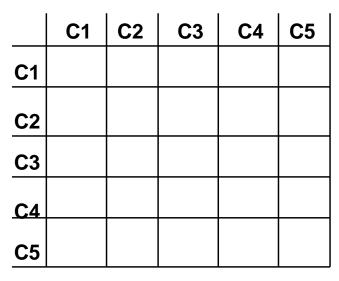


**Proximity Matrix** 

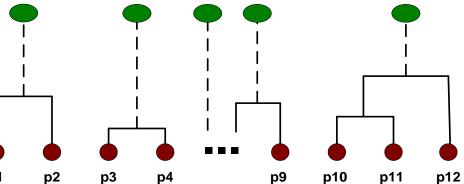


## **Intermediate Situation**

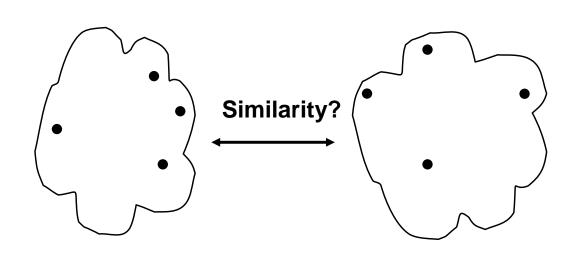




**Proximity Matrix** 



## How to Define Inter-Cluster Similarity

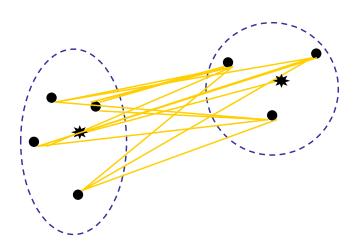


	<b>p</b> 1	<b>p2</b>	рЗ	p4	р5	_ <u>.</u>
<b>p1</b>						
p2						
рЗ						
<b>p4</b>						
р5						

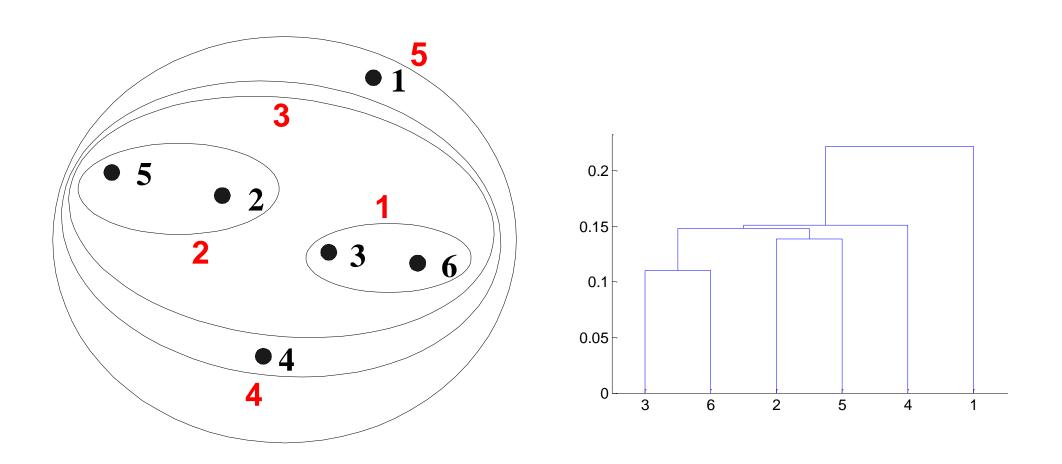
**Proximity Matrix** 

#### Distance Between Clusters

- Single Link: smallest distance between points
- Complete Link: largest distance between points
- Average Link: average distance between points
- Centroid: distance between centroids



# Hierarchical Clustering: MIN

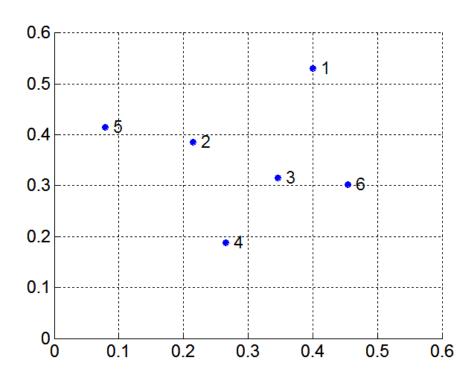


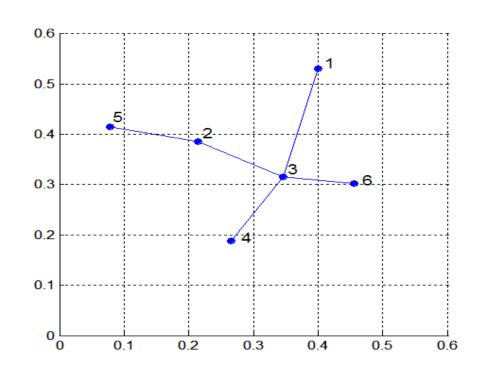
**Nested Clusters** 

**Dendrogram** 

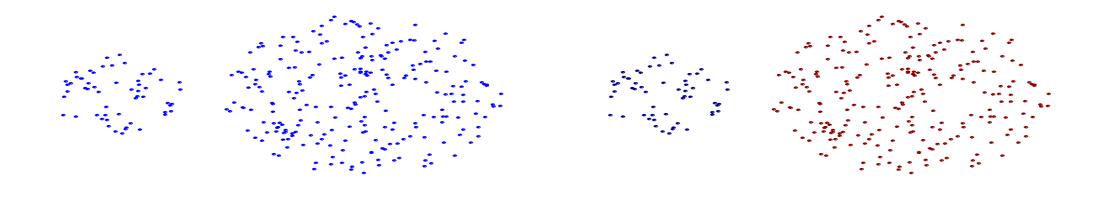
# MST (Minimum Spanning Tree)

- An aggolomerative algorithm using minimum distance can be also called a minimal spanning tree (MST) algorithm
- Start with a tree that consists of any point
- In successive steps, look for the closest pair of points (p, q) such that one point (p) is in the current tree but the other (q) is not; add q to the tree and put an edge between p and q





# Strength of MIN

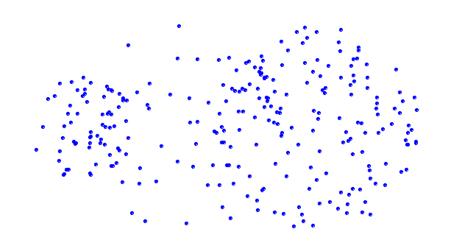


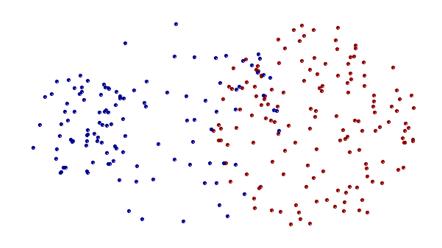
**Original Points** 

**Two Clusters** 

Can handle non-elliptical shapes

## **Limitations of MIN**



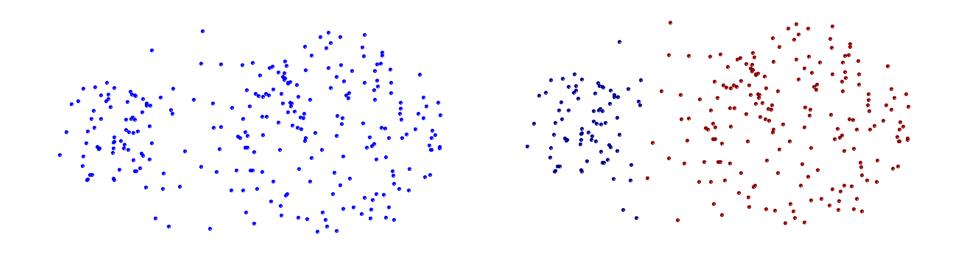


**Original Points** 

**Two Clusters** 

Sensitive to noise and outliers

# Strength of MAX

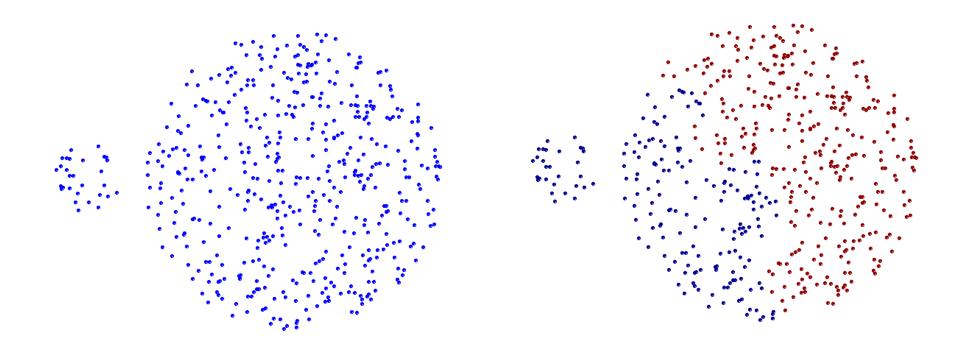


**Original Points** 

**Two Clusters** 

Less susceptible to noise and outliers

## Limitations of MAX



**Original Points** 

**Two Clusters** 

- Tends to break large clusters
- Biased towards globular clusters

## Hierarchical Clustering: Group Average

 Compromise between Single and Complete Link

- Strengths
  - Less susceptible to noise and outliers
  - Can handle categorical and numerical data
- Limitations
  - Biased towards globular clusters

#### Hierarchical Clustering: Major Weaknesses

- Do not scale well (N: number of points)
  - Space complexity: O(N²)
  - Time complexity: O(N³)

O(N<sup>2</sup> log(N)) for some cases/approaches

- Cannot undo what was done previously
- Quality varies in terms of distance measures
  - MIN (single link): susceptible to noise/outliers
  - MAX/GROUP AVERAGE: may not work well with nonglobular clusters

# Cluster Analysis

- Overview
- Partitioning methods
- Hierarchical methods and graph-based methods
  - Classical methods
  - Recent methods
- Density-based methods
- Other Methods
- Outlier analysis
- Summary

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