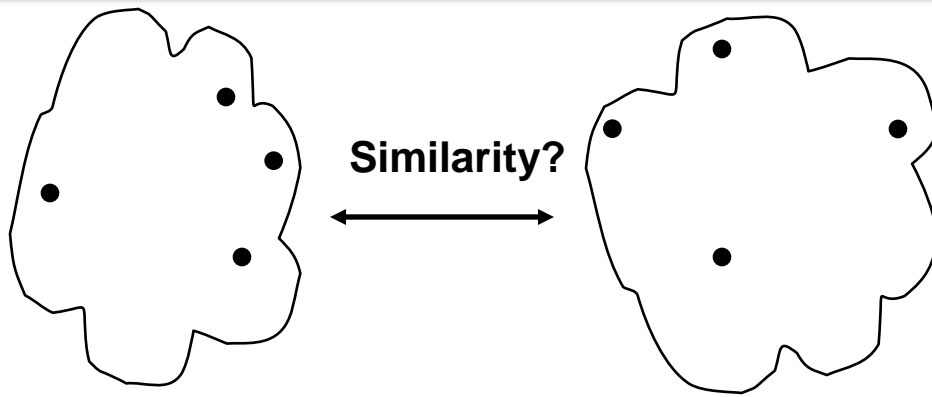


# Quick Review

# How to Define Inter-Cluster Similarity



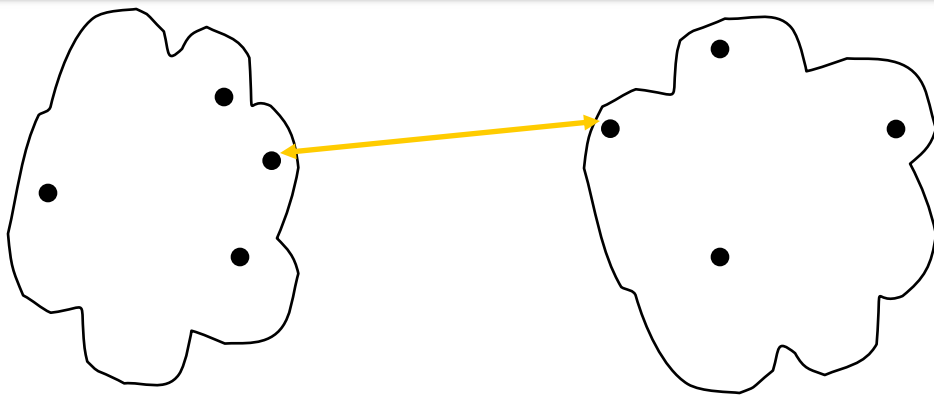
- MIN
- MAX
- Group Average

	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						
.						

.

· **Proximity Matrix**

# How to Define Inter-Cluster Similarity

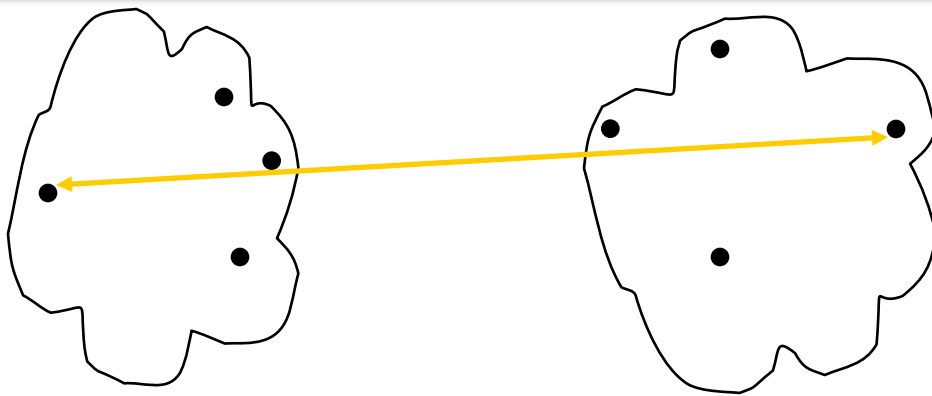


- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
  - Ward's Method uses squared error

	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						
.						

Proximity Matrix

# How to Define Inter-Cluster Similarity

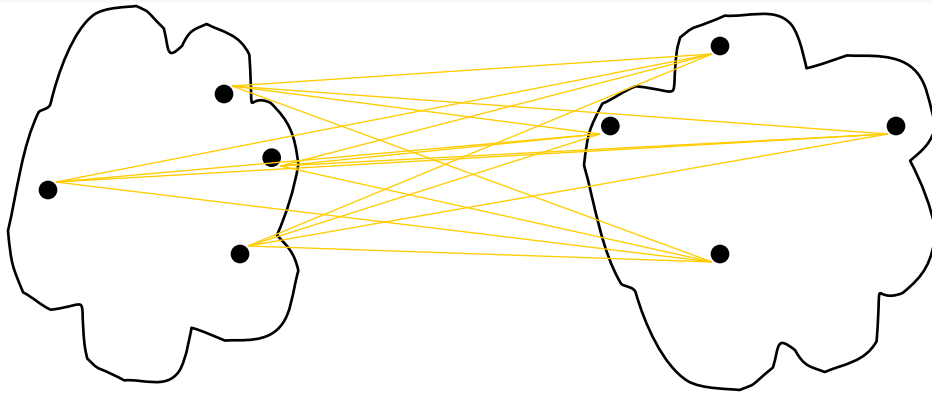


- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
  - Ward's Method uses squared error

	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						
.						
.						
.						

**Proximity Matrix**

# How to Define Inter-Cluster Similarity



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

	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						
.						
.						
.						

**Proximity Matrix**

# Single Link

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
<i>a</i>	0	12	6	3	25	4
<i>b</i>		0	19	8	14	15
<i>c</i>			0	12	5	18
<i>d</i>				0	11	9
<i>e</i>					0	7
<i>f</i>						0

	<i>ad</i>	<i>b</i>	<i>c</i>	<i>e</i>	<i>f</i>
<i>ad</i>	0	8	6	11	4
<i>b</i>		0	19	14	15
<i>c</i>			0	5	18
<i>e</i>				0	7
<i>f</i>					0

	<i>adf</i>	<i>b</i>	<i>ce</i>
<i>adf</i>	0	8	6
<i>b</i>		0	14
<i>ce</i>			0

	<i>adfce</i>	<i>b</i>
<i>adfce</i>	0	8
<i>b</i>	8	0

# Complete Link min(max distances)

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
<i>a</i>	0	12	6	3	25	4
<i>b</i>		0	19	8	14	15
<i>c</i>			0	12	5	18
<i>d</i>				0	11	9
<i>e</i>					0	7
<i>f</i>						0

	<i>ad</i>	<i>b</i>	<i>c</i>	<i>e</i>	<i>f</i>
<i>ad</i>	0	12	12	25	9
<i>b</i>		0	19	14	15
<i>c</i>			0	5	18
<i>e</i>				0	7
<i>f</i>					0

	<i>ad</i>	<i>b</i>	<i>ce</i>	<i>f</i>
<i>ad</i>	0	12	25	9
<i>b</i>		0	19	15
<i>ce</i>			0	18
<i>f</i>				0

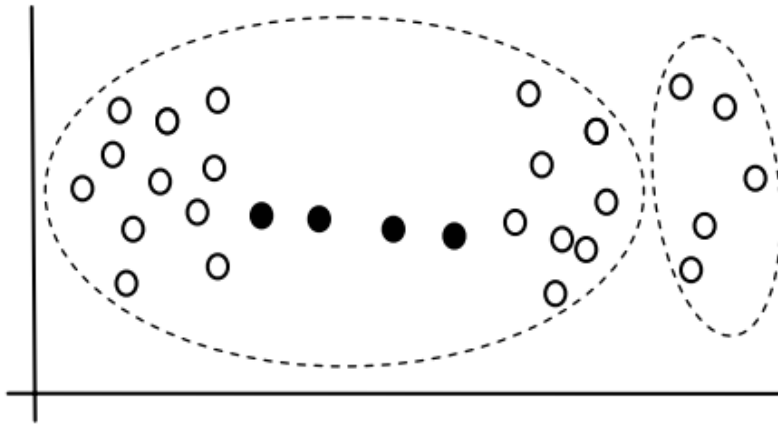
# Properties of intergroup similarity

- Single linkage
  - can produce “chaining,” where a sequence of close observations in different groups cause early merges of those groups
  
- Complete linkage has the opposite problem.
  - It might not merge close groups because of outlier members that are far apart.
  
- Group average represents a natural compromise,
  - but depends on the scale of the similarities. Applying a monotone transformation to the similarities can change the results.

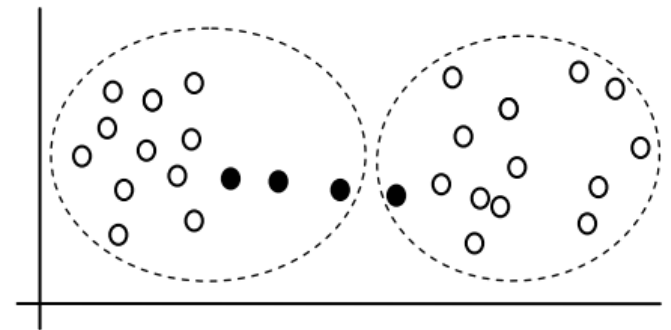


# Properties of intergroup similarity

- Single-link is suitable for non-elliptical shape clusters
- But, it is sensitive to noise and may cause a chain effect and produce straggly clusters



Chain effect of the single link method



Clustering using the complete link method

# Hierarchical Clustering: Time and Space

## □ SPACE

- $O(N^2)$  space since it uses the proximity matrix.
  - ◆  $N$  is the number of points.

## □ TIME

- $O(N^3)$  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 if we use a special structure like a heap or sorted lists

# Hierarchical Clustering: Limitations

- Once a decision is made to combine two clusters, it cannot be undone
- Do not scale well: time complexity of at least  $O(N^2 \log N)$ , where  $n$  is the number of total objects
- Different schemes have problems with one or more of the following:
  - Sensitivity to noise and outliers
  - Difficulty handling different sized clusters and convex shapes
  - Breaking large clusters

# 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

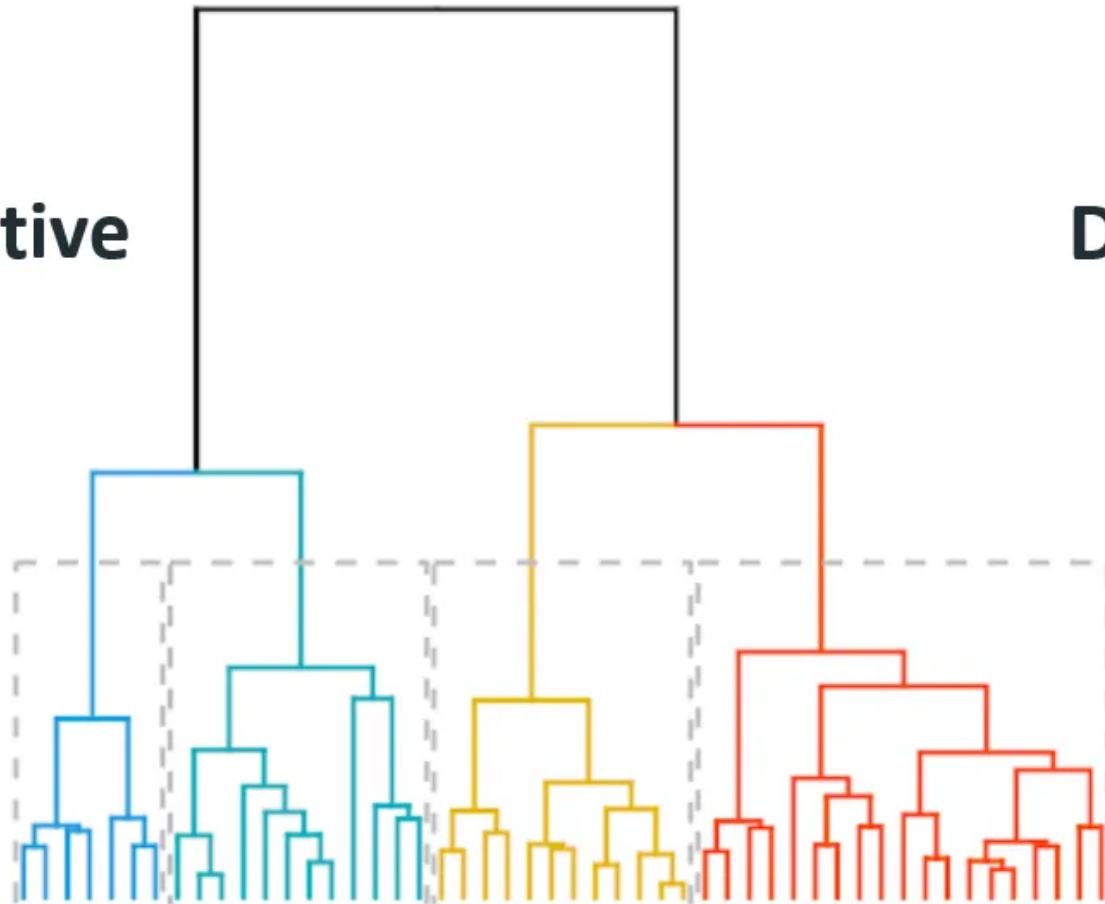
- Start with one, all-inclusive cluster
- At each step, split a cluster until each cluster contains a point (or there are  $k$  clusters)

**Traditional hierarchical algorithms use a similarity or distance matrix to Merge or split one cluster at a time**

# Hierarchical Clustering

**Agglomerative**

**Divisive**



# Agglomerative Clustering Algorithm

## More popular hierarchical clustering technique

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

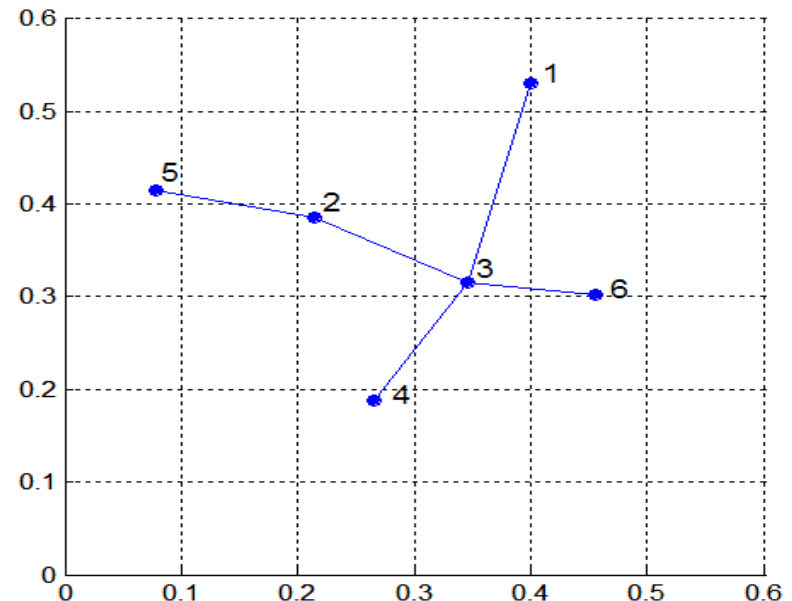
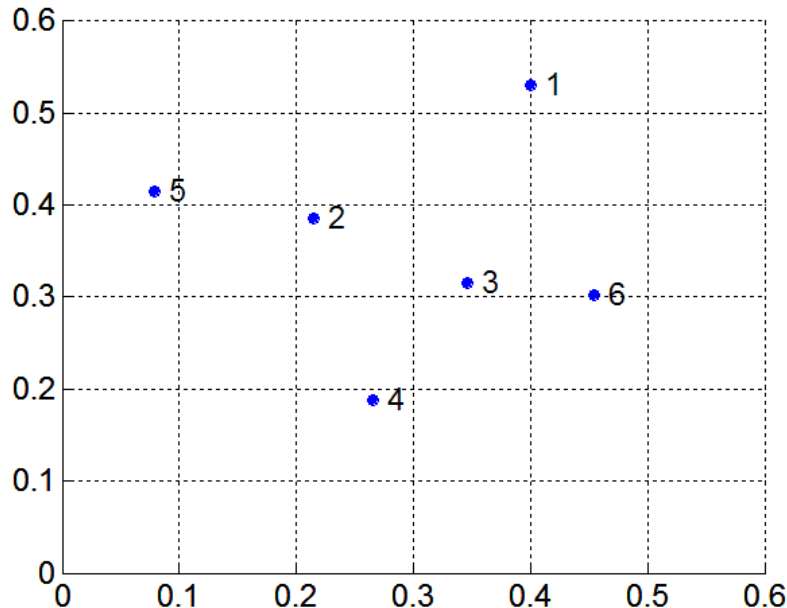
**Key operation is the computation of the proximity of two clusters**

Different approaches exist to define the distance between clusters

# MST: Divisive Hierarchical Clustering

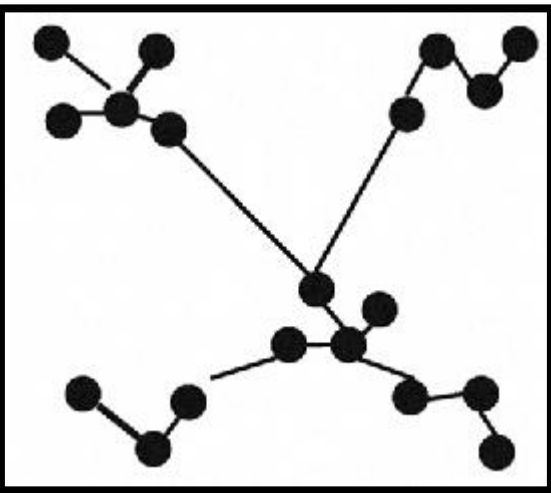
## Build MST (Minimum Spanning Tree)

- Start with an arbitrary vertex (consider it a tree with one vertex)
- 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  $q$

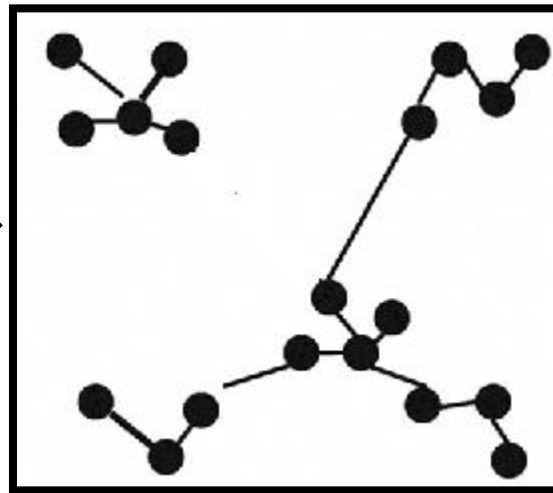


# Algorithm MST Divisive Hierarchical Clustering

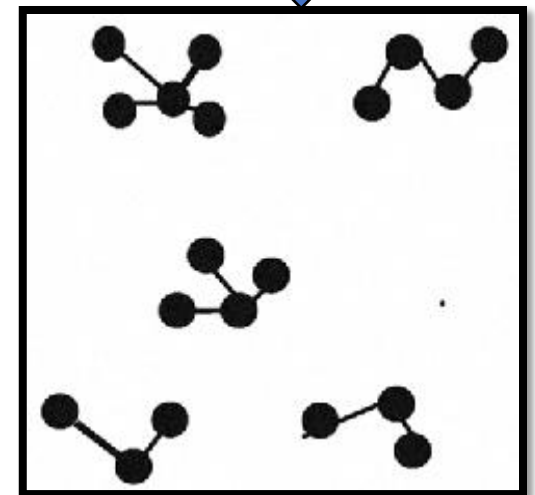
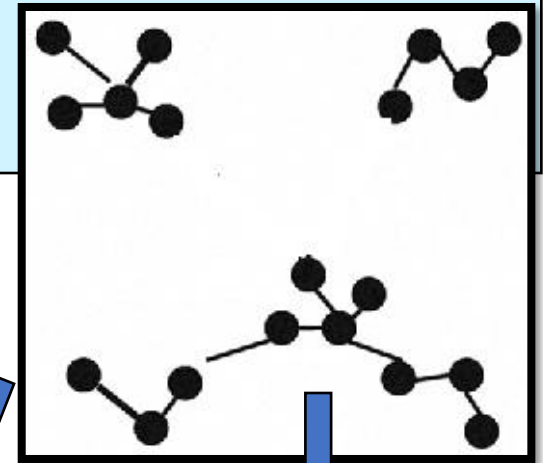
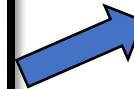
- Compute MST for the proximity graph
- **Repeat**
  - ◆ Create a new cluster by breaking the link corresponding to the largest distance (smallest similarity)
- **Until only singleton clusters remain**



**MST**



**Clusters after removing  
longest edges**

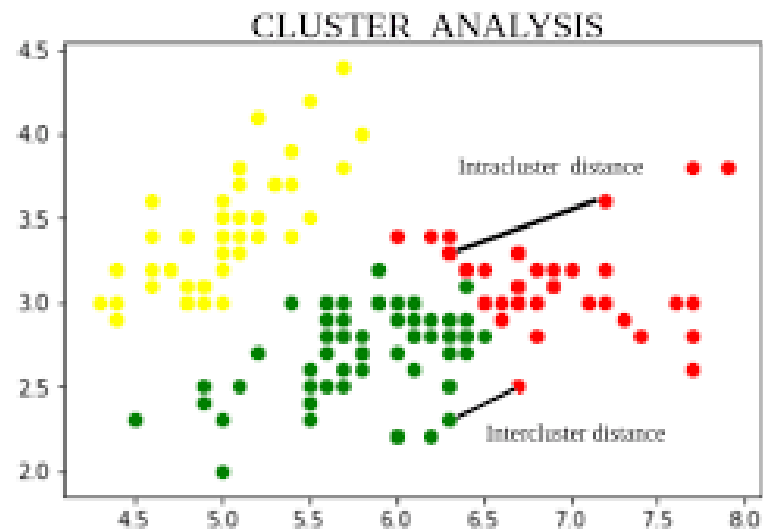




# Remarks

	Partitioning Clustering	Hierarchical Clustering
Time Complexity	$O(n)$	$O(n^2 \log n)$
Pros	Easy to use and Relatively efficient	Outputs a dendrogram that is desired in many applications.
Cons	Sensitive to initialization; bad initialization might lead to bad results. Need to store all data in memory.	higher time complexity; Need to store all data in memory.

# Cluster Validity

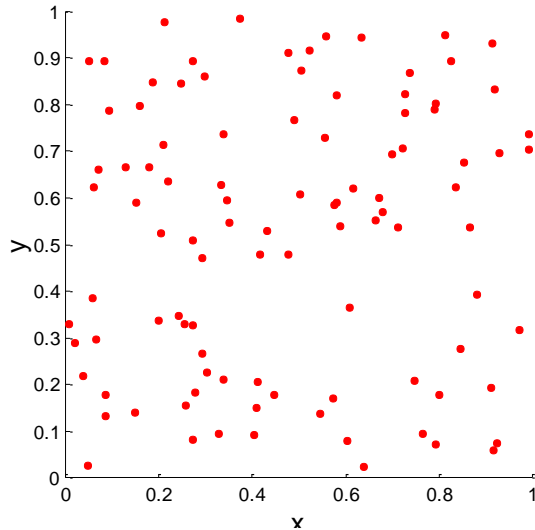


# Cluster Validity

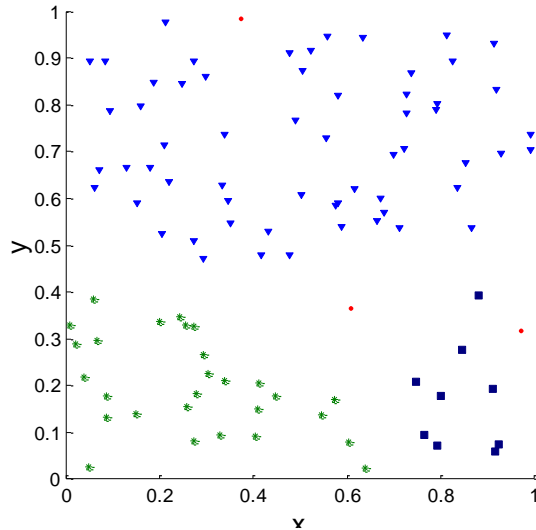
- For cluster analysis, we want to evaluate the “goodness” of the resulting clusters?
- But “clusters are in the eye of the beholder”!
- **Then why do we want to evaluate them?**
  - To avoid finding patterns in noise
  - To compare clustering algorithms
  - To compare two sets of clusters
  - To compare two clusters

# Clusters found in Random Data

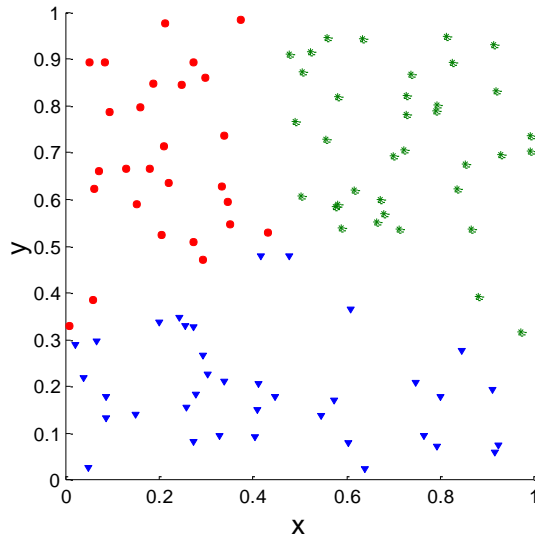
**Random  
Points**



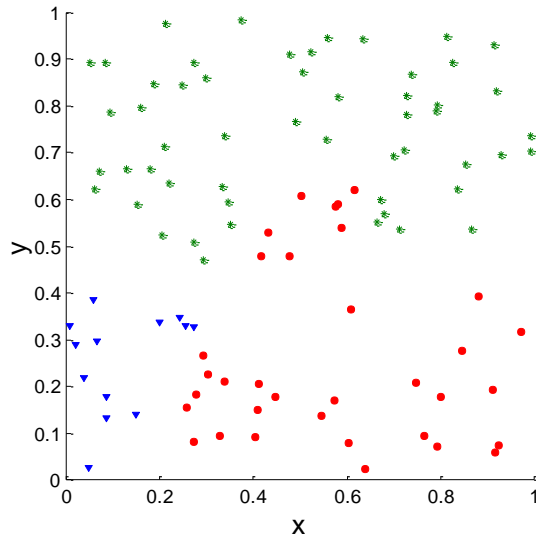
**DBSCAN**



**K-means**



**Complete  
Link**



# Measures of Cluster Validity

Numerical measures to judge cluster validity:

- **External Index:** measure the extent to which cluster labels match externally supplied class labels.
  - Entropy
- **Internal Index:** measure the goodness of a clustering structure *without* respect to external information.
  - Sum of Squared Error (SSE)
- **Relative Index:** Used to compare two different clusterings or clusters.
  - Often, an external or internal index is used for this function, e.g., SSE or entropy

# Unsupervised Cluster Evaluation

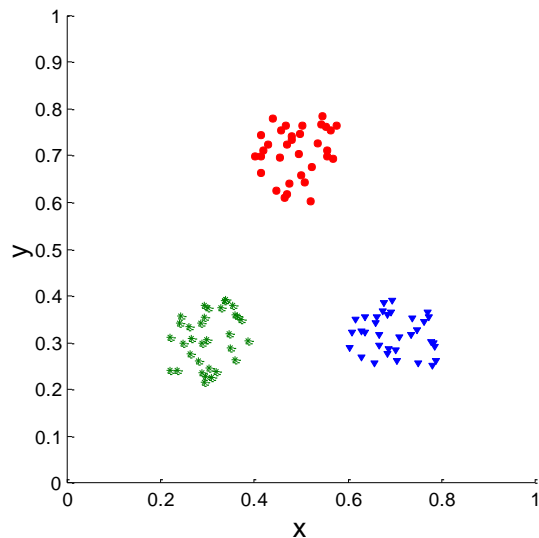
- Consider two unsupervised approaches for cluster evaluation using the Proximity Matrix
  - **Correlation of actual and Ideal Proximity matrices**
  - **Visualization**
- Two matrices
  - **Similarity Matrix for the data set**
  - **Ideal Similarity Matrix** (cluster label from cluster analysis)
    - ◆ One row and one column for each data point
    - ◆ An entry is 1 if the associated pair of points belong to the same cluster
    - ◆ An entry is 0 if the associated pair of points belongs to different clusters

# Measuring Cluster Validity Via Correlation

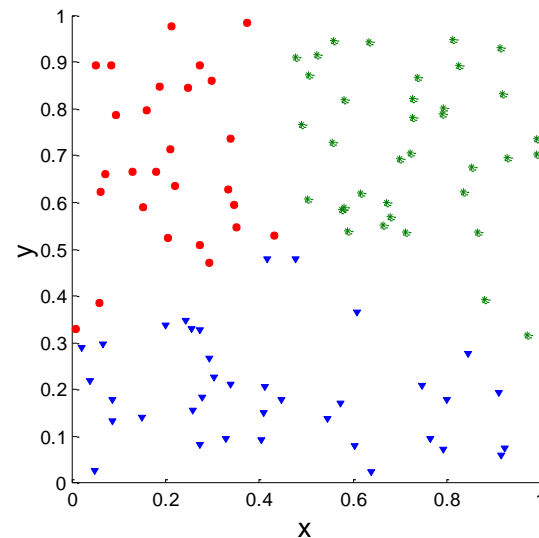
- Compute the correlation between the two matrices
  - Since the matrices are symmetric, only the correlation between  $n(n-1) / 2$  entries needs to be calculated.
- High correlation indicates that points that belong to the same cluster are close to each other.
- Not a good measure for some density or contiguity based clusters.

# Measuring Cluster Validity Via Correlation

- Correlation of Ideal and proximity matrices for the K-means clusterings of the following two data sets.



**Corr = 0.9235**

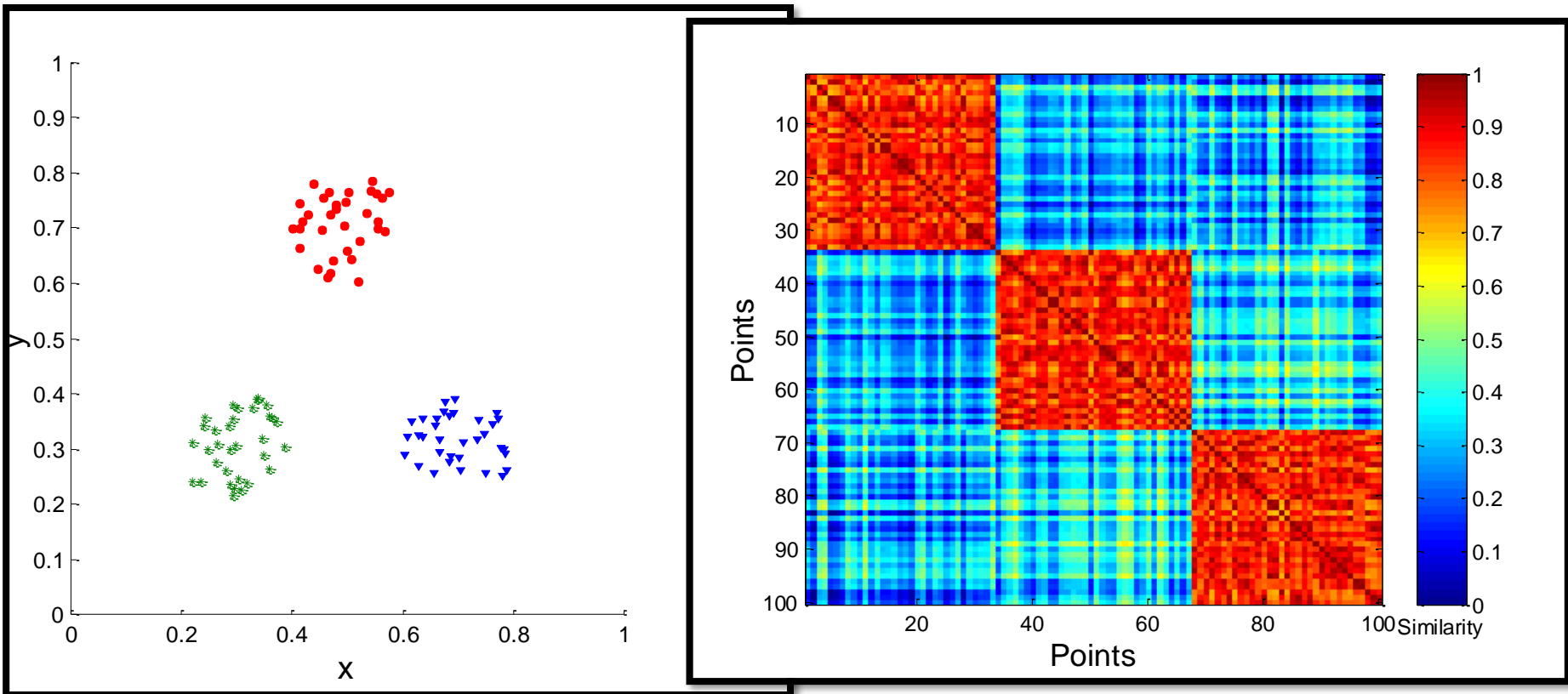


**Corr = 0.5810**



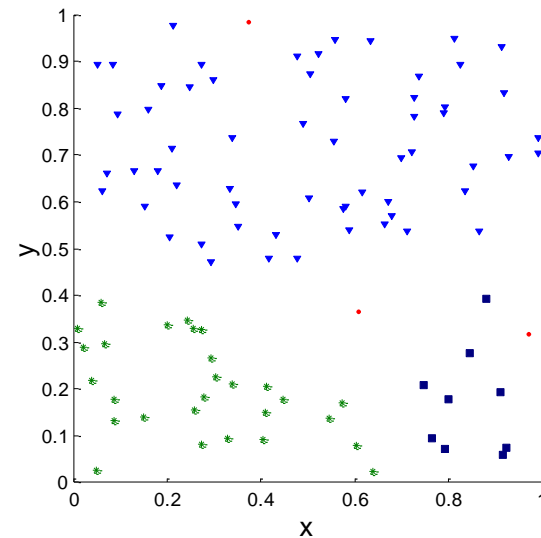
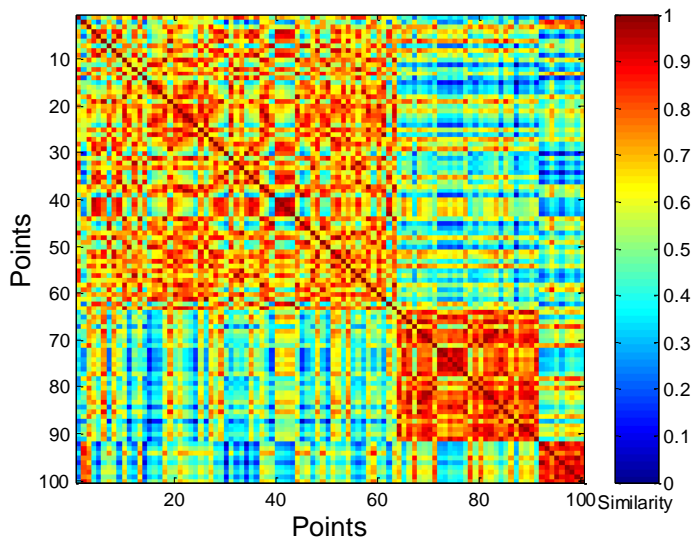
# Using Similarity Matrix for Cluster Validation

- Order the similarity matrix with respect to cluster labels and inspect visually.



# Using Similarity Matrix for Cluster Validation

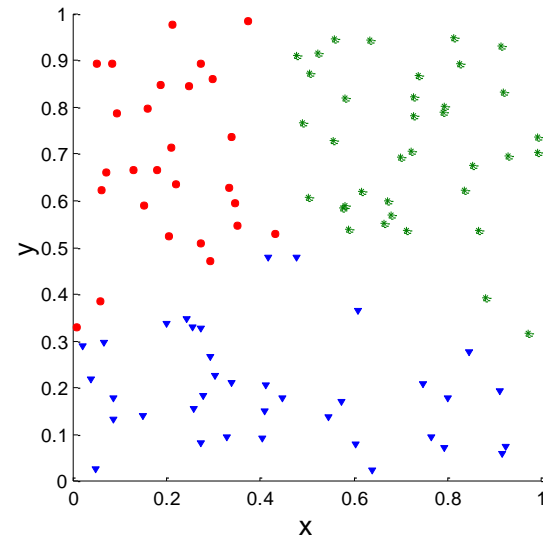
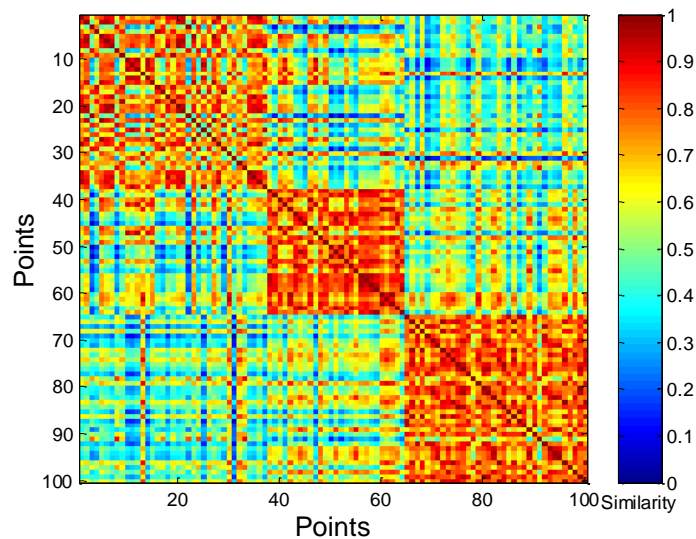
- Clusters in random data are not so crisp



**DBSCAN**

# Using Similarity Matrix for Cluster Validation

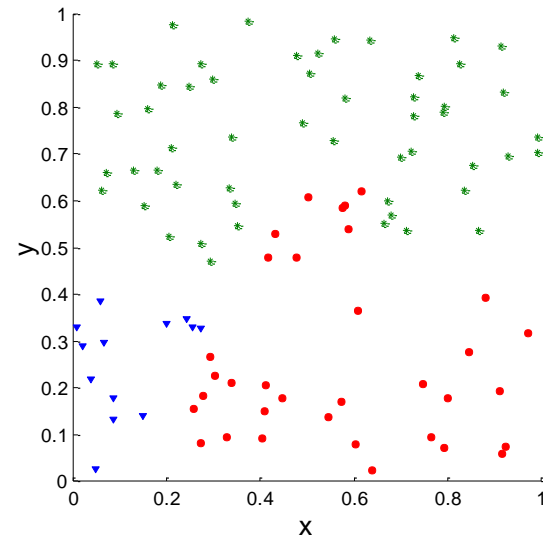
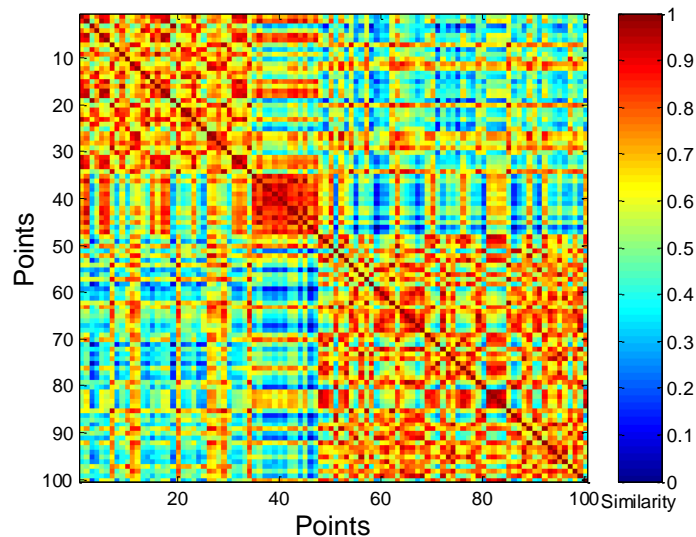
- Clusters in random data are not so crisp



**K-means**

# Using Similarity Matrix for Cluster Validation

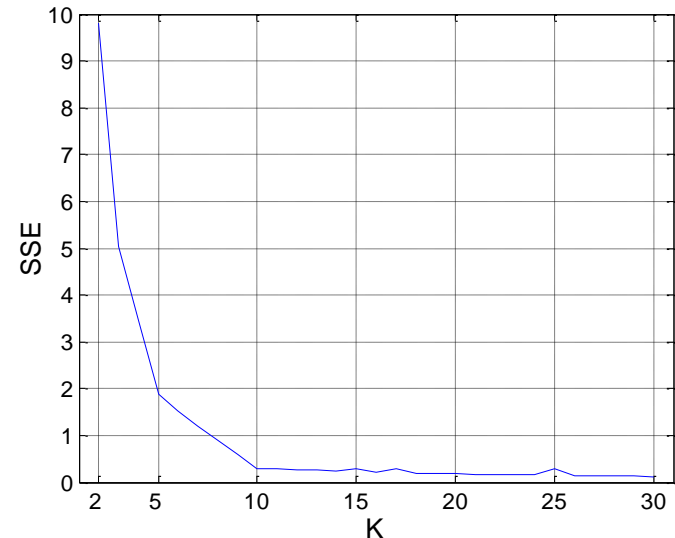
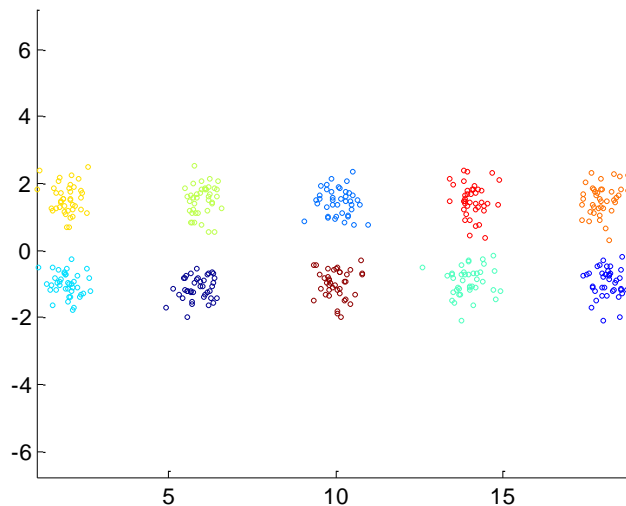
- Clusters in random data are not so crisp



**Complete Link**

# Internal Measures: SSE

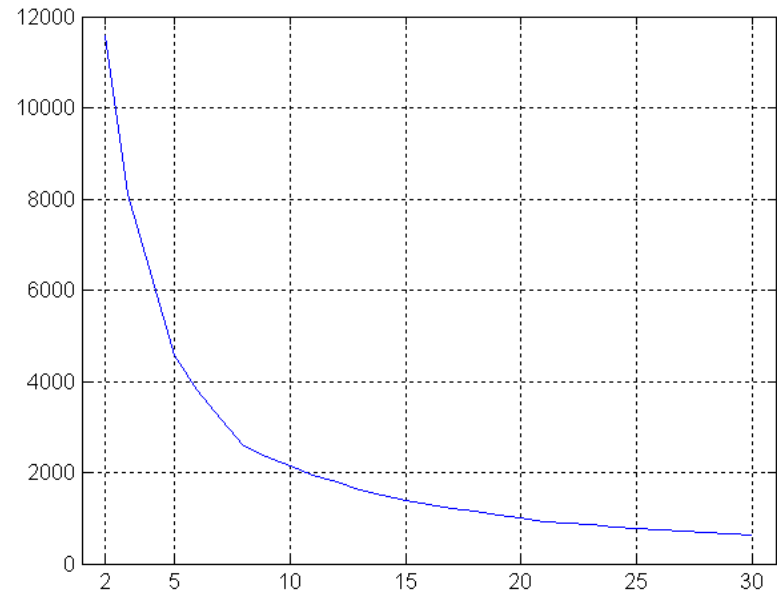
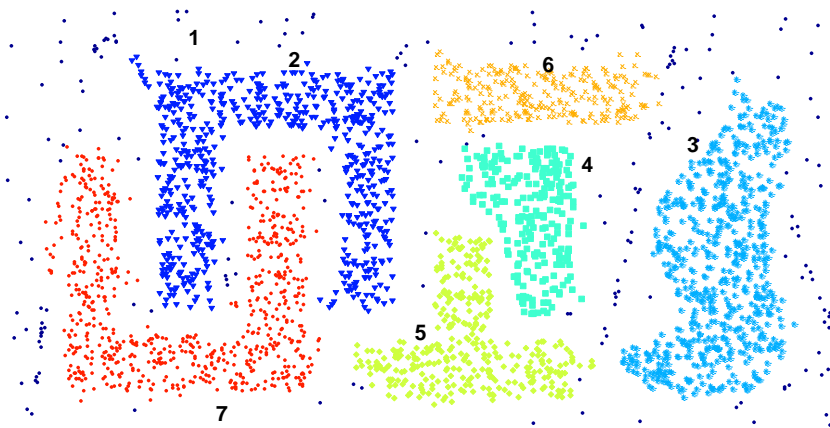
- ❑ Clusters in more complicated data aren't well separated
- ❑ **Internal Index:** Used to measure the goodness of a clustering structure without respect to external information
  - SSE
- ❑ SSE is good for comparing two clusterings or two clusters (average SSE).
- ❑ ***Can also be used to estimate the number of clusters***



Shows a plot of the SSE vs the no of clusters for a (bisecting) k-means clustering of the data

# Internal Measures: SSE

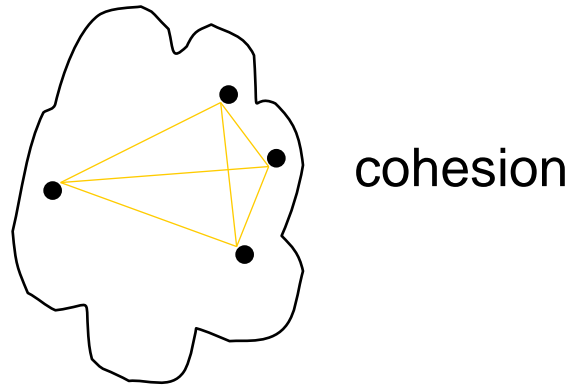
- SSE curve for a more complicated data set



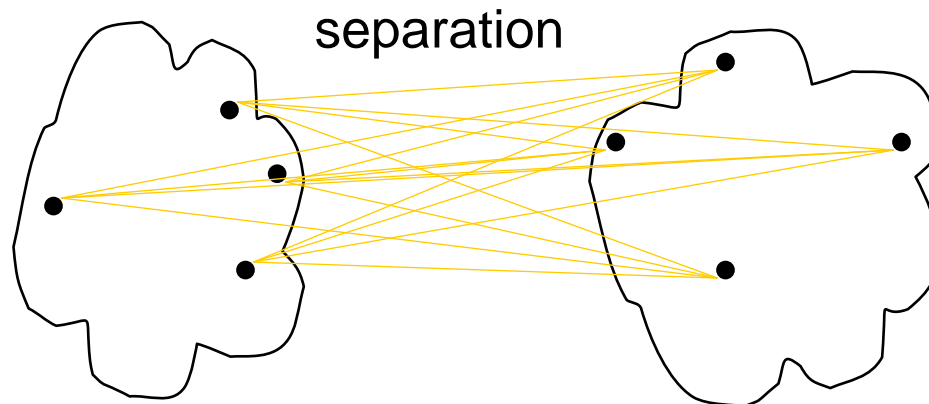
**SSE of clusters found using K-means**

# Internal Measures: Cohesion and Separation

**Cluster Cohesion:** Measures how closely related are objects in a cluster



**Cluster Separation:** Measure how distinct or well-separated a cluster is from other clusters

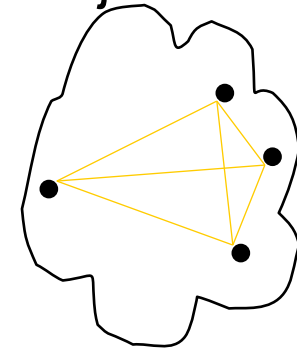


# Internal Measures: Cluster Cohesion

- **Cohesion**: Measures how closely related are objects in a cluster

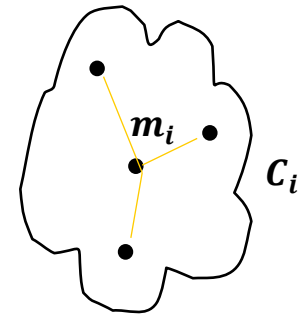
$$\text{Cohesion } (C_i) = \sum_{x, y \in C_i} \text{proximity}(\mathbf{x}, \mathbf{y})$$

The proximity function can be a similarity or a dissimilarity.



- **Cohesion can be centroid based**

$$\text{Cohesion } (C_i) = \sum_{x \in C_i} \text{proximity}(\mathbf{x}, \mathbf{m}_i)$$



Cohesion is within cluster sum of squares (SSE) if we let proximity to be squared Euclidean distance

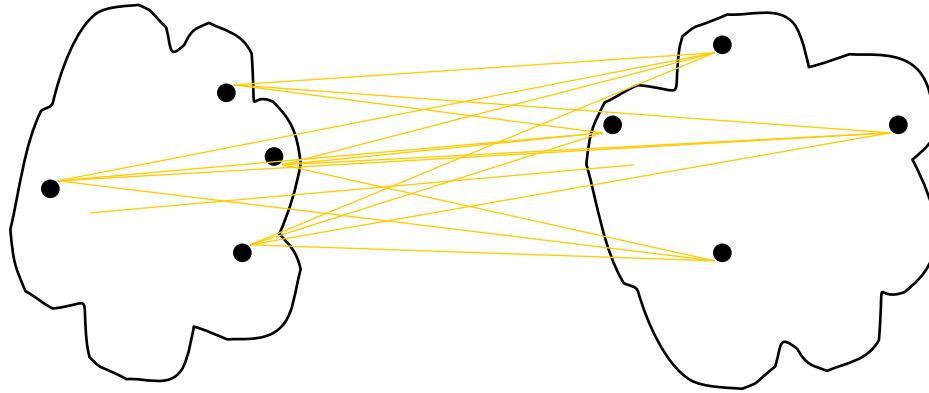
$$WSS = \sum_i \sum_{x \in C_i} (x - m_i)^2$$

where  $C_i$  is the cluster  $i$ ,  $m_i$  is the mean of cluster  $i$

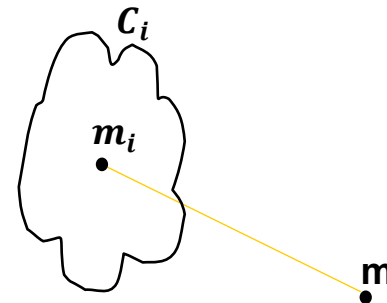
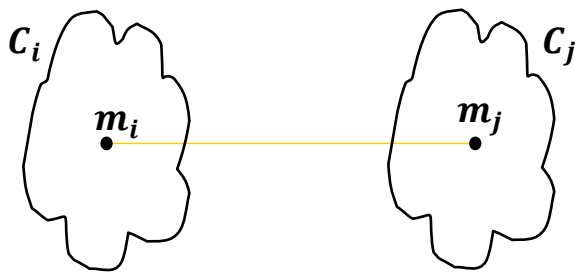


# Internal Measures: Separation

**Separation:** Measure how distinct or well-separated a cluster is from other clusters



$$\text{Separation}(C_i, C_j) = \sum_{x \in C_i, y \in C_j} \text{proximity}(x, y)$$



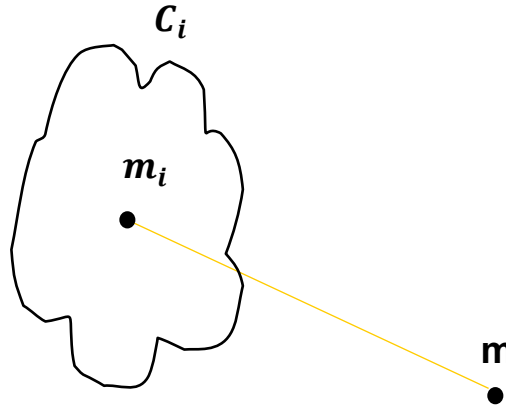
$$\text{Separation}(C_i, C_j) = \text{proximity}(m_i, m_j)$$

$$\text{Separation}(C_i) = \text{proximity}(m_i, m)$$

where  $|C_i|$  is the size of cluster  $i$ ,  $m_i$  is the mean of cluster  $i$  and  $m$  is the overall mean of all data points

## Internal Measures: Separation

**Separation:** Measure how distinct or well-separated a cluster is from other clusters



$$\text{Separation}(C_i) = \text{proximity}(C_i, C_j)$$

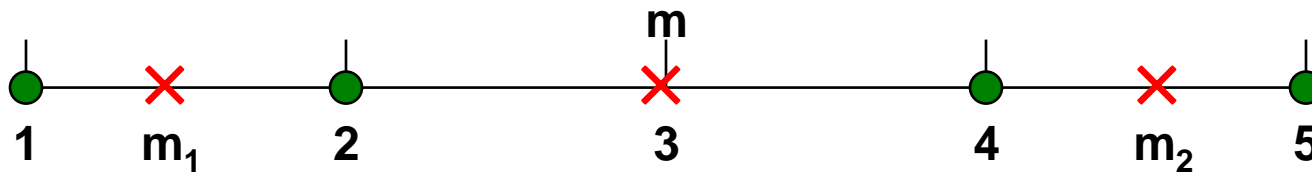
$$BSS = \sum_i |C_i| (m - m_i)^2$$

**Separation is measured by the between cluster sum of squares**  
where  $|C_i|$  is the size of cluster  $i$ ,  $m_i$  is the mean of cluster  $i$  and  $m$  is the overall mean of all data points

# Internal Measures: Cohesion and Separation

- Example:  $SSE = WSS(\text{Cohesion}) + BSS(\text{separation}) = \text{constant}$

$$WSS = \sum_i \sum_{x \in C_i} (x - m_i)^2 \quad BSS = \sum_i |C_i| (m - m_i)^2$$



**K=1 cluster:**

$$WSS = (1 - 3)^2 + (2 - 3)^2 + (4 - 3)^2 + (5 - 3)^2 = 10$$

$$BSS = 4 \times (3 - 3)^2 = 0$$

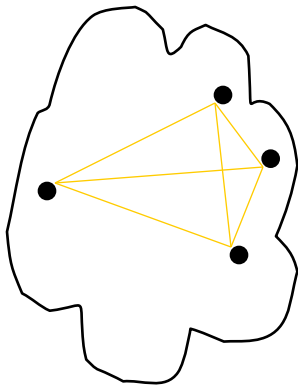
**K=2 clusters:**

$$WSS = (1 - 1.5)^2 + (2 - 1.5)^2 + (4 - 4.5)^2 + (5 - 4.5)^2 = 1$$

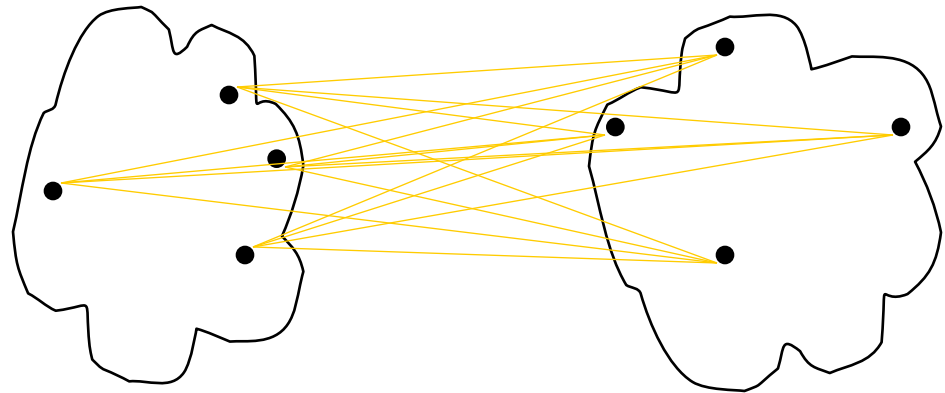
$$BSS = 2 \times (3 - 1.5)^2 + 2 \times (3 - 4.5)^2 = 9$$

# Internal Measures: Combine Cohesion & Separation

- We can combine the idea of cohesion and separation.
  - Cluster cohesion is the sum of the weight of all links within a cluster.
  - Cluster separation is the sum of the weights between nodes in the cluster and nodes outside the cluster.



cohesion



separation

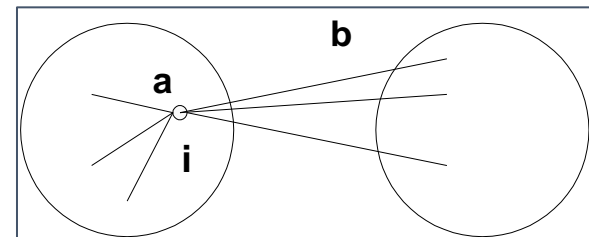
# Internal Measures: Silhouette Coefficient

- Silhouette Coefficient combines ideas of cohesion & separation
- Silhouette Coefficient for an individual point,  $i$ 
  - Calculate  $a$  = average distance of  $i$  to the points in its cluster
  - Calculate  $b$  = min (average distance of  $i$  to points in another cluster)

**The silhouette coefficient for a point is**  
 **$s = 1 - a/b$  if  $a < b$**

**(or  $s = b/a - 1$  if  $a \geq b$ , not the usual case)**

Typically between 0 and 1.  
The closer to 1 the better.



**Average Silhouette width for a cluster is the average of silhouette coefficients of points in the cluster**