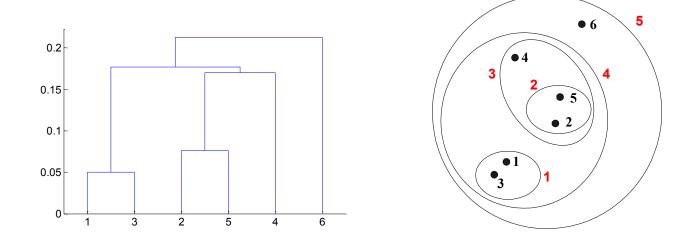
### Data Mining Cluster Analysis: Basic Concepts and Algorithms

### Lecture Notes for Chapter 7

# Introduction to Data Mining, 2<sup>nd</sup> Edition by Tan, Steinbach, Karpatne, Kumar

## **Hierarchical Clustering**

- Produces a set of nested clusters organized as a hierarchical tree
- Can be visualized as a dendrogram
  - A tree like diagram that records the sequences of merges or splits



# **Strengths of Hierarchical Clustering**

- Do not have to assume any particular number of clusters
  - Any desired number of clusters can be obtained by **'cutting' the dendrogram** at the proper level
- They may correspond to meaningful taxonomies
  - Example in biological sciences (e.g., animal kingdom, phylogeny reconstruction, ...)

# **Hierarchical Clustering**

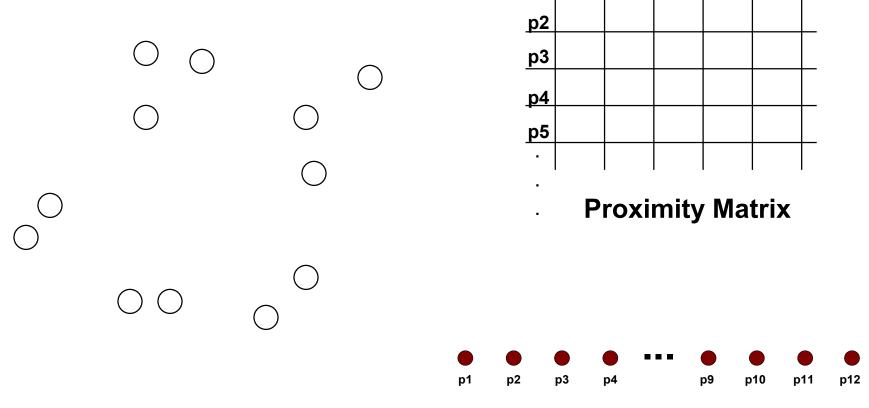
- Two main types of hierarchical clustering
  - Agglomerative:
    - Start with the points as individual clusters
    - At each step, merge the closest pair of clusters until only one cluster (or k clusters) left
  - Divisive:
    - Start with one, all-inclusive cluster
    - At each step, split a cluster until each cluster contains an individual point (or there are k clusters)
- Traditional hierarchical algorithms use a similarity or distance matrix
  - Merge or split one cluster at a time

# **Agglomerative Clustering Algorithm**

- Most popular hierarchical clustering technique
- Basic algorithm is straightforward
  - 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
- Key operation is the computation of the proximity of two clusters
  - Different approaches to defining the distance between clusters distinguish the different algorithms

## **Starting Situation**

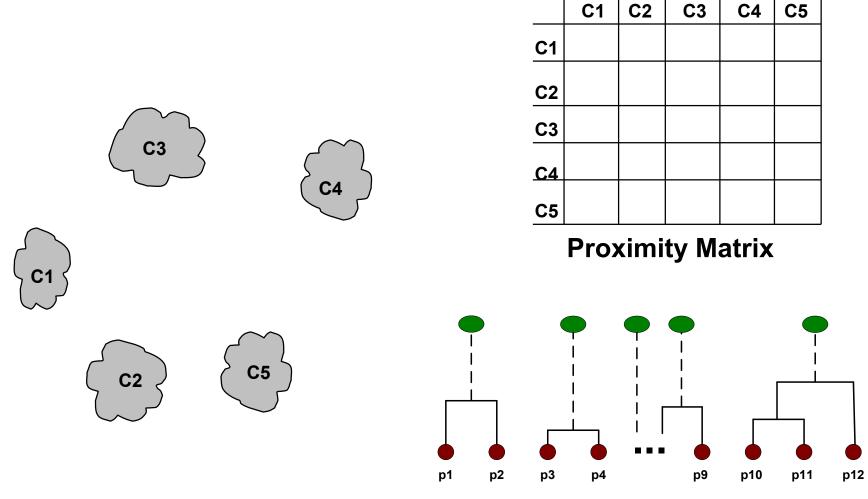
Start with clusters of individual points and a proximity matrix



р1

### **Intermediate Situation**

• After some merging steps, we have some clusters



## **Intermediate Situation**

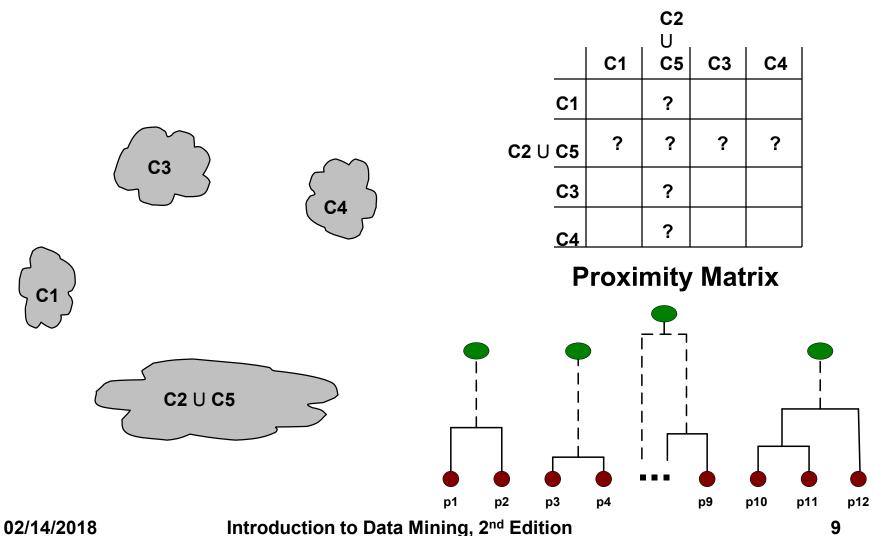
 We want to merge the two closest clusters (C2 and C5) and update the proximity matrix.
| c1 | c2 | c3 | c4 | c5 |

**C1** 

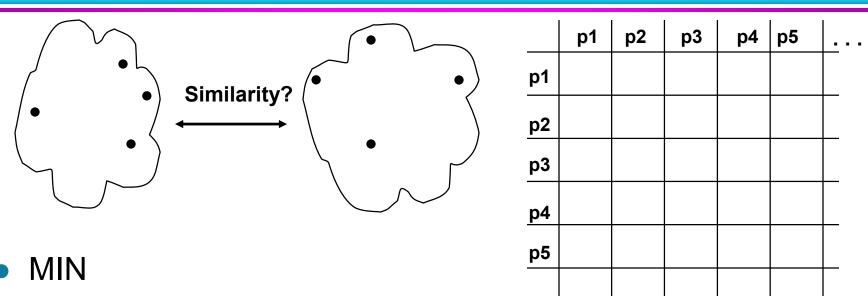
**C2 C**3 **C**3 **C4 C4 C5 Proximity Matrix C1 C5 C2** p2 p10 p11 p1 p3 p4 p9 p12

## **After Merging**

The question is "How do we update the proximity matrix?"

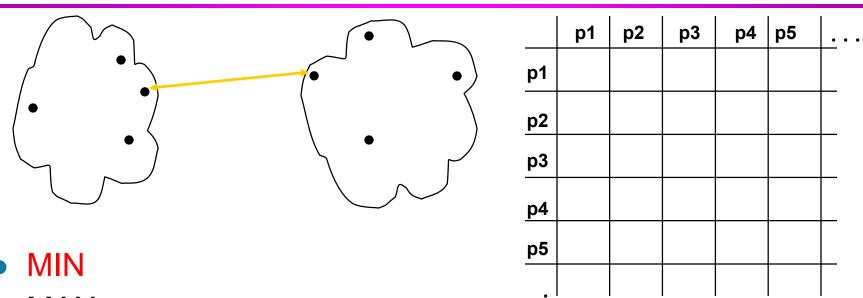


#### **How to Define Inter-Cluster Distance**



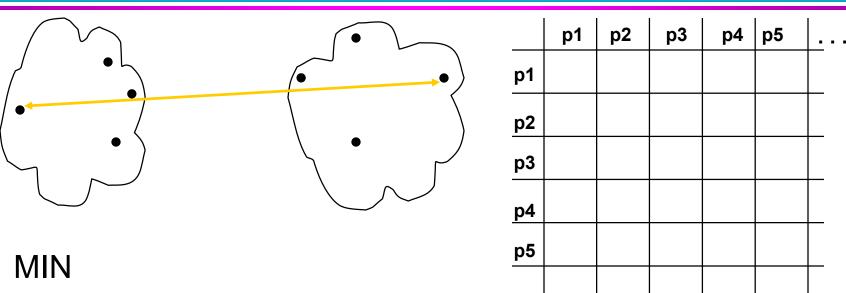
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
  - Ward's Method uses squared error

#### **Proximity Matrix**



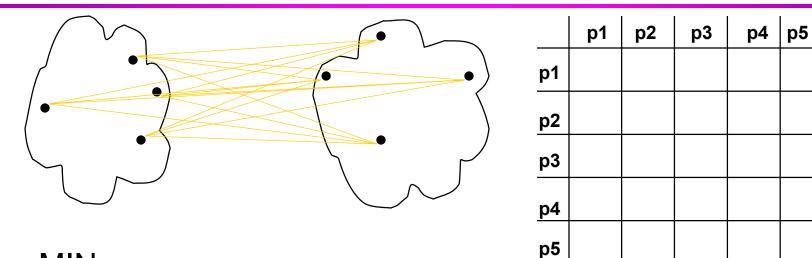
- MAX
- Group Average
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**Proximity Matrix** 



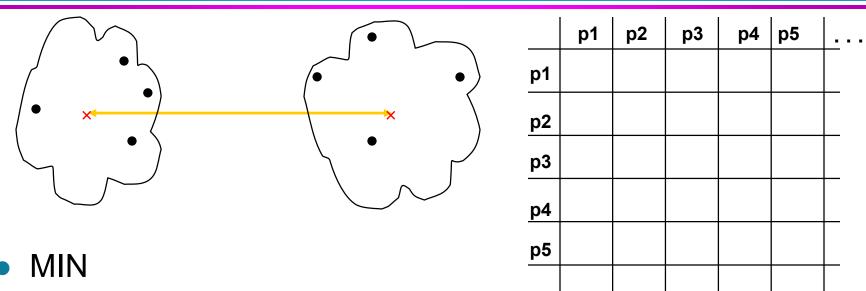
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
  - Ward's Method uses squared error

#### **Proximity Matrix**



- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
  - Ward's Method uses squared error

**Proximity Matrix** 

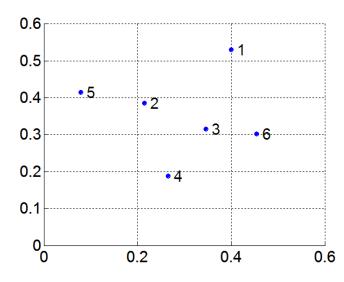


- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
  - Ward's Method uses squared error

**Proximity Matrix** 

## **MIN or Single Link**

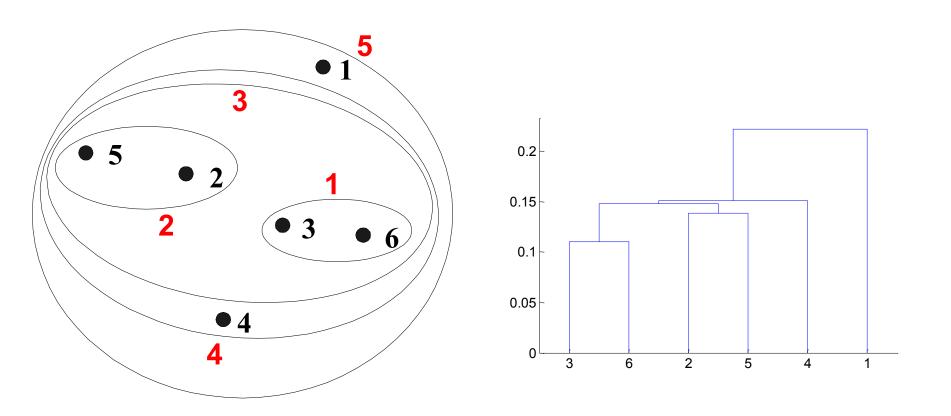
- Proximity of two clusters is based on the two closest points in the different clusters
  - Determined by one pair of points, i.e., by one link in the proximity graph
- Example:



#### **Distance Matrix:**

	p1	p2	p3	p4	p5	p6
p1	0.00	0.24	0.22	0.37	0.34	0.23
p2	0.24	0.00	0.15	0.20	0.14	0.25
p3	0.22	0.15	0.00	0.15	0.28	0.11
p4	0.37	0.20	0.15	0.00	0.29	0.22
p5	0.34	0.14	0.28	0.29	0.00	0.39
p6	0.23	0.25	0.11	0.22	0.39	0.00

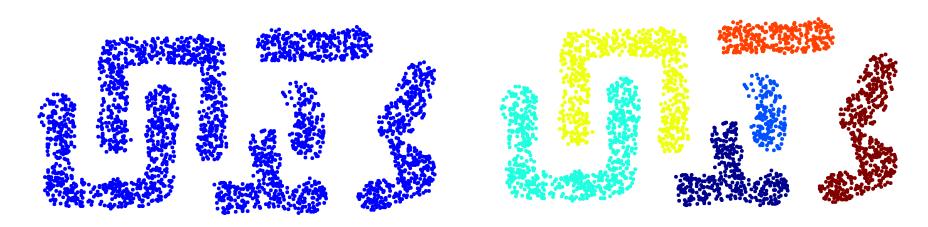
## **Hierarchical Clustering: MIN**



#### **Nested Clusters**

Dendrogram

### **Strength of MIN**



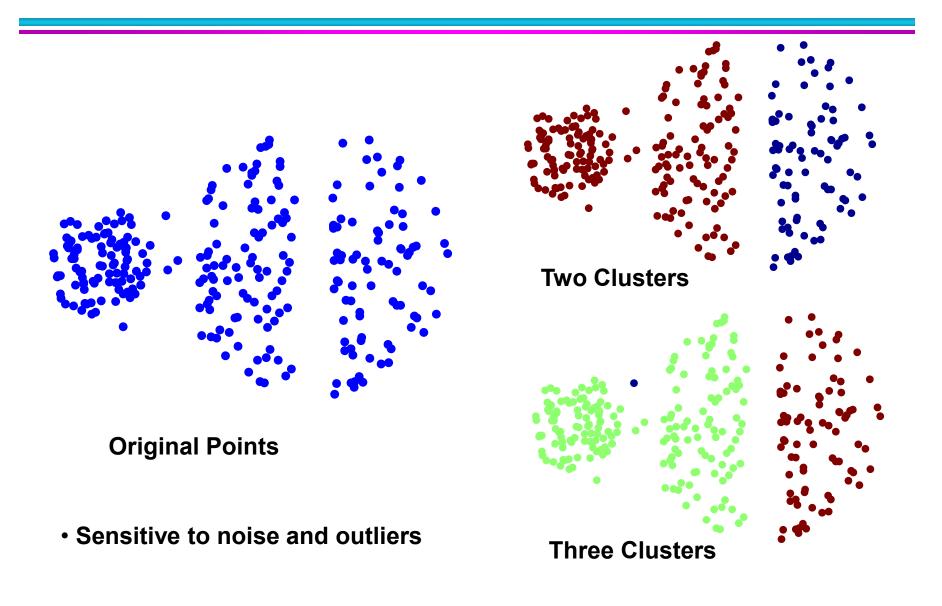
#### **Original Points**

**Six Clusters** 

Can handle non-elliptical shapes

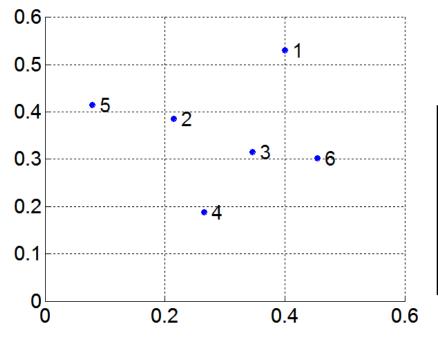
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## **Limitations of MIN**



#### **MAX or Complete Linkage**

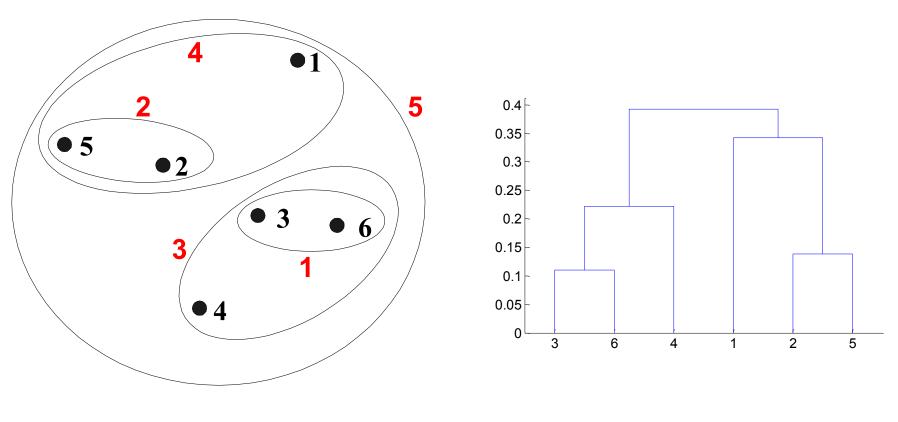
- Proximity of two clusters is based on the two most distant points in the different clusters
  - Determined by all pairs of points in the two clusters



#### **Distance Matrix:**

	p1	p2	p3	p4	p5	p6
p1	0.00	0.24	0.22	0.37	0.34	0.23
p2	0.24	0.00	0.15	0.20	0.14	0.25
p3	0.22	0.15	0.00	0.15	0.28	0.11
p4	0.37	0.20	0.15	0.00	0.29	0.22
p5	0.34	0.14	0.28	0.29	0.00	0.39
p6	0.23	0.25	0.11	0.22	0.39	0.00

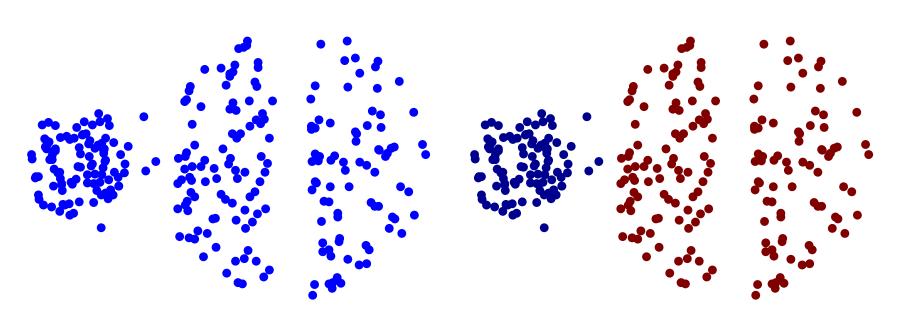
### **Hierarchical Clustering: MAX**



**Nested Clusters** 

Dendrogram

### **Strength of MAX**



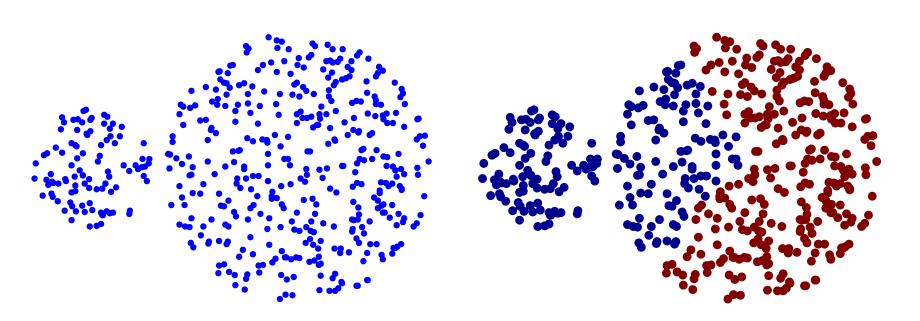
**Original Points** 

**Two Clusters** 

Less susceptible to noise and outliers

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### **Limitations of MAX**



**Original Points** 

**Two Clusters** 

- Tends to break large clusters
- Biased towards globular clusters

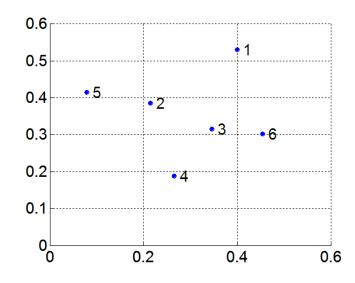
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### **Group Average**

 Proximity of two clusters is the average of pairwise proximity between points in the two clusters.

proximity( Cluster , Cluster , Cluster , Cluster , Cluster , Cluster , 
$$\mathbf{p}_{j} \in Cluster , \mathbf{p}_{j} \in Cluster ,$$

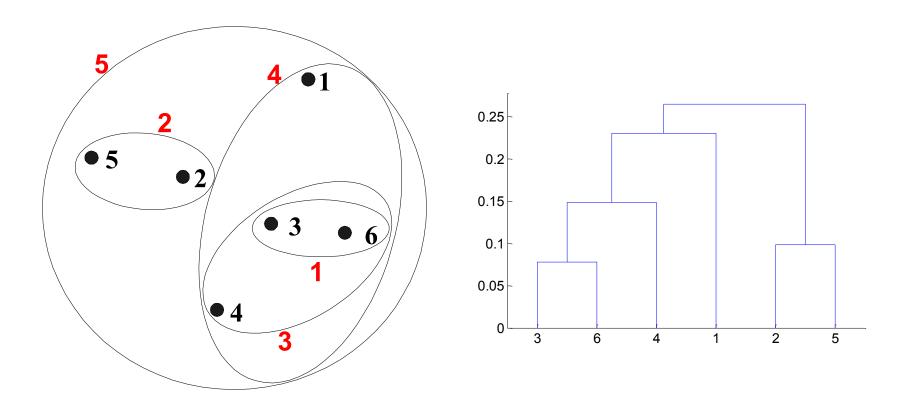
 Need to use average connectivity for scalability since total proximity favors large clusters



#### **Distance Matrix:**

	p1	p2	p3	p4	p5	p6
p1	0.00	0.24	0.22	0.37	0.34	0.23
p2	0.24	0.00	0.15	0.20	0.14	0.25
p3	0.22	0.15	0.00	0.15	0.28	0.11
p4	0.37	0.20	0.15	0.00	0.29	0.22
p5	0.34	0.14	0.28	0.29	0.00	0.39
p6	0.23	0.25	0.11	0.22	0.39	0.00

## **Hierarchical Clustering: Group Average**



**Nested Clusters** 

Dendrogram

# **Hierarchical Clustering: Group Average**

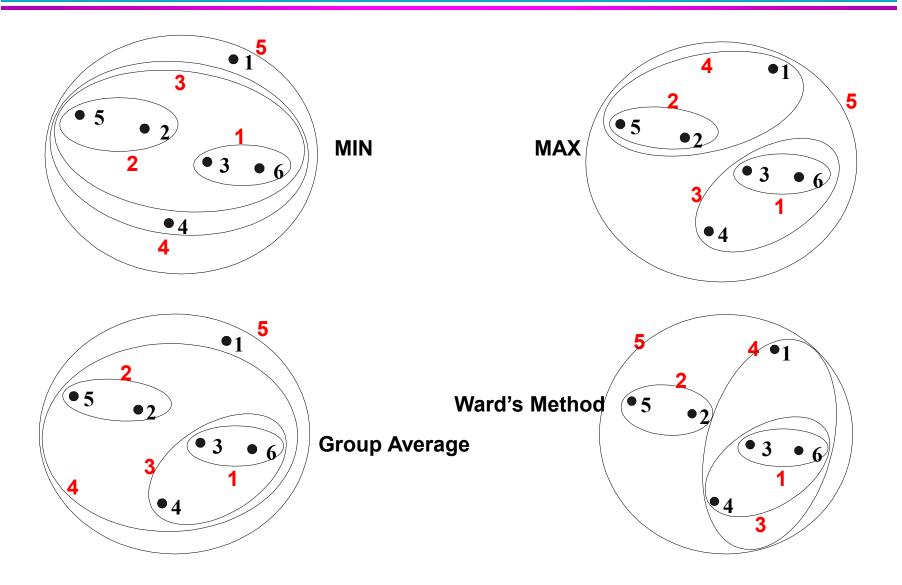
 Compromise between Single and Complete Link

- Strengths
  - Less susceptible to noise and outliers
- Limitations
  - Biased towards globular clusters

## **Cluster Similarity: Ward's Method**

- Similarity of two clusters is based on the increase in squared error when two clusters are merged
  - Similar to group average if distance between points is distance squared
- Less susceptible to noise and outliers
- Biased towards globular clusters
- Hierarchical analogue of K-means
  - Can be used to initialize K-means

#### **Hierarchical Clustering: Comparison**

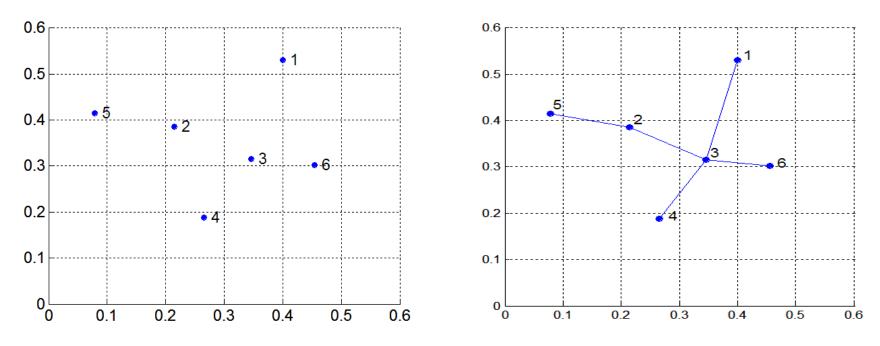


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## **MST: Divisive Hierarchical Clustering**

#### • Build MST (Minimum Spanning Tree)

- 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



02/14/2018

# **MST: Divisive Hierarchical Clustering**

#### Use MST for constructing hierarchy of clusters

#### Algorithm 7.5 MST Divisive Hierarchical Clustering Algorithm

- 1: Compute a minimum spanning tree for the proximity graph.
- 2: repeat
- 3: Create a new cluster by breaking the link corresponding to the largest distance (smallest similarity).
- 4: until Only singleton clusters remain

- O(N<sup>2</sup>) space since it uses the proximity matrix.
  - N is the number of points.
- O(N<sup>3</sup>) time in many cases
  - There are N steps and at each step the size, N<sup>2</sup>, proximity matrix must be updated and searched
  - Complexity can be reduced to O(N<sup>2</sup> log(N)) time with some cleverness

#### **Hierarchical Clustering: Problems and Limitations**

- Once a decision is made to combine two clusters, it cannot be undone
- No global objective function is directly minimized
- Different schemes have problems with one or more of the following:
  - Sensitivity to noise and outliers
  - Difficulty handling clusters of different sizes and nonglobular shapes
  - Breaking large clusters