

Master Degree in Artificial Intelligence for Science and Technology

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# Cluster Analysis: Hierarchical Clustering



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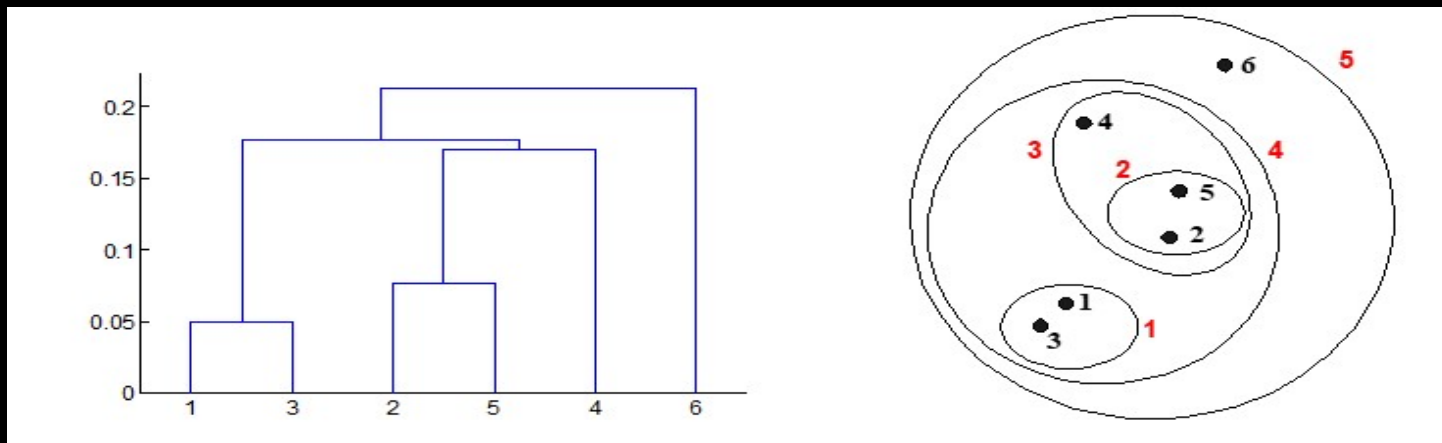
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## OUTLOOK

- Concept
- Strengths
- Types
  - Agglomerative
    - single linkage
    - complete linkage
    - average linkage
    - Ward's method
  - Divisive
- Complexity
- Limitations

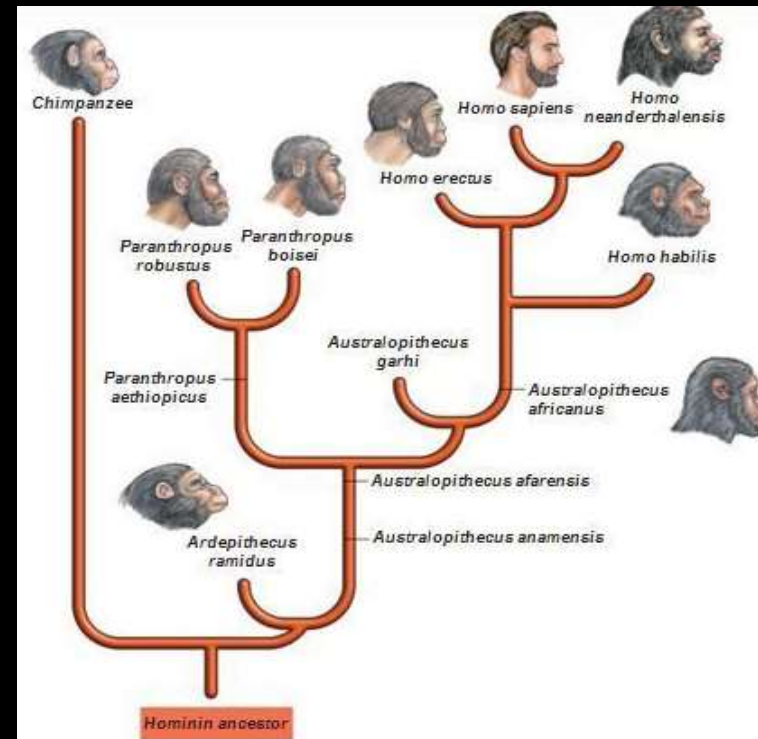
## CONCEPT

- 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

- 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, ...)



## TYPES OF 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

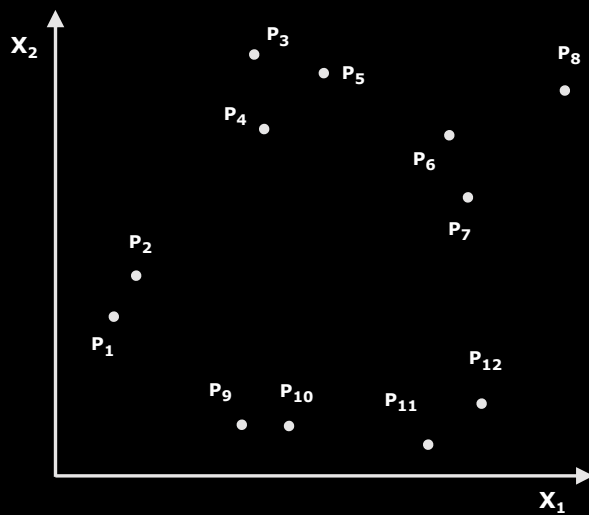
— **KEY IDEA:** successively merge closest clusters

### BASIC 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

- Key operation is the computation of the proximity of two clusters
- Different approaches to defining the distance between clusters distinguish the different algorithms

## AGGLOMERATIVE CLUSTERING



- **STEP 1:** compute the proximity matrix

	p1	p2	p3	p4	p5	p6	p7	p8	p9	p10	p11	p12
p1												
p2												
p3												
p4												
p5												
p6												
p7												
p8												
p9												
p10												
p11												
p12												

•••••••••••••

$p_1$

$p_2$

$p_3$

$p_4$

$p_5$

$p_6$

$p_7$

$p_8$

$p_9$

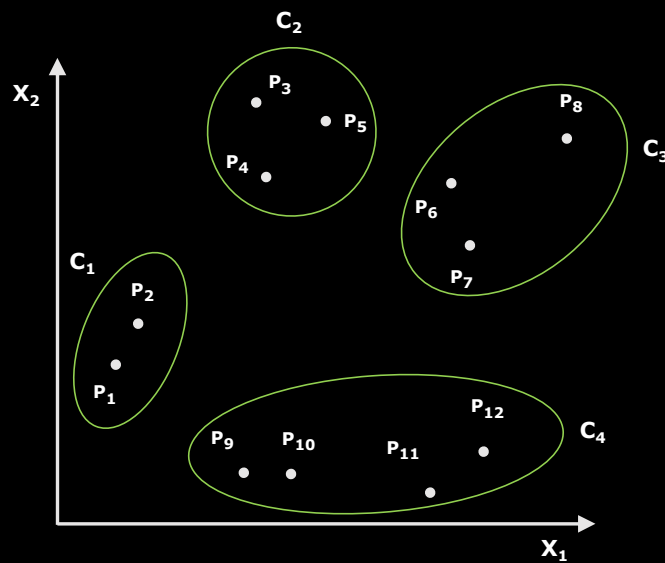
$p_{10}$

$p_{11}$

$p_{12}$

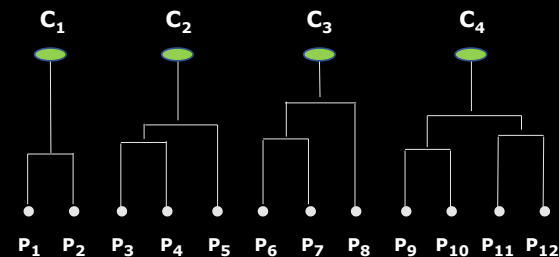
- **STEP 2:** each point is a cluster

## AGGLOMERATIVE CLUSTERING



	C1	C2	C3	C4
C1				
C2				
C3				
C4				

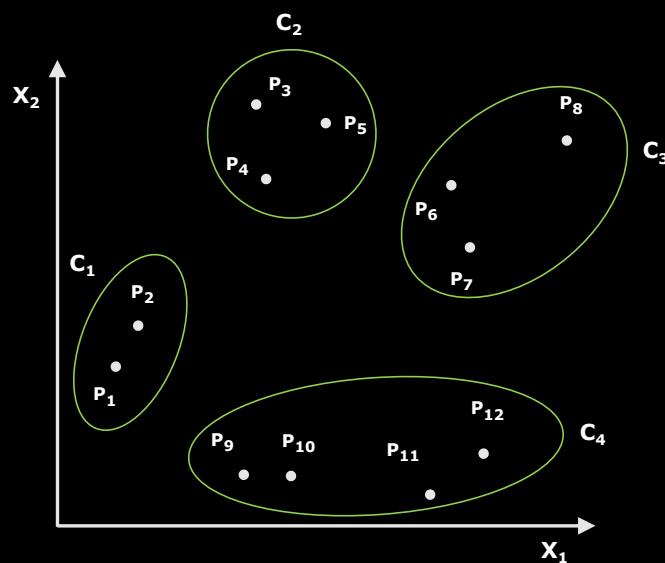
PROXIMITY MATRIX



After some merging steps we have some clusters

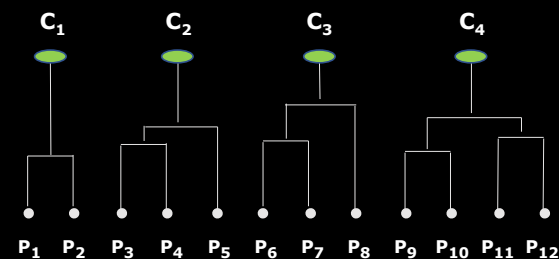


## AGGLOMERATIVE CLUSTERING



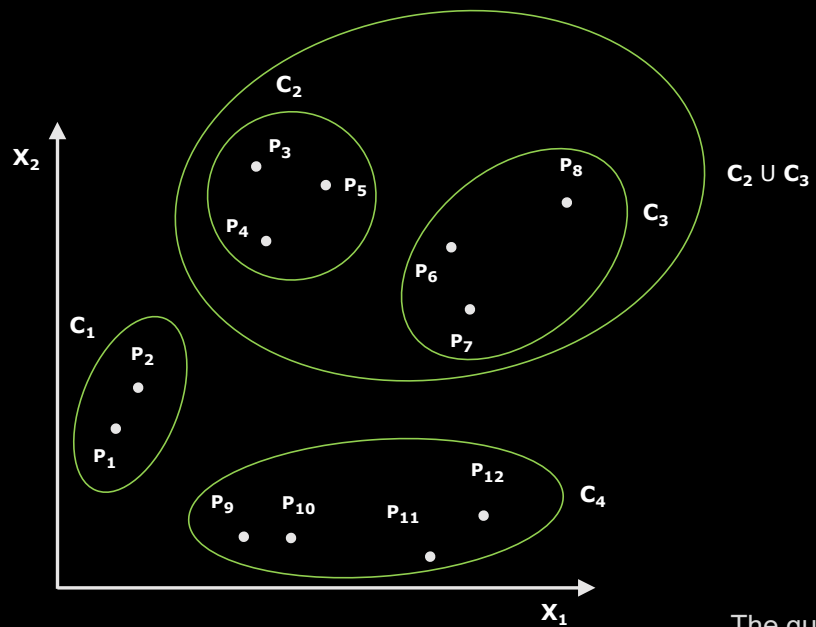
	C1	C2	C3	C4
C1				
C2				
C3				
C4				

PROXIMITY MATRIX



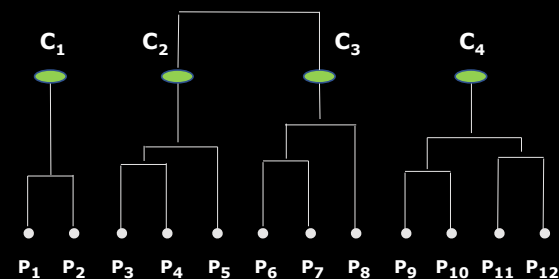
We want to merge the two closest clusters ( $C_2$  and  $C_3$ ) and update the proximity matrix.

## AGGLOMERATIVE CLUSTERING



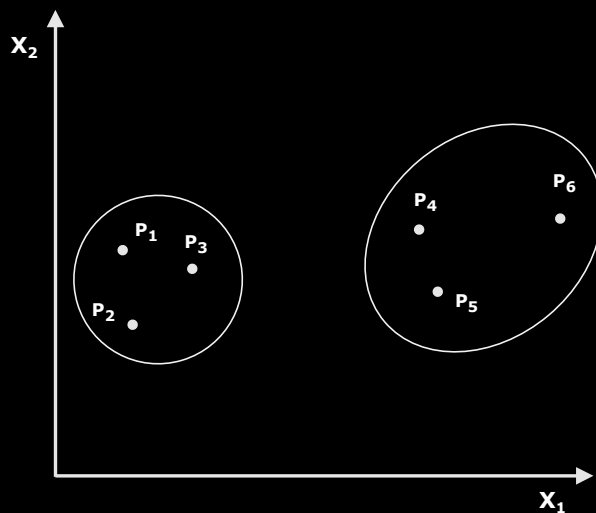
	C1	C2 C3	C4
C1		?	
C2 C3	?	?	?
C4		?	

PROXIMITY MATRIX



The question is “How do we update the proximity matrix?”

## HOW TO DEFINE INTER-CLUSTER PROXIMITY?

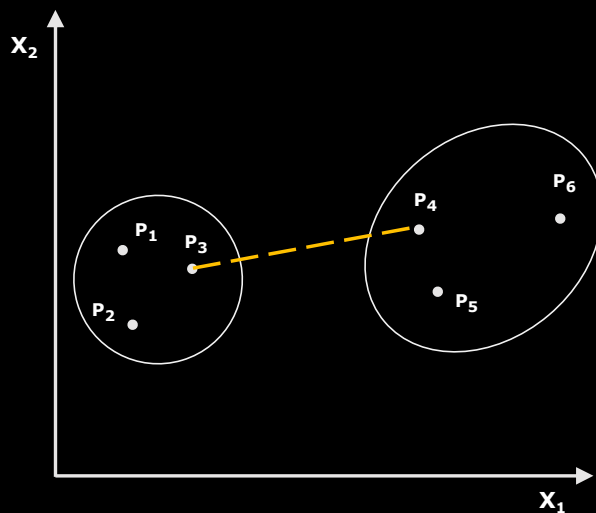


	C1	C2	C3	C4
C1				
C2				
C3				
C4				

PROXIMITY MATRIX

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

## HOW TO DEFINE INTER-CLUSTER PROXIMITY?

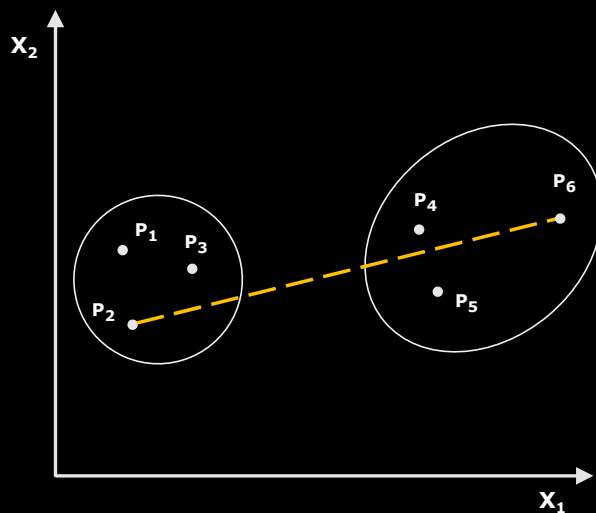


	C1	C2	C3	C4
C1				
C2				
C3				
C4				

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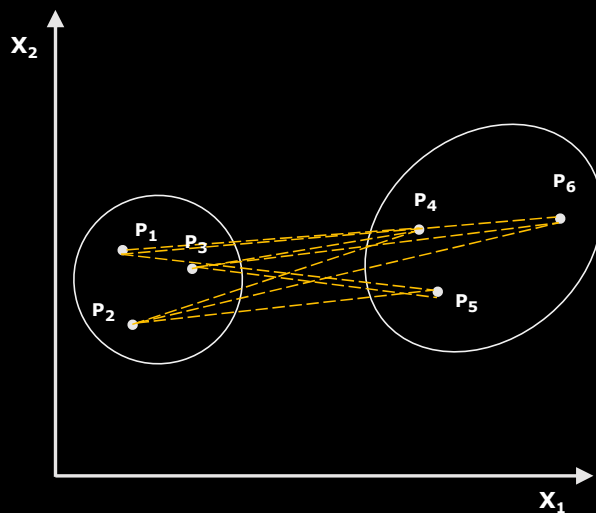


	C1	C2	C3	C4
C1				
C2				
C3				
C4				

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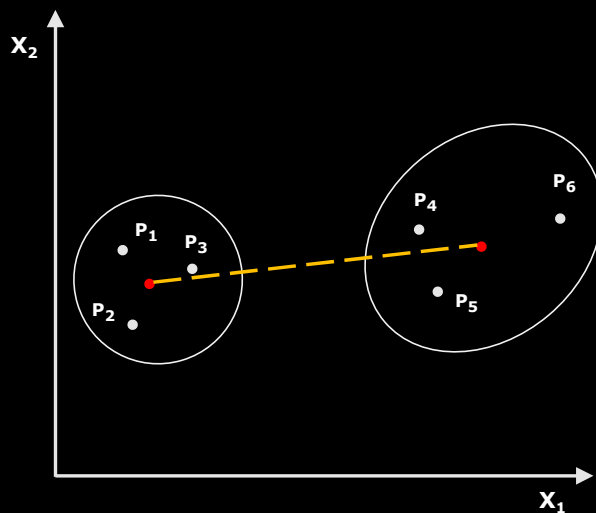


	C1	C2	C3	C4
C1				
C2				
C3				
C4				

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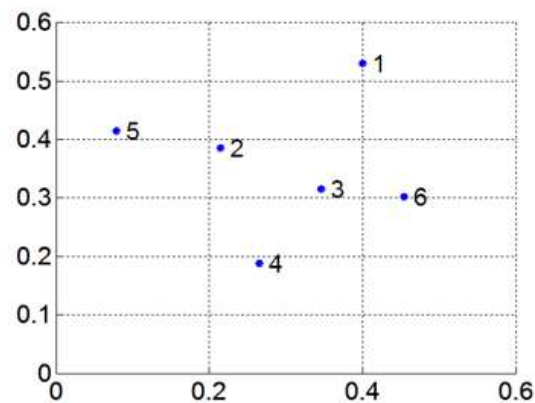
	C1	C2	C3	C4
C1				
C2				
C3				
C4				

PROXIMITY MATRIX

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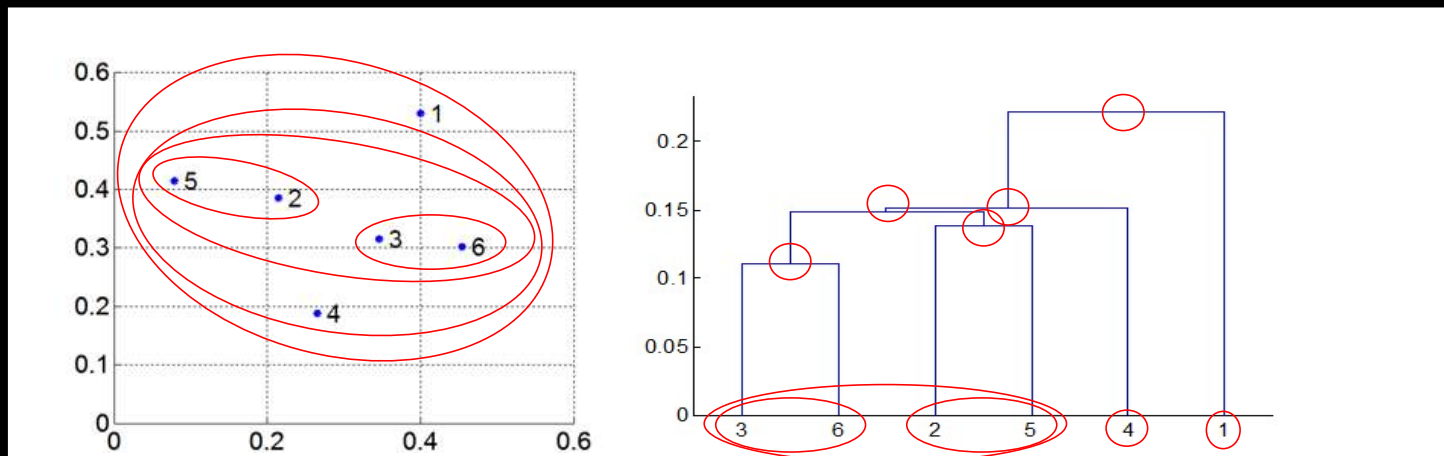
**MIN OR SINGLE LINKAGE**

- 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

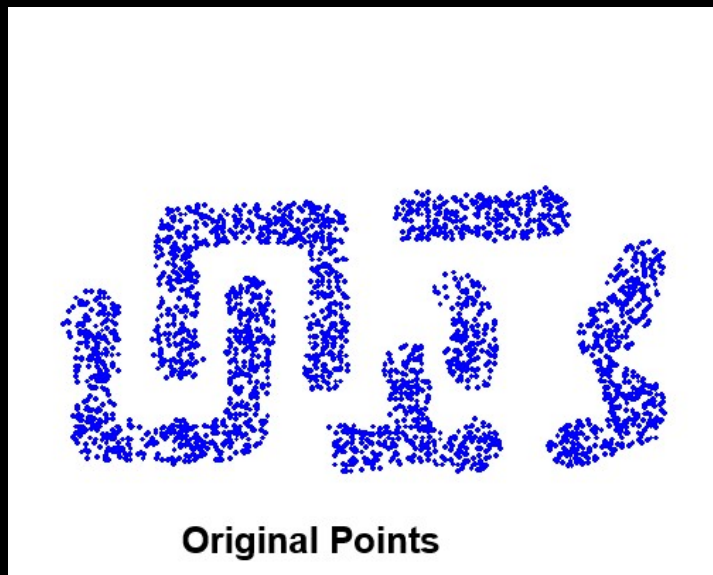
**Distance Matrix:**

	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

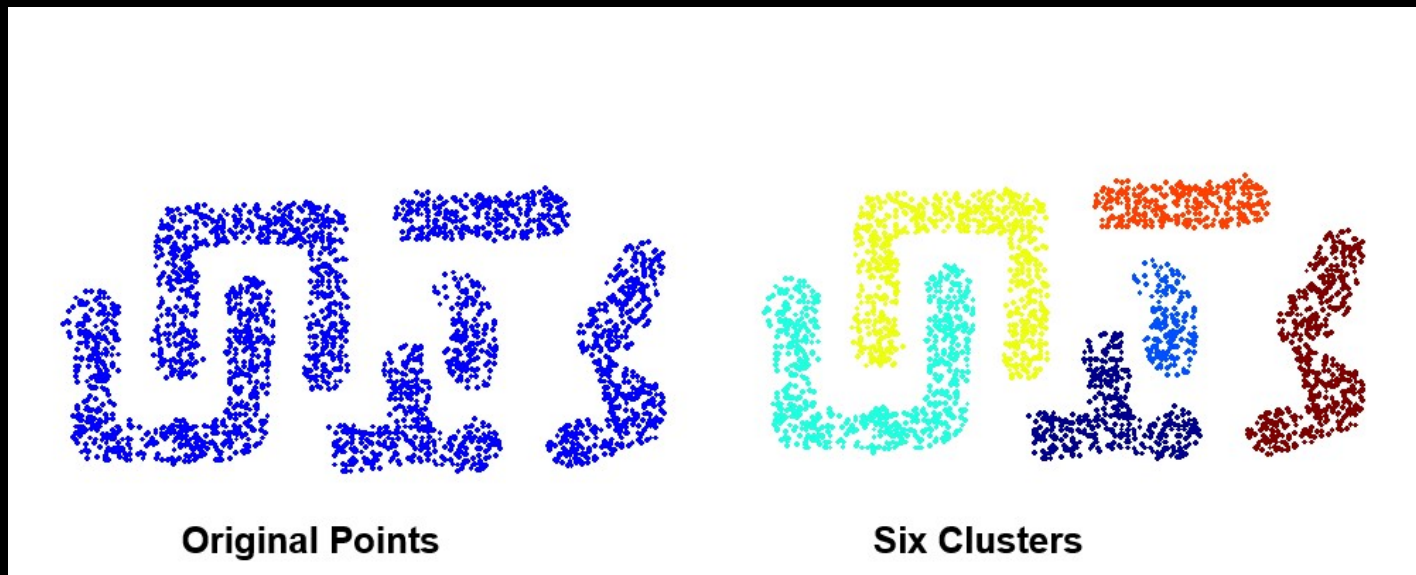


**MIN OR SINGLE LINKAGE**

## MIN OR SINGLE LINKAGE (STRENGTHS)

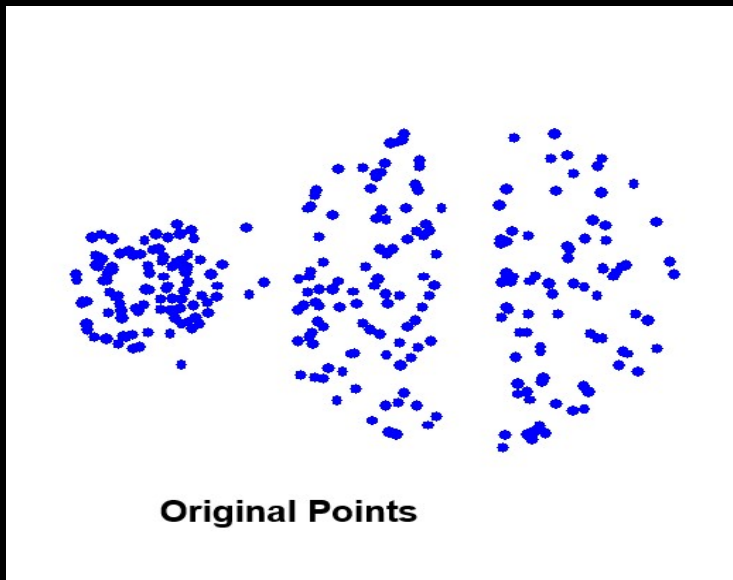


## MIN OR SINGLE LINKAGE (STRENGTHS)

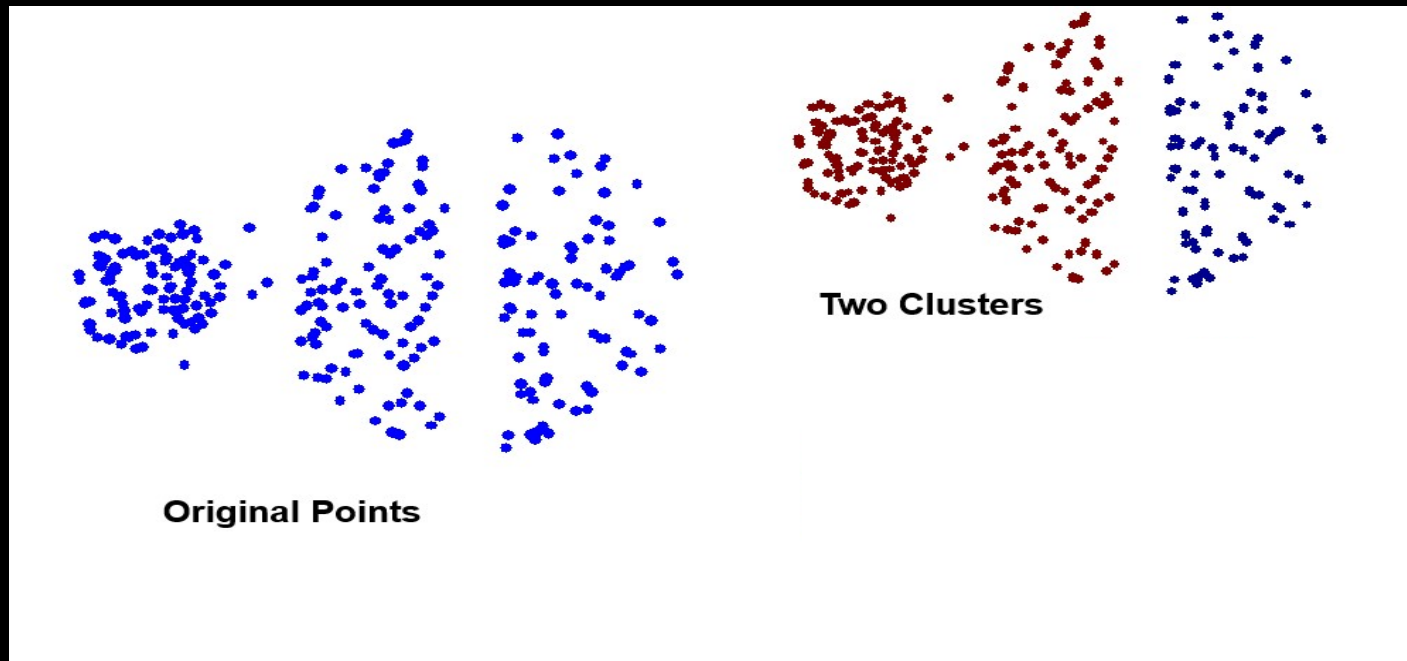


Can handle non-elliptical shapes

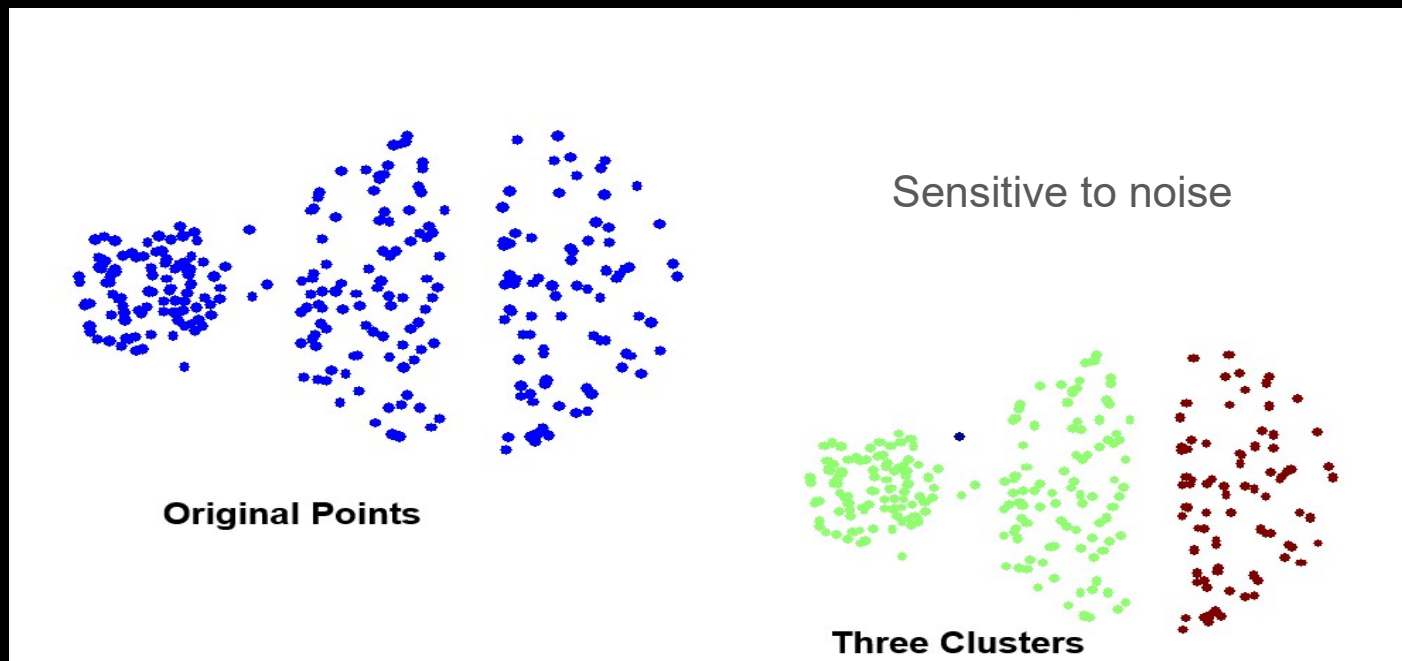
## MIN OR SINGLE LINKAGE (LIMITATIONS)



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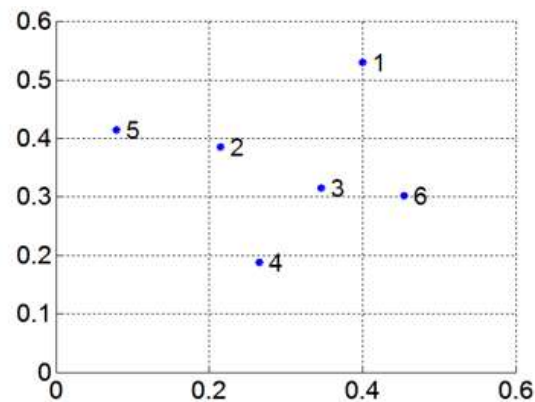


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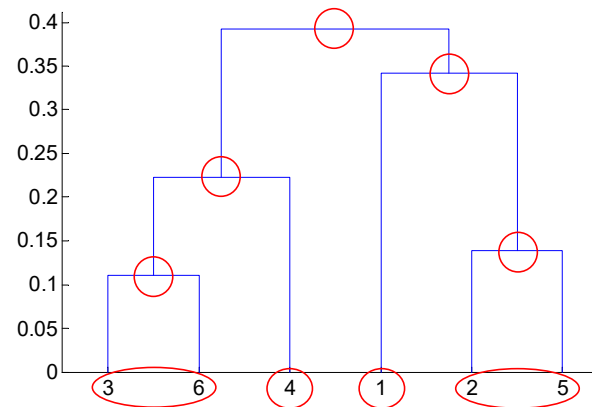
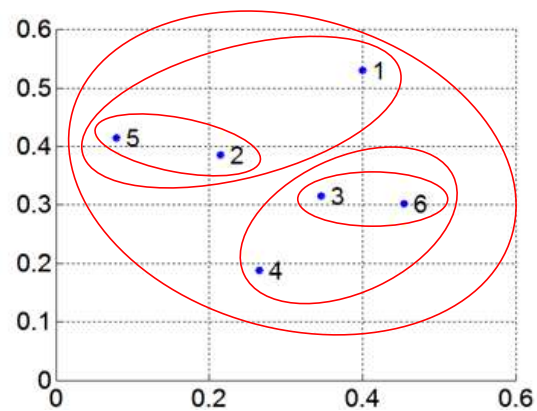


**MAX OR COMPLETE LINKAGE**

- 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

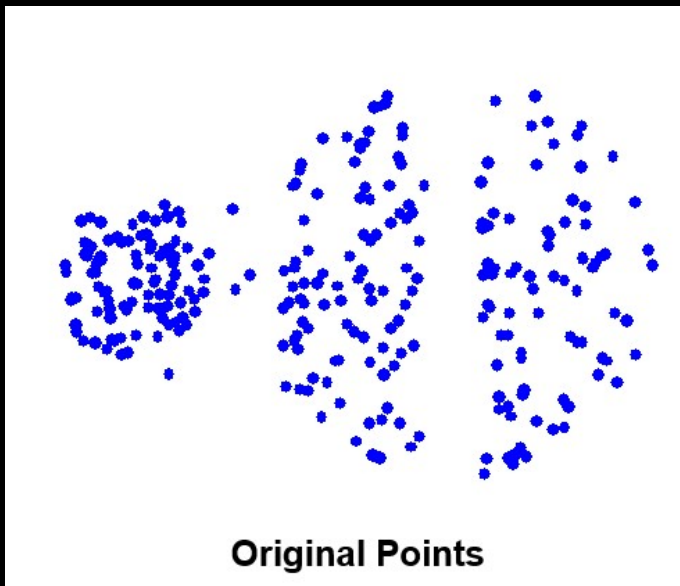
**Distance Matrix:**

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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

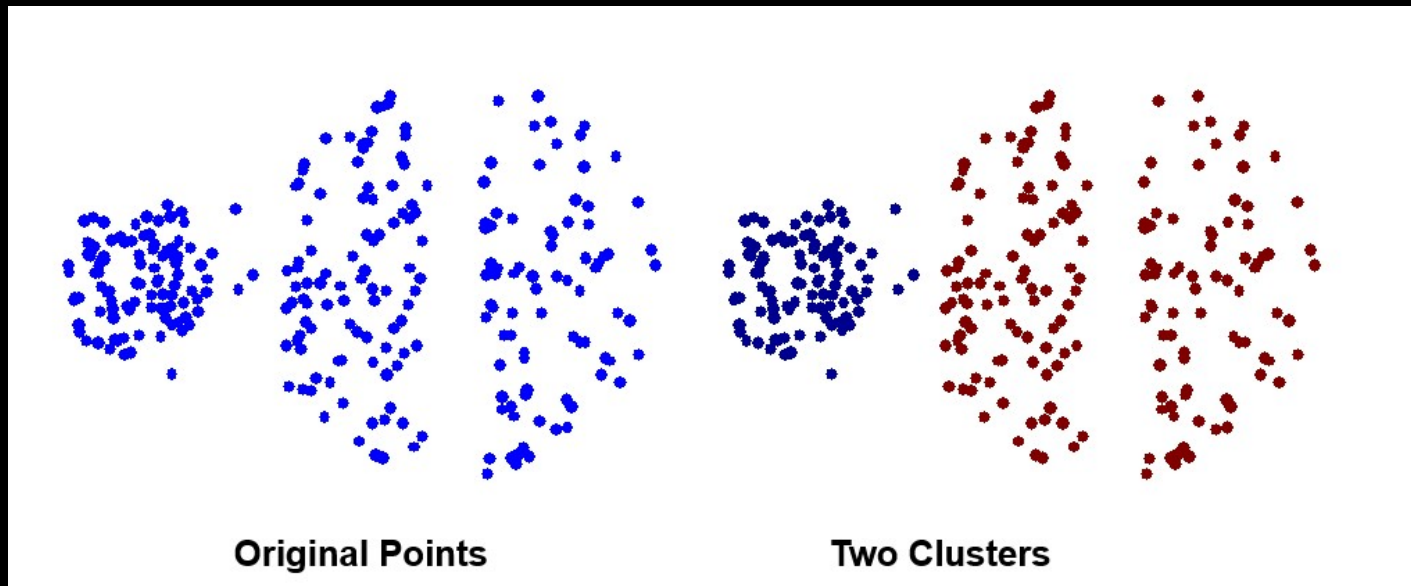
**MAX OR COMPLETE LINKAGE**



## MAX OR COMPLETE LINKAGE (STRENGTHS)

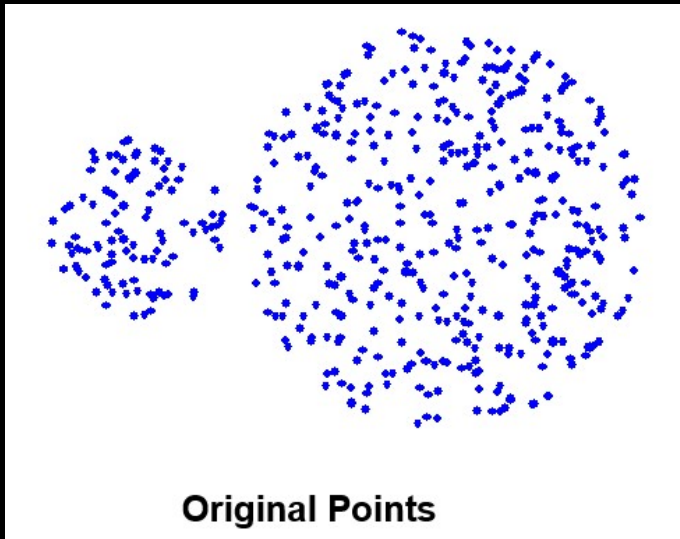


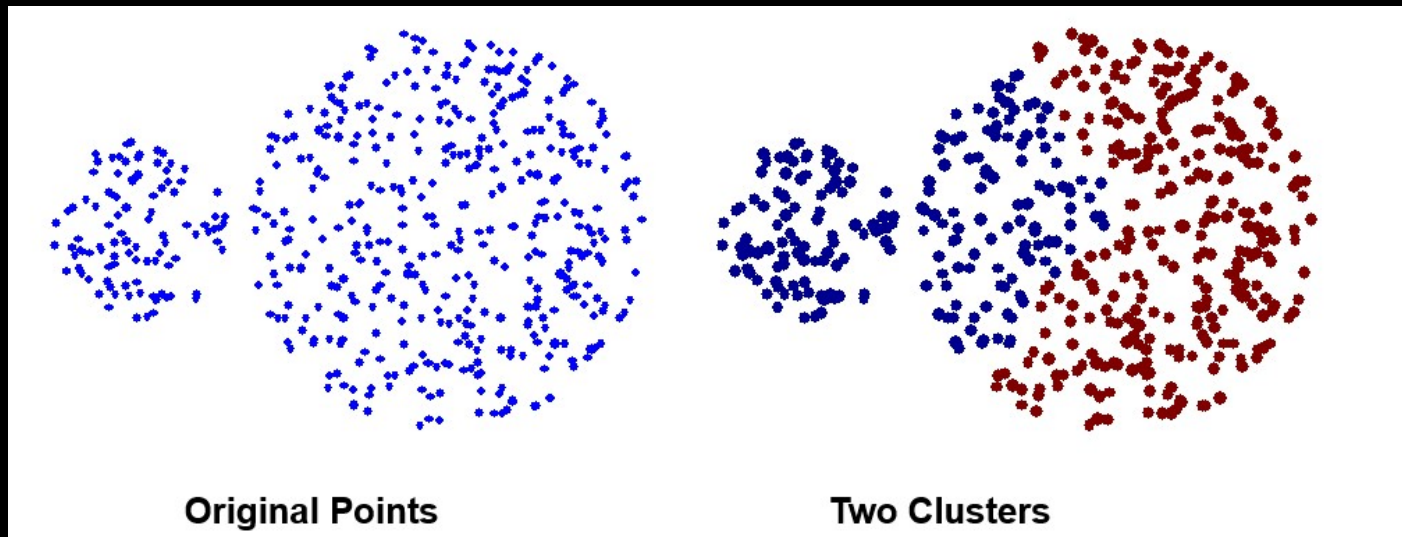
## MAX OR COMPLETE LINKAGE (STRENGTHS)



Less susceptible to noise

## MAX OR COMPLETE LINKAGE (LIMITATIONS)



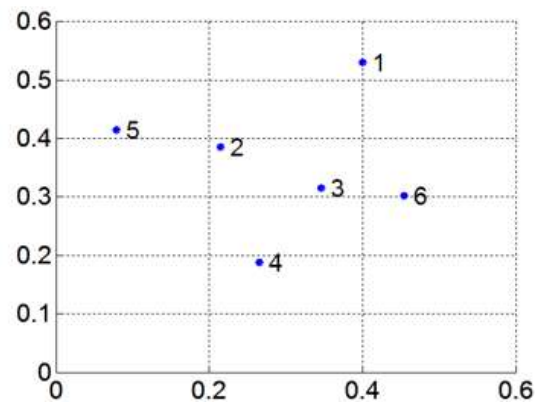
**MAX OR COMPLETE LINKAGE (LIMITATIONS)**

- Tends to break large clusters
- Biased towards globular clusters

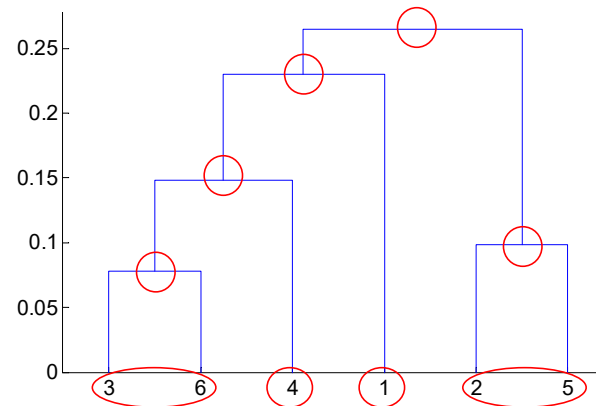
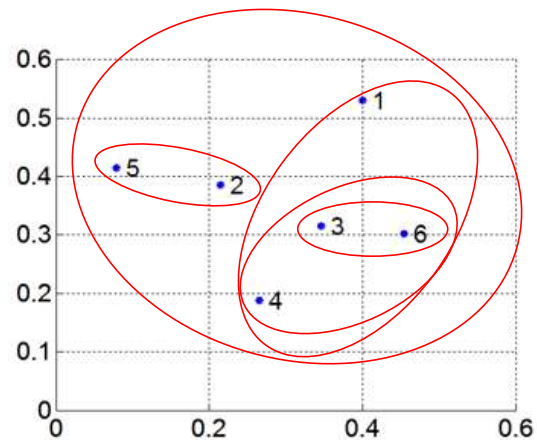
**GROUP AVERAGE**

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

$$\text{proximity}(C_i, C_j) = \frac{\sum_{p_k \in C_i, p_m \in C_j} \text{proximity}(p_k, p_m)}{|C_i||C_j|}$$

**Distance Matrix:**

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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

**GROUP AVERAGE**

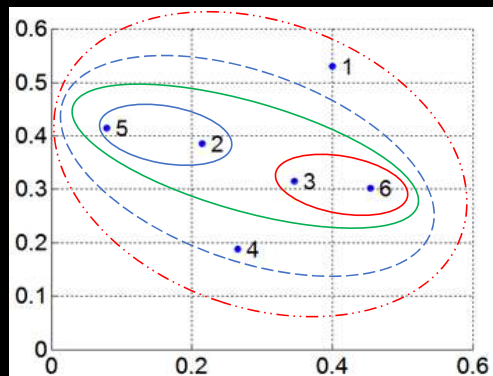
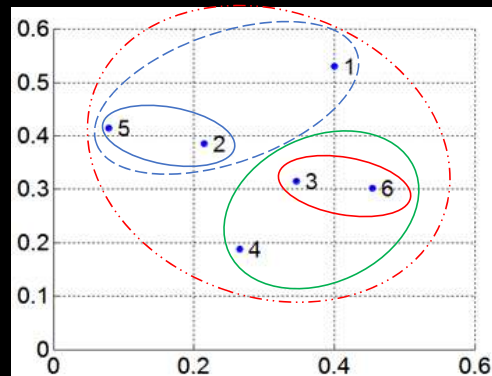
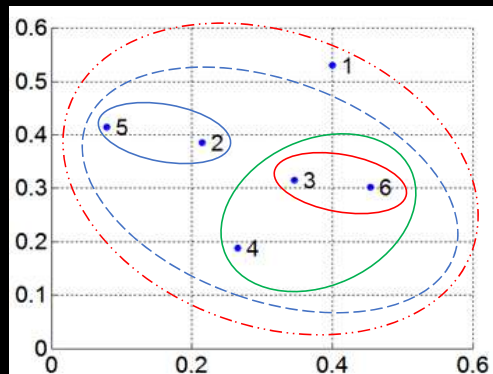
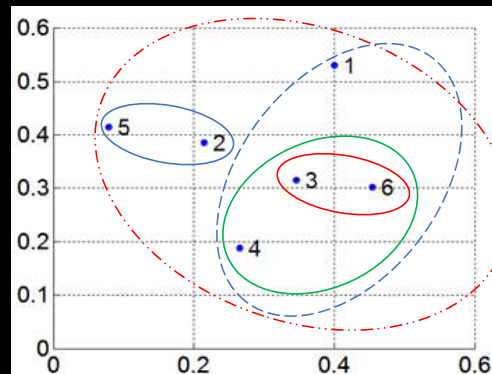
## GROUP AVERAGE

- Compromise between single and complete link
- Strengths
  - less susceptible to noise
- Limitations
  - biased towards globular clusters

## 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
- Biased towards globular clusters
- Hierarchical analogue of K-means
  - can be used to initialize K-means



**MIN****MAX****AVERAGE****WARD'S**

**HIERARCHICAL CLUSTERING: TIME AND SPACE REQUIREMENTS**

- $O(N^2)$  space since it uses the proximity matrix.
  - $N$  is the number of points.
  
- $O(N^3)$  time in many cases
  - there are  $N$  steps and at each step the size,  $N^2$  proximity matrix must be updated and searched
  - complexity can be reduced to  $O(N^2 \log(N))$  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
  - difficulty handling clusters of different sizes and non-globular shapes
  - breaking large clusters

## RECAP

- Concept
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