Clustering fundamentals



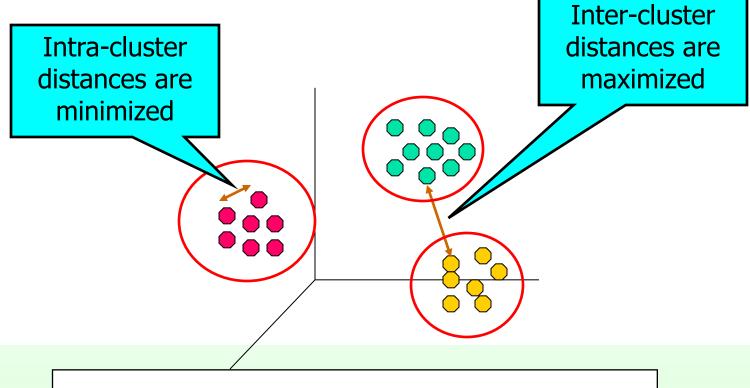
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Politecnico di Torino

What is Cluster Analysis?



 Finding groups of objects such that the objects in a group will be similar (or related) to one another and different from (or unrelated to) the objects in other groups





Applications of Cluster Analysis



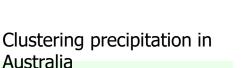
Understanding

 Group related documents for browsing, group genes and proteins that have similar functionality, or group stocks with similar price fluctuations

Summarization

Reduce the size of large data sets

	Discovered Clusters	Industry Group
1	Applied-Matl-DOWN,Bay-Network-Down,3-COM-DOWN, Cabletron-Sys-DOWN,CISCO-DOWN,HP-DOWN, DSC-Comm-DOWN,INTEL-DOWN,LSI-Logic-DOWN, Micron-Tech-DOWN,Texas-Inst-Down,Tellabs-Inc-Down, Natl-Semiconduct-DOWN,Oracl-DOWN,SGI-DOWN, Sun-DOWN	Technology1-DOWN
2	Apple-Comp-DOWN,Autodesk-DOWN,DEC-DOWN, ADV-Micro-Device-DOWN,Andrew-Corp-DOWN, Computer-Assoc-DOWN,Circuit-City-DOWN, Compaq-DOWN, EMC-Corp-DOWN, Gen-Inst-DOWN, Motorola-DOWN,Microsoft-DOWN,Scientific-Atl-DOWN	Technology2-DOWN
3	Fannie-Mae-DOWN,Fed-Home-Loan-DOWN, MBNA-Corp-DOWN,Morgan-Stanley-DOWN	Financial-DOWN
4	Baker-Hughes-UP,Dresser-Inds-UP,Halliburton-HLD-UP, Louisiana-Land-UP,Phillips-Petro-UP,Unocal-UP, Schlumberger-UP	Oil-UP



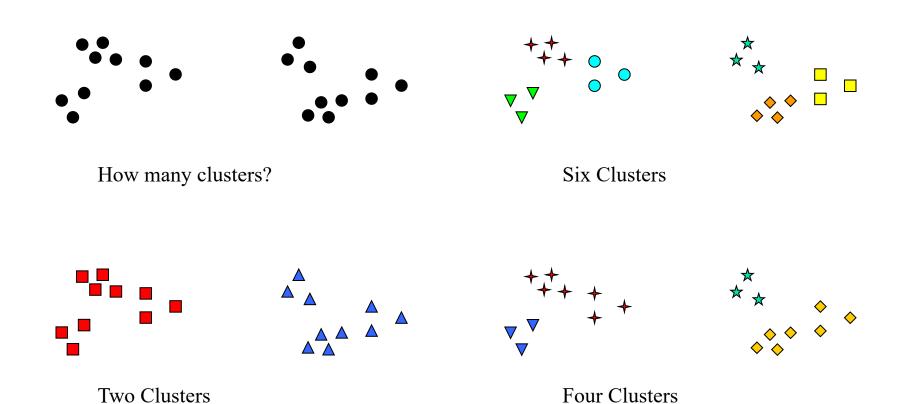




Australia

Notion of a Cluster can be Ambiguous







Types of Clusterings

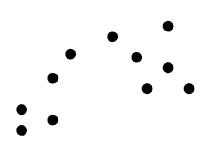


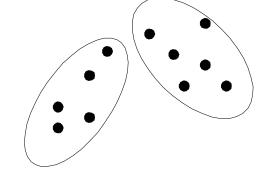
- A clustering is a set of clusters
- Important distinction between hierarchical and partitional sets of clusters
- Partitional Clustering
 - Divides data objects into non-overlapping subsets (clusters) such that each data object is in exactly one subset
- Hierarchical clustering
 - A set of nested clusters organized as a hierarchical tree

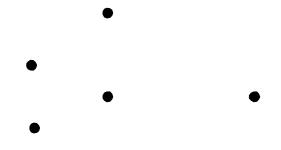


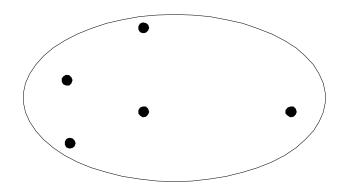
Partitional Clustering











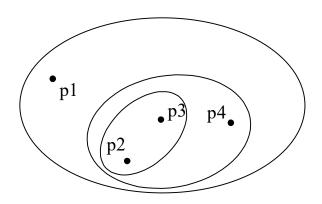
Original Points

A Partitional Clustering

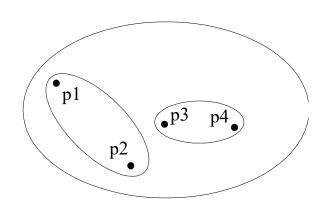


Hierarchical Clustering

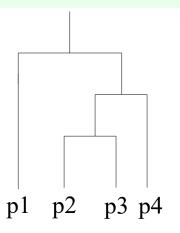




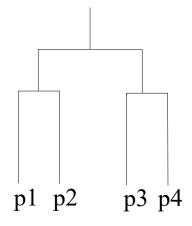
Traditional Hierarchical Clustering



Non-traditional Hierarchical Clustering



Traditional Dendrogram



Non-traditional Dendrogram



Other Distinctions Between Sets of Clusters



Exclusive versus non-exclusive

In non-exclusive clustering, points may belong to multiple clusters.

Fuzzy versus non-fuzzy

- In fuzzy clustering, a point belongs to every cluster with some weight between 0 and 1
- Weights must sum to 1
- Probabilistic clustering has similar characteristics

Partial versus complete

In some cases, we only want to cluster some of the data

Heterogeneous versus homogeneous

Cluster of widely different sizes, shapes, and densities



Types of Clusters



- Well-separated clusters
- Center-based clusters
- Contiguous clusters
- Density-based clusters
- Property or Conceptual
- Described by an Objective Function

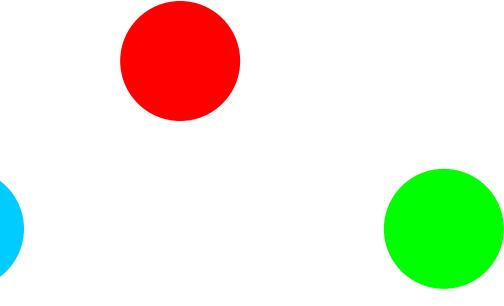


Types of Clusters: Well Separated



Well-Separated Clusters:

 A cluster is a set of points such that any point in a cluster is closer (or more similar) to every other point in the cluster than to any point not in the cluster.





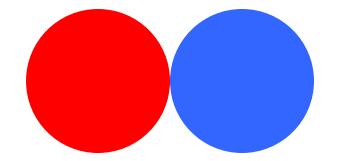
3 well-separated clusters

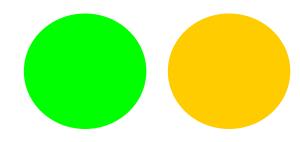
Types of Clusters: Center-Based



Center-based

- A cluster is a set of objects such that an object in a cluster is closer (more similar) to the "center" of a cluster, than to the center of any other cluster
- The center of a cluster is often a centroid, the average of all the points in the cluster, or a medoid, the most "representative" point of a cluster





4 center-based clusters



Types of Clusters: Contiguity-Based



- Contiguous Cluster (Nearest neighbor or Transitive)
 - A cluster is a set of points such that a point in a cluster is closer (or more similar) to one
 or more other points in the cluster than to any point not in the cluster.





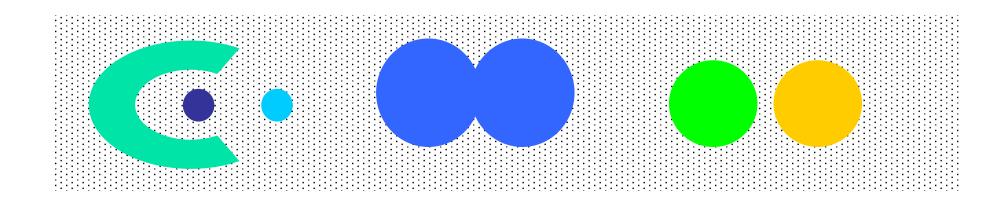
8 contiguous clusters

Types of Clusters: Density-Based



Density-based

- A cluster is a dense region of points, which is separated by low-density regions, from other regions of high density.
- Used when the clusters are irregular or intertwined, and when noise and outliers are present.



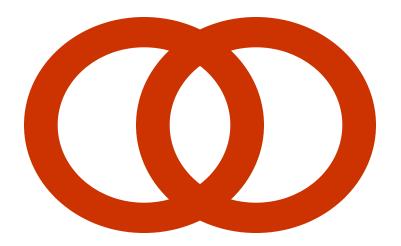


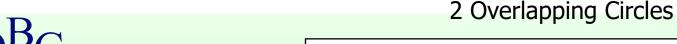
6 density-based clusters

Types of Clusters: Conceptual Clusters



- Shared Property or Conceptual Clusters
 - Finds clusters that share some common property or represent a particular concept.







Clustering Algorithms



- K-means and its variants
- Hierarchical clustering
- Density-based clustering



K-means Clustering



- Partitional clustering approach
- Each cluster is associated with a centroid (center point)
- Each point is assigned to the cluster with the closest centroid
- Number of clusters, K, must be specified
- The basic algorithm is very simple

1: Select K points as the initial centroids.

2: repeat

Form K clusters by assigning all points to the closest centroid.

4: Recompute the centroid of each cluster.

5: **until** The centroids don't change



K-means Clustering – Details

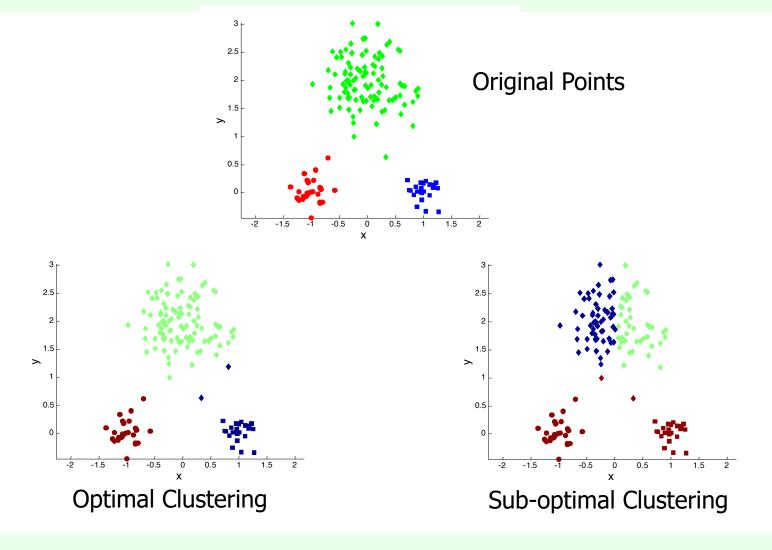


- Initial centroids are often chosen randomly.
 - Clusters produced vary from one run to another.
- The centroid is (typically) the mean of the points in the cluster.
- 'Closeness' is measured by Euclidean distance, cosine similarity, correlation, etc.
- K-means will converge for common similarity measures mentioned above.
- 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(n * K * I * d)
 - n = number of points, K = number of clusters,
 I = number of iterations, d = number of attributes



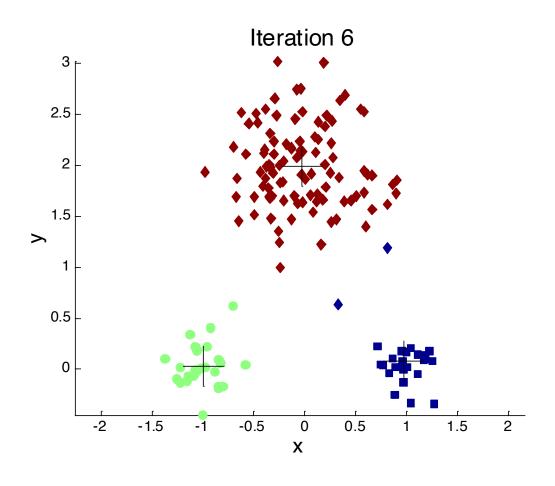
Two different K-means Clusterings





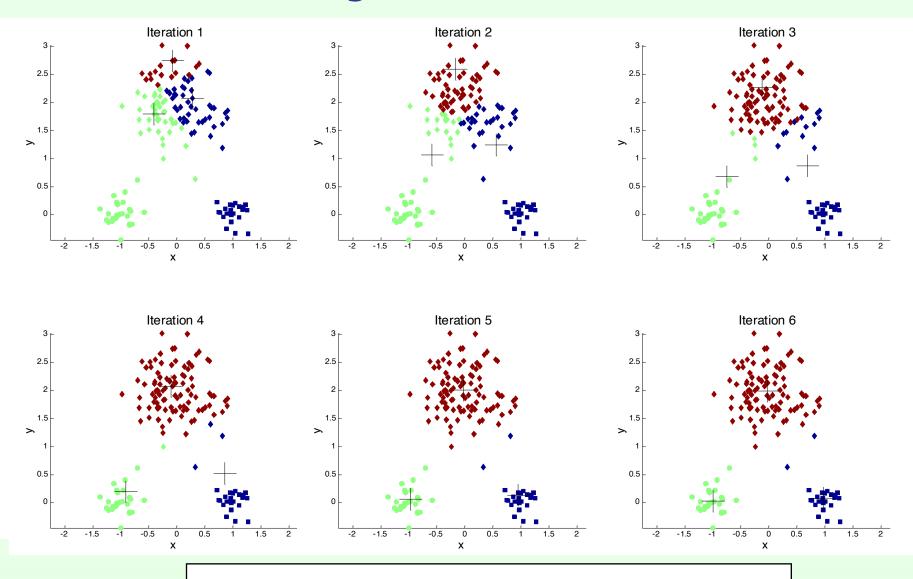






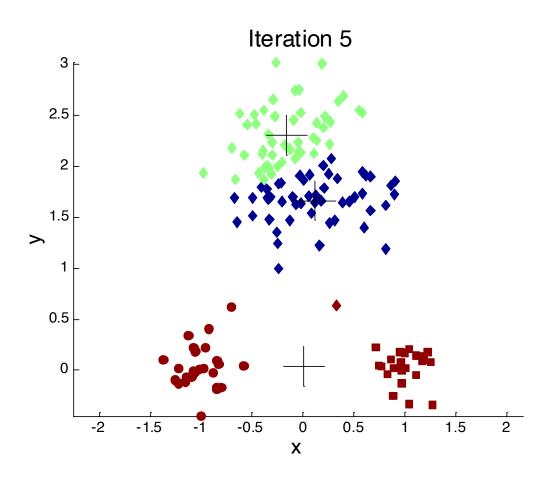






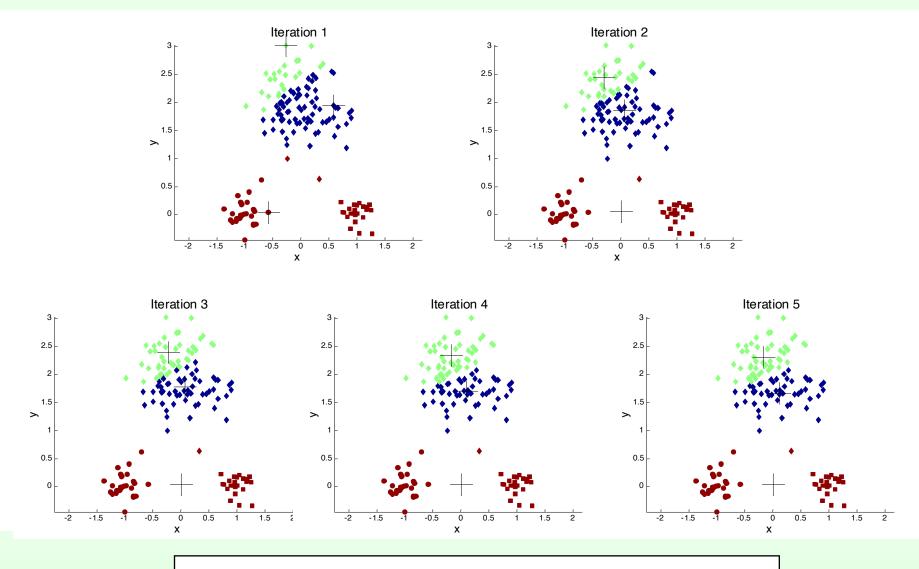














Evaluating K-means Clusters



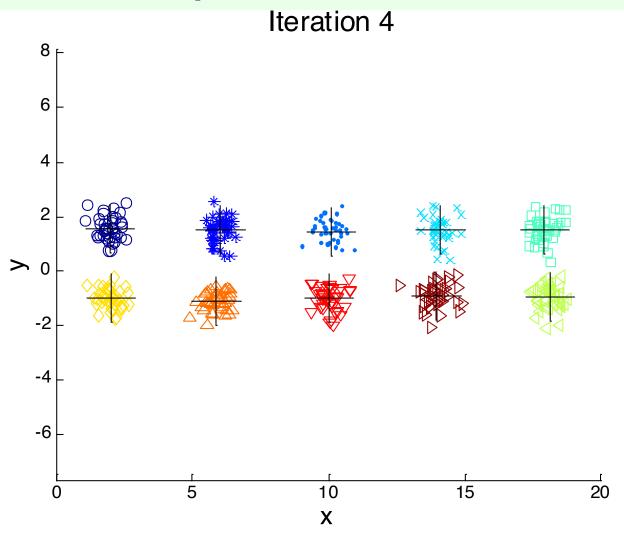
- Most common measure is Sum of Squared Error (SSE)
 - For each point, the error is the distance to the nearest cluster
 - To get SSE, we square these errors and sum them.

$$SSE = \sum_{i=1}^{K} \sum_{x \in C_i} dist^2(m_i, x)$$

- x is a data point in cluster C_i and m_i is the representative point for cluster C_i
 - can show that m_i corresponds to the center (mean) of the cluster
- Given two clusters, we can choose the one with the smallest error
- One easy way to reduce SSE is to increase K, the number of clusters
 - A good clustering with smaller K can have a lower SSE than a poor clustering with higher K



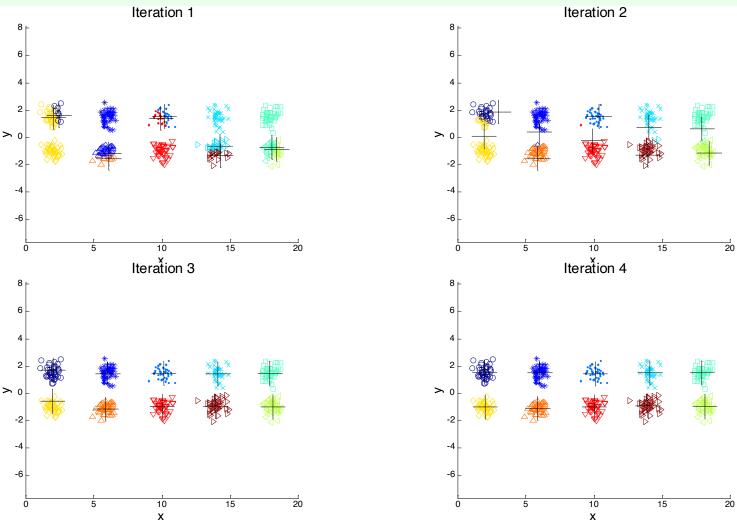






Starting with two initial centroids in one cluster of each pair of clusters



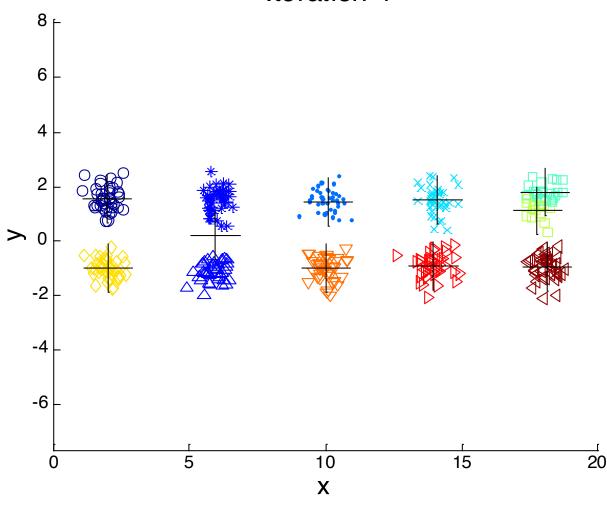




Starting with two initial centroids in one cluster of each pair of clusters



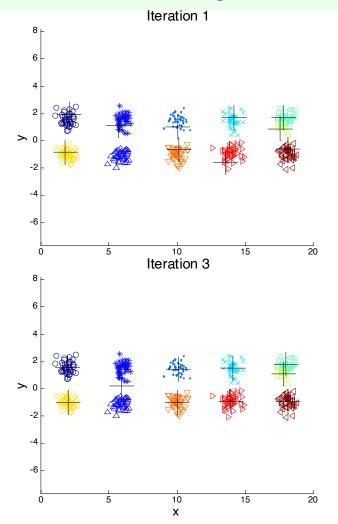


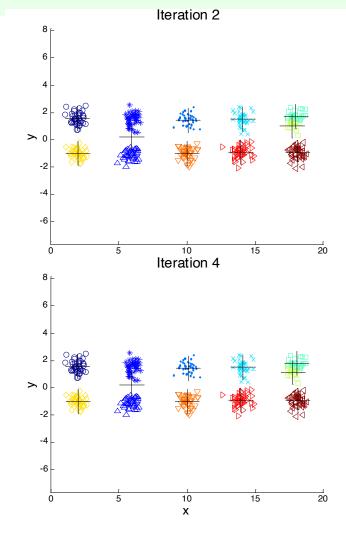




Starting with some pairs of clusters having three initial centroids, while other have only one.









Starting with some pairs of clusters having three initial centroids, while other have only one.

Solutions to Initial Centroids Problem



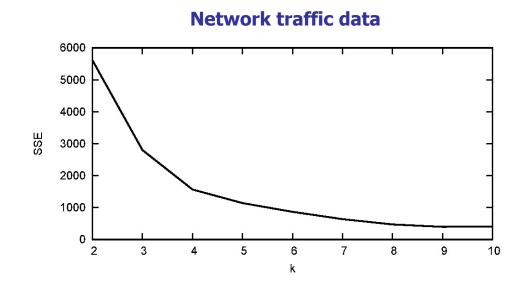
- Multiple runs
 - Helps, but probability is not on your side
- Sample and use hierarchical clustering to determine initial centroids
- Select more than k initial centroids and then select among these initial centroids
 - Select most widely separated
- Postprocessing
- Bisecting K-means
 - Not as susceptible to initialization issues



K-means parameter setting



- Elbow graph (Knee approach)
 - Plotting the quality measure trend (e.g., SSE) against K
 - Choosing the value of K
 - the gain from adding a centroid is negligible
 - The reduction of the quality measure is not interesting anymore



Medical records





Handling Empty Clusters



- Basic K-means algorithm can yield empty clusters
- Several strategies
 - Choose the point that contributes most to SSE
 - Choose a point from the cluster with the highest SSE
 - If there are several empty clusters, the above can be repeated several times.



Pre-processing and Post-processing



- Pre-processing
 - Normalize the data
 - Eliminate outliers
- Post-processing
 - Eliminate small clusters that may represent outliers
 - Split 'loose' clusters, i.e., clusters with relatively high SSE
 - Merge clusters that are 'close' and that have relatively low SSE
 - Can use these steps during the clustering process



Bisecting K-means



Bisecting K-means algorithm

Variant of K-means that can produce a partitional or a hierarchical clustering

1: Initialize the list of clusters to contain the cluster containing all points.

2: repeat

3: Select a cluster from the list of clusters

4: **for** i = 1 to $number_of_iterations$ **do**

5: Bisect the selected cluster using basic K-means

6: end for

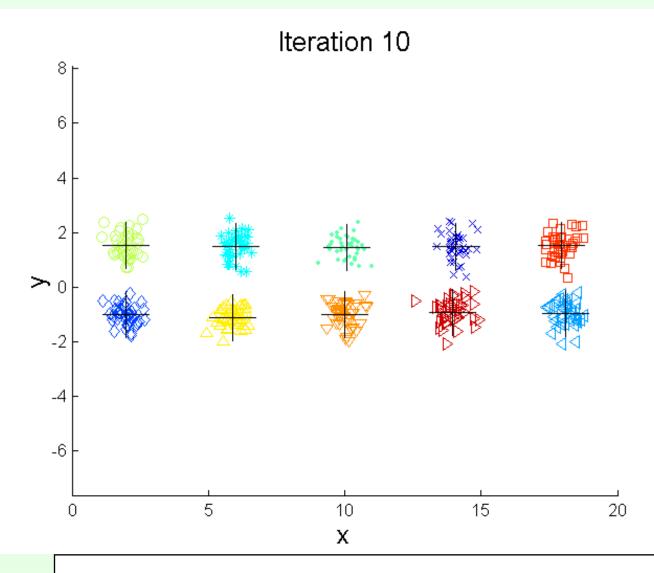
7: Add the two clusters from the bisection with the lowest SSE to the list of clusters.

8: until Until the list of clusters contains K clusters



Bisecting K-means Example







Limitations of K-means



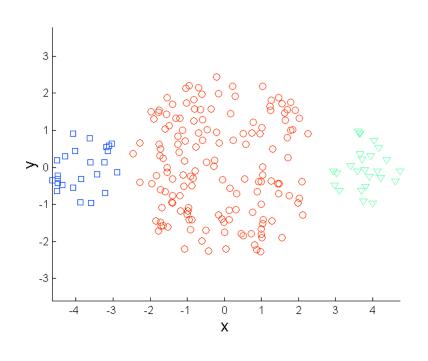
- K-means has problems when clusters are of differing
 - Sizes
 - Densities
 - Non-globular shapes

K-means has problems when the data contains outliers.



Limitations of K-means: Differing Sizes





3 - 2 - 1 0 1 2 3 4 X

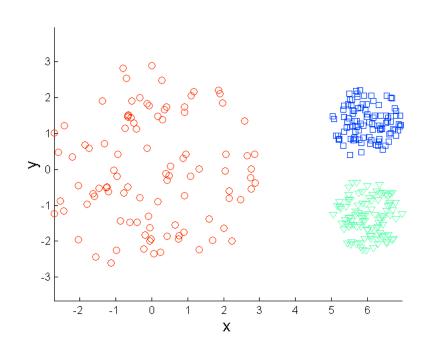
Original Points

K-means (3 Clusters)



Limitations of K-means: Differing Density





3 - 2 - 1 0 1 2 3 4 5 6 X

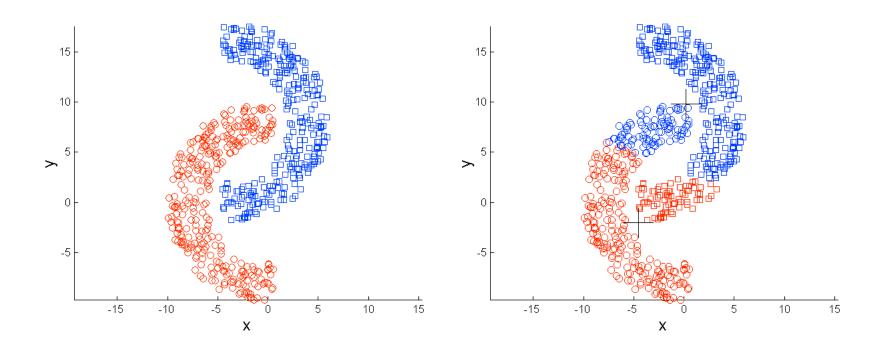
Original Points

K-means (3 Clusters)



Limitations of K-means: Non-globular Shapes





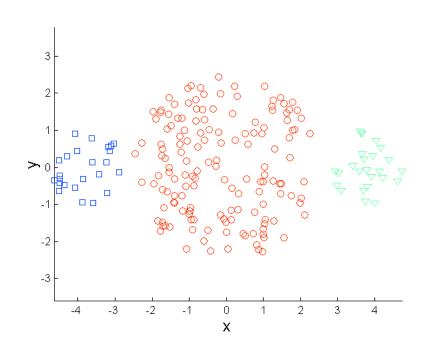
Original Points

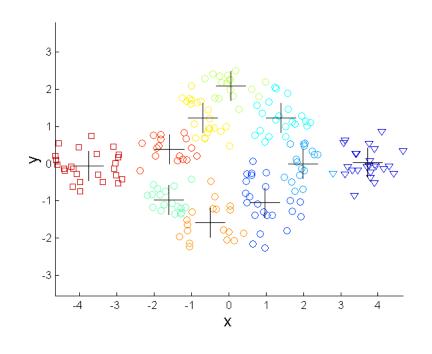
K-means (2 Clusters)



Overcoming K-means Limitations







Original Points

K-means Clusters

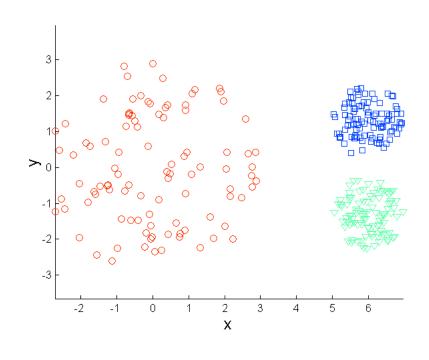
One solution is to use many clusters.

Find parts of clusters, but need to put together.



Overcoming K-means Limitations





X

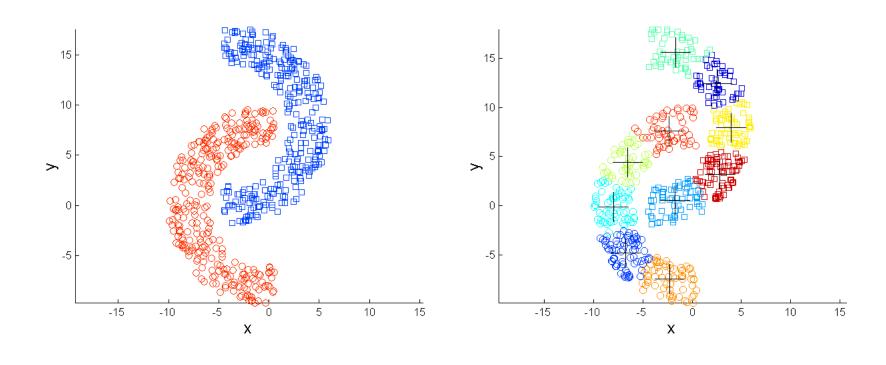
Original Points

K-means Clusters



Overcoming K-means Limitations





Original Points

K-means Clusters



Hierarchical Clustering

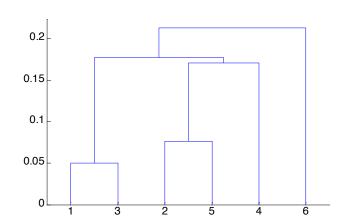


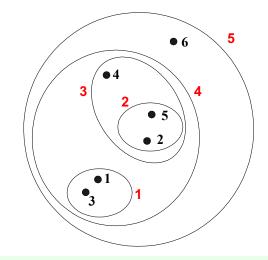
 Produces a set of nested clusters organized as a hierarchical tree

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 dendogram 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 a 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



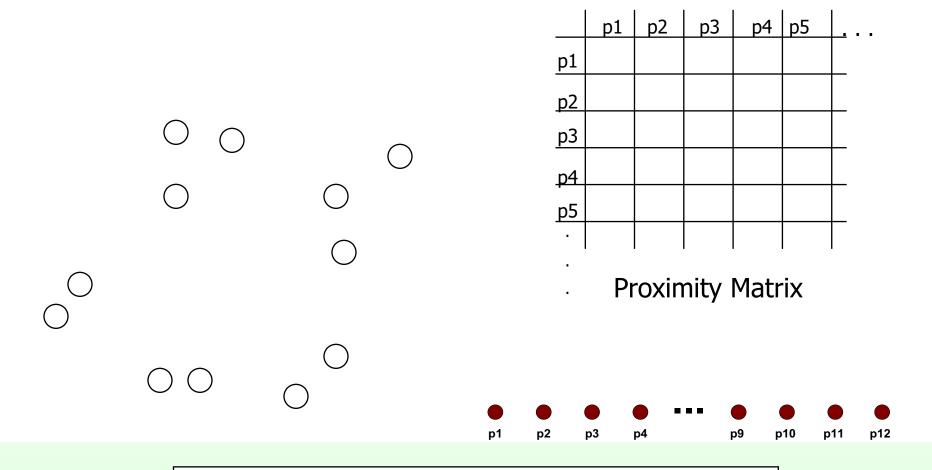
- More popular hierarchical clustering technique
- Basic algorithm is straightforward
 - 1. Compute the proximity matrix
 - Let each data point be a cluster
 - 3. Repeat
 - 4. Merge the two closest clusters
 - 5. Update the proximity matrix
 - **6. Until** only a single cluster remains
- Key operation 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

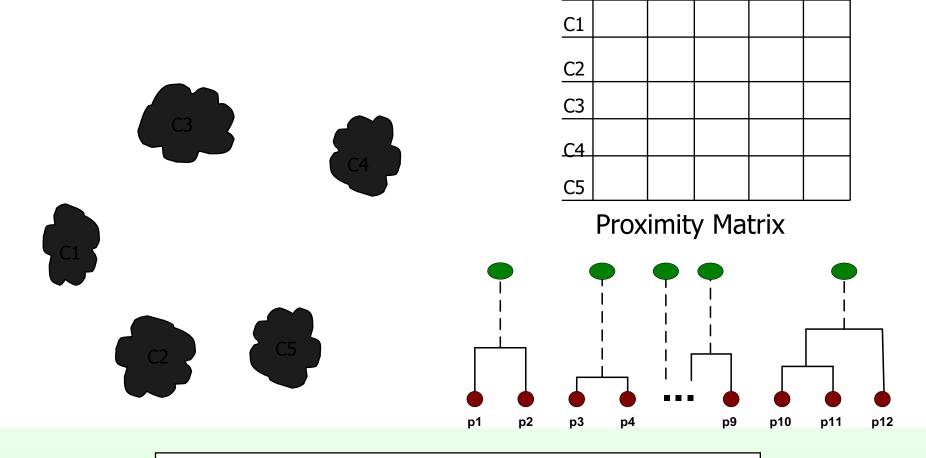




Intermediate Situation



After some merging steps, we have some clusters



C2

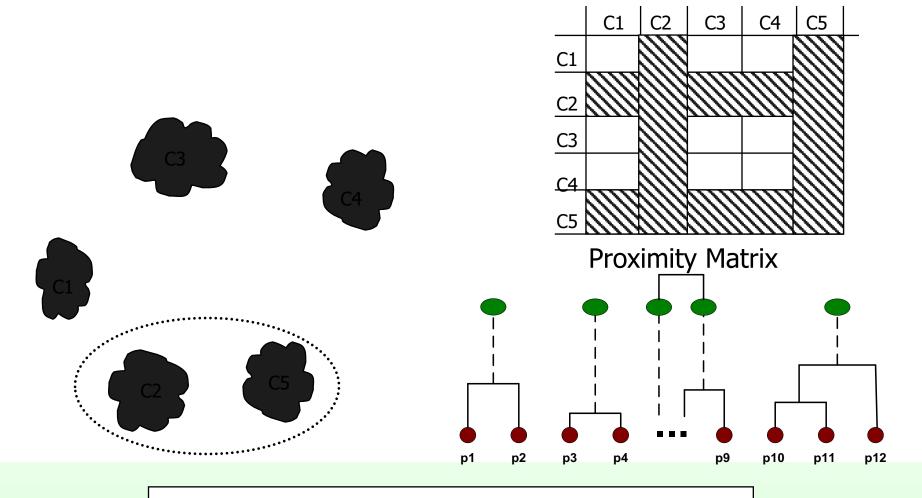
C4



Intermediate Situation



We want to merge the two closest clusters (C2 and C5) and update the proximity matrix.

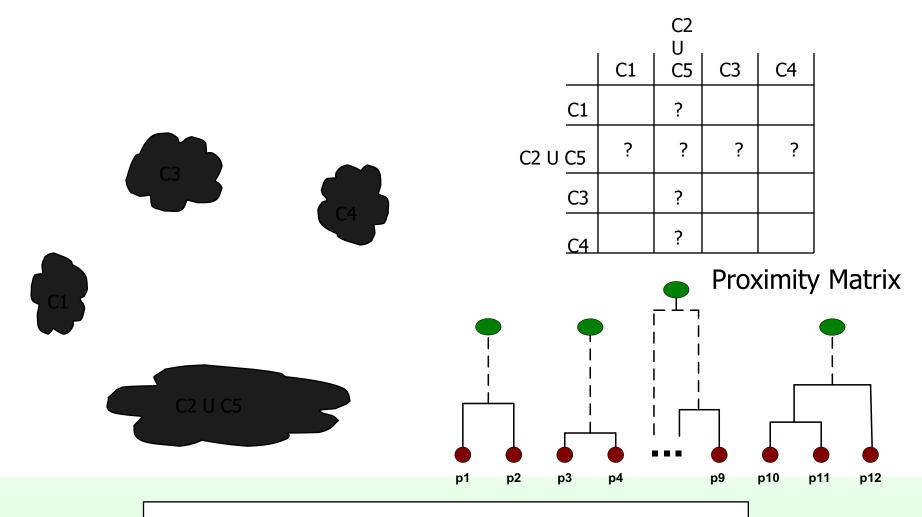




After Merging

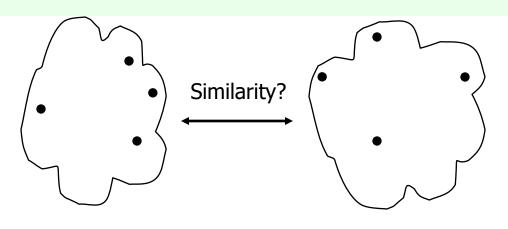


The question is "How do we update the proximity matrix?"









	p1	p2	р3	p4	p5	<u> </u>
p1						
p2						
p2 p3						
<u>p4</u> p5						_
_						

- MIN
- MAX
- Group Average

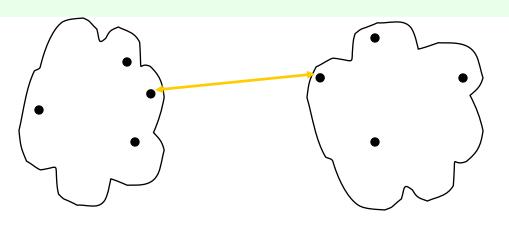
function

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

 $D_{M}^{B}G$

Proximity Matrix





	p1	p2	р3	p4	p5	<u> </u>
p1						
p2						
p2 p3						
<u>p4</u> p5						

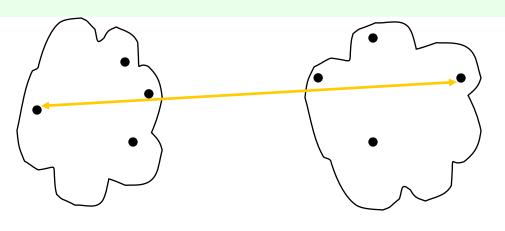
Proximity Matrix

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



50





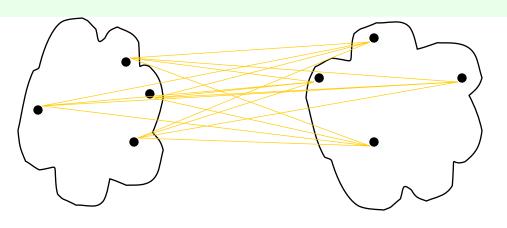
	p1	p2	р3	p4	p5	<u> </u>
p1						
p2						
p2 p3						
<u>p4</u> p5						

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



Proximity Matrix





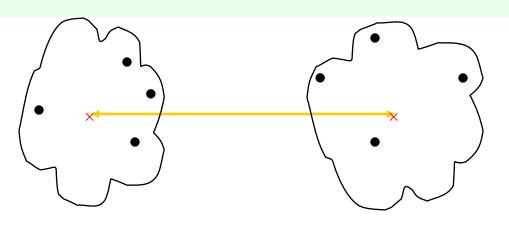
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	p1	p2	рЗ	p4	р5	<u> </u>
p1						
p2						
p2 p3						
						_
<u>p4</u> p5						

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

 D_{MG}^{B}

Proximity Matrix





	p1	p2	р3	p4	p5	
	Ρ-	ρ2	PS	Ρ'	PS	<u> </u>
p1						
p2						
<u>р2</u> р3						
<u>р4</u> р5						

Proximity Matrix

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



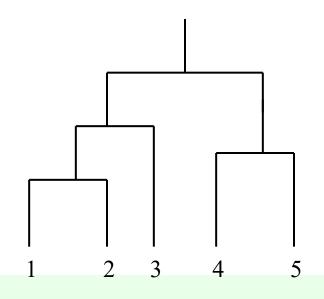
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Cluster Similarity: MIN or Single Link



- Similarity of two clusters is based on the two most similar (closest) points in the different clusters
 - Determined by one pair of points, i.e., by one link in the proximity graph.

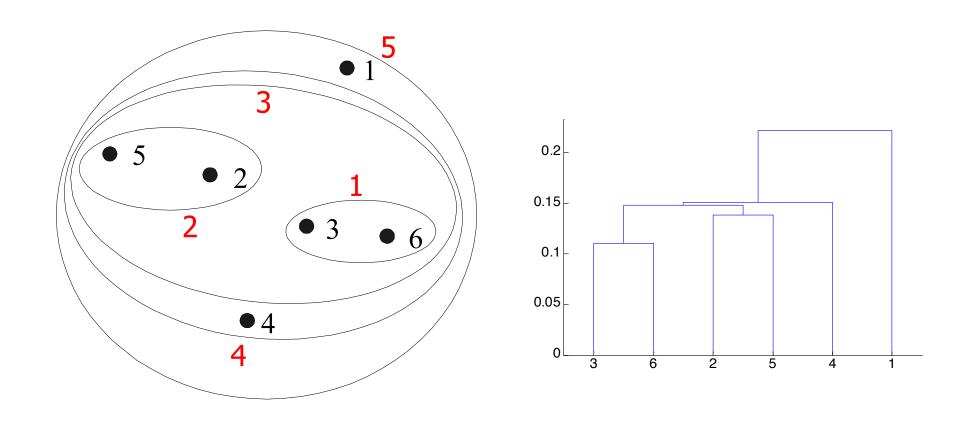
	I 1	12	13	14	15
11	1.00	0.90	0.10	0.65	0.20
12	0.90	1.00	0.70	0.60	0.50
13	0.10	0.70	1.00	0.40	0.30
14	0.65	0.60	0.40	1.00	0.80
15	0.20	0.50	0.30	0.80	0.20 0.50 0.30 0.80 1.00





Hierarchical Clustering: MIN





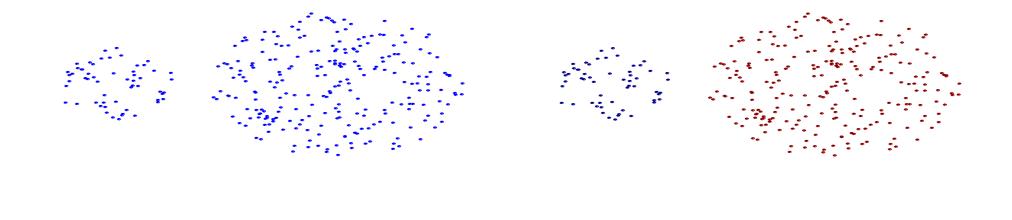
Nested Clusters

Dendrogram



Strength of MIN





Two Clusters

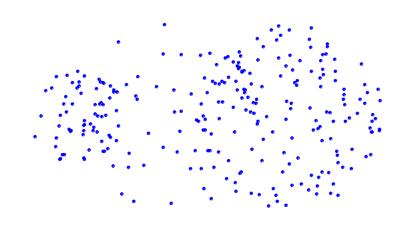
• Can handle non-elliptical shapes

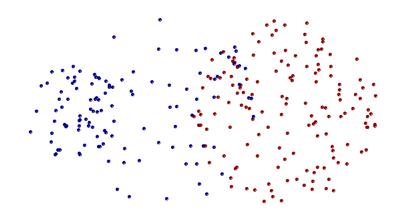
Original Points



Limitations of MIN







Original Points

Two Clusters

• Sensitive to noise and outliers

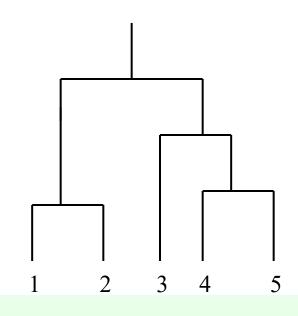


Cluster Similarity: MAX or Complete Linkage



- Similarity of two clusters is based on the two least similar (most distant) points in the different clusters
 - Determined by all pairs of points in the two clusters

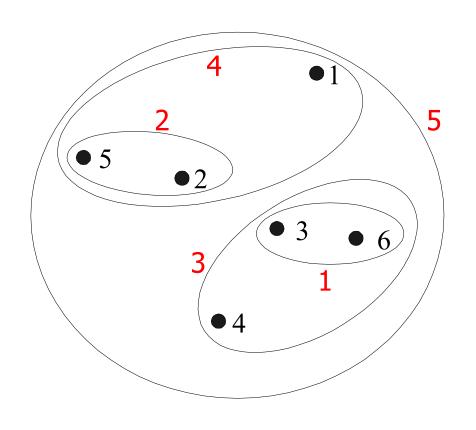
_				14	
11	1.00	0.90	0.10	0.65	0.20
12	0.90	1.00	0.70	0.60	0.50
13	0.10	0.70	1.00	0.40	0.30
14	0.65	0.60	0.40	1.00	0.80
15	0.20	0.50	0.30	0.80	0.20 0.50 0.30 0.80 1.00

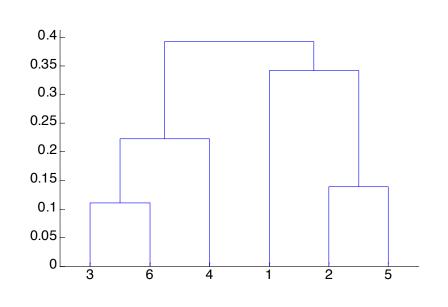




Hierarchical Clustering: MAX







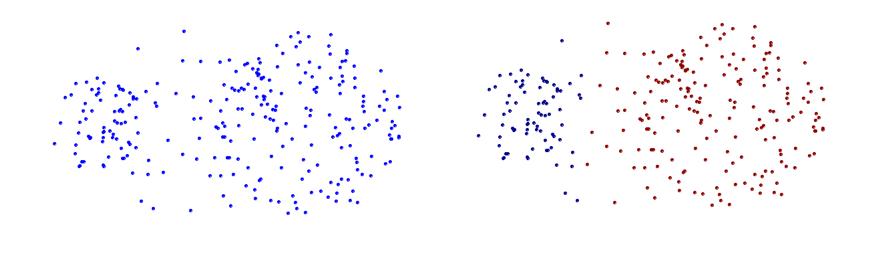
Nested Clusters

Dendrogram



Strength of MAX





• Less susceptible to noise and outliers

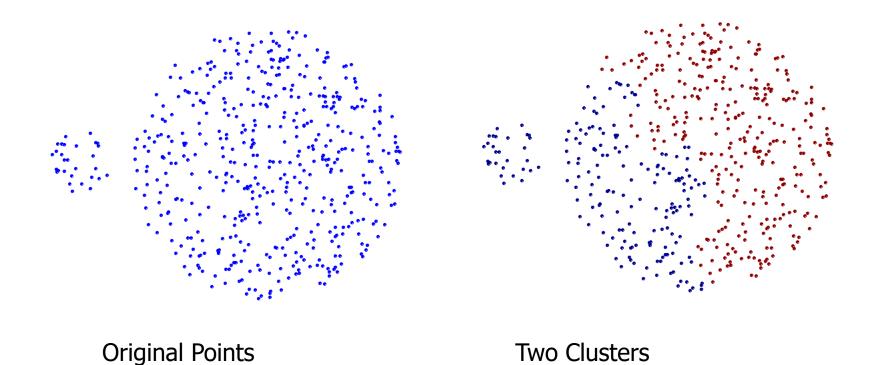
Original Points



Two Clusters

Limitations of MAX





- •Tends to break large clusters
- •Biased towards globular clusters



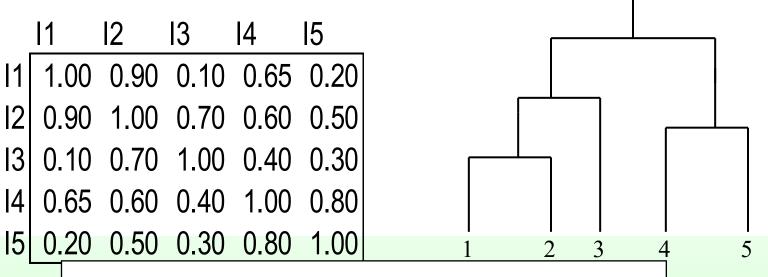
Cluster Similarity: Group Average



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

$$proximity(Cluster_{i}, Cluster_{j}) = \frac{\sum_{\substack{p_{i} \in Cluster_{i} \\ p_{j} \in Cluster_{j}}}}{|Cluster_{i}| * |Cluster_{i}|}$$

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

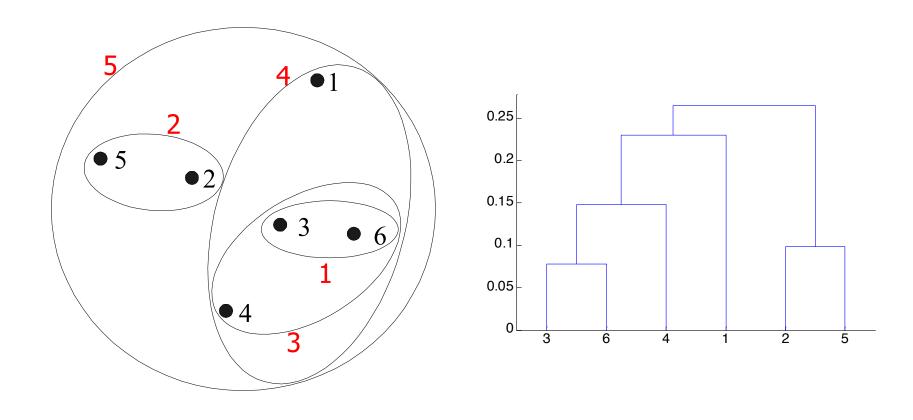




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Hierarchical Clustering: Group Average





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

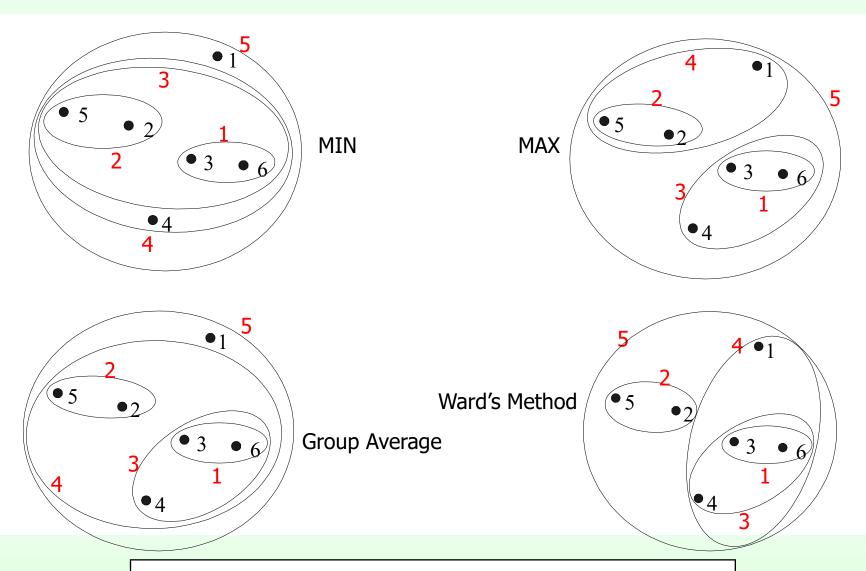


- Similarity of two clusters is based on the increase in squared error when two clusters are merged
 - Similar to group average if distance between points is distance squared
- Less susceptible to noise and outliers
- Biased towards globular clusters
- Hierarchical analogue of K-means
 - Can be used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



Hierarchical Clustering: Comparison







Hierarchical Clustering: Time and Space requirements



- O(N²) space since it uses the proximity matrix.
 - N is the number of points.

- O(N³) time in many cases
 - There are N steps and at each step the size, N², proximity matrix must be updated and searched
 - Complexity can be reduced to O(N² log(N)) time for some approaches



DBSCAN

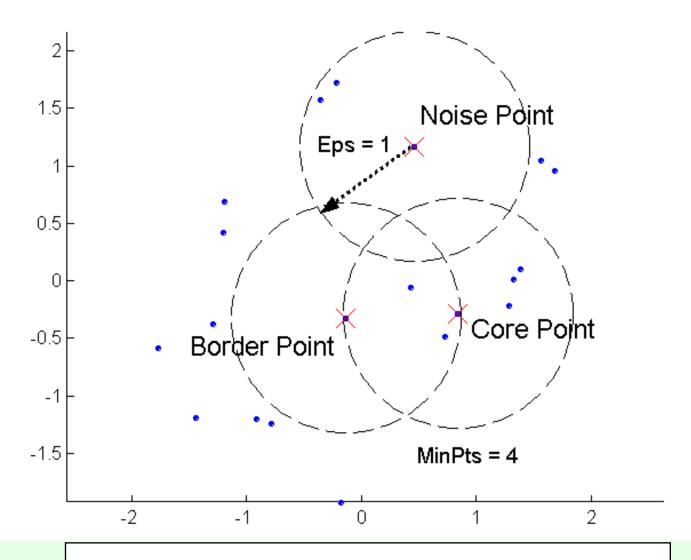


- DBSCAN is a density-based algorithm
 - Density = number of points within a specified radius (Eps)
 - A point is a core point if it has more than a specified number of points (MinPts) within Eps
 - These are points that are at the interior of a cluster
 - A border point has fewer than MinPts within Eps, but is in the neighborhood of a core point
 - A noise point is any point that is not a core point or a border point.



DBSCAN: Core, Border, and Noise Points







DBSCAN Algorithm



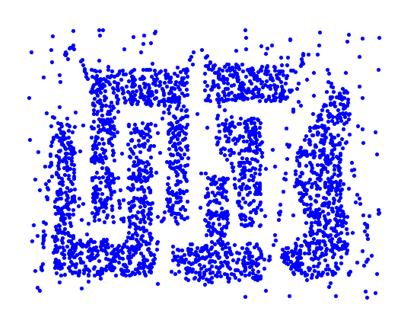
- Eliminate noise points
- Perform clustering on the remaining points

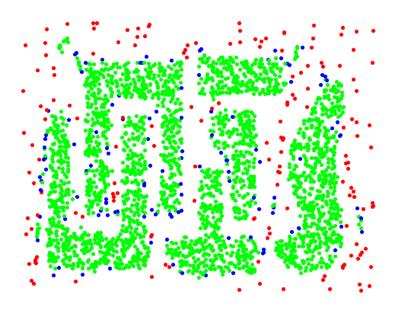
```
current\ cluster\ label \leftarrow 1
for all core points do
  if the core point has no cluster label then
     current\_cluster\_label \leftarrow current\_cluster\_label + 1
     Label the current core point with cluster label current\_cluster\_label
  end if
  for all points in the Eps-neighborhood, except i^{th} the point itself do
    if the point does not have a cluster label then
       Label the point with cluster label current_cluster_label
     end if
  end for
end for
```



DBSCAN: Core, Border, and Noise Points







Original Points

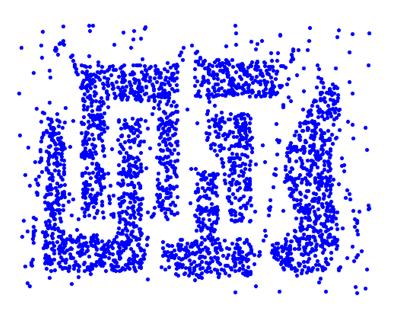
Point types: core, border and noise

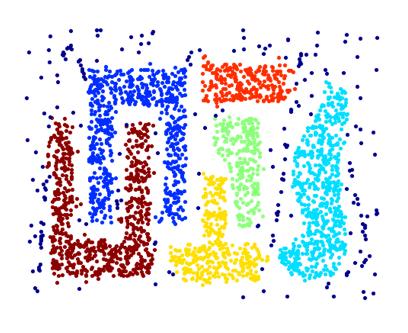


Eps = 10, MinPts = 4

When DBSCAN Works Well







Original Points

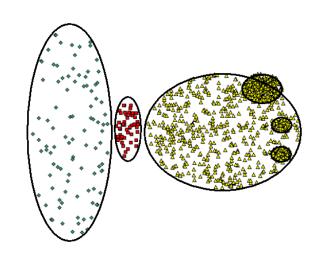
Clusters

- Resistant to Noise
- Can handle clusters of different shapes and sizes



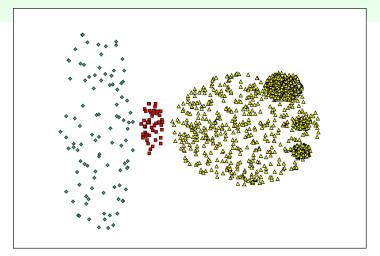
When DBSCAN Does NOT Work Well



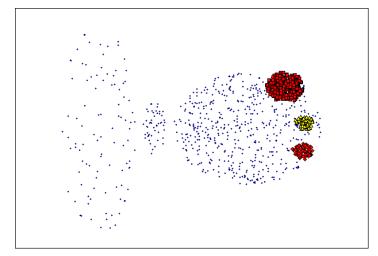


Original Points

- Varying densities
- High-dimensional data



(MinPts=4, Eps=9.75).



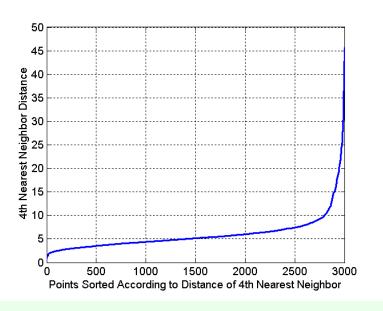
(MinPts=4, Eps=9.62)



DBSCAN: Determining EPS and MinPts



- Idea is that for points in a cluster, their kth nearest neighbors are at roughly the same distance
- Noise points have the kth nearest neighbor at farther distance
- So, plot sorted distance of every point to its kth nearest neighbor







Elena Baralis, Tania Cerquitelli

Politecnico di Torino

Cluster Validity

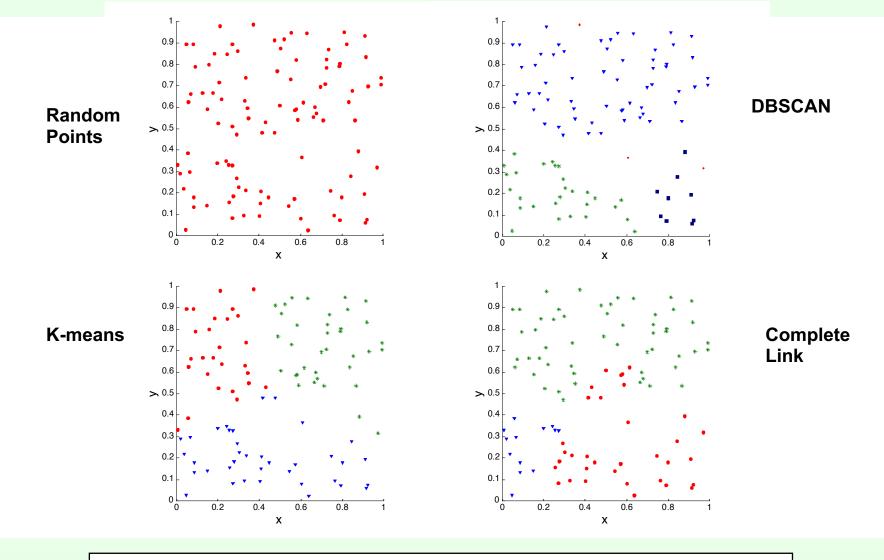


- For supervised classification we have a variety of measures to evaluate how good our model is
 - Accuracy, precision, recall
- For cluster analysis, the analogous question is how to evaluate the "goodness" of the resulting clusters?
- But "clusters are in the eye of the beholder"!
- Then why do we 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







Different Aspects of Cluster Validation



- Determining the clustering tendency of a set of data, i.e., distinguishing whether nonrandom structure actually exists in the data.
- 2. Comparing the results of a cluster analysis to externally known results, e.g., to externally given class labels.
- Evaluating how well the results of a cluster analysis fit the data without reference to external information.
 - Use only the data
- 4. Comparing the results of two different sets of cluster analyses to determine which is better.
- 5. Determining the 'correct' number of clusters.

For 2, 3, and 4, we can further distinguish whether we want to evaluate the entire clustering or just individual clusters.



Measures of Cluster Validity



- Numerical measures are applied to judge various aspects of cluster validity
- Numerical measures can be classified into three classes
 - External Index: Used to measure the extent to which cluster labels match externally supplied class labels.
 - e.g., entropy, purity
 - Internal Index: Used to measure the goodness of a clustering structure without respect to external information.
 - e.g., Sum of Squared Error (SSE), cluster cohesion, cluster separation, Rand-Index, adjusted rand-index, Silhouette index
 - 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



Internal Measures: Cohesion and Separation



- Cluster Cohesion: Measures how closely related are objects in a cluster
 - Cohesion is measured by the within cluster sum of squares (SSE)

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

- Cluster Separation: Measure how distinct or well-separated a cluster is from other clusters
 - Separation is measured by the between cluster sum of squares

$$BSS = \sum_{i} |C_i| (m - m_i)^2$$

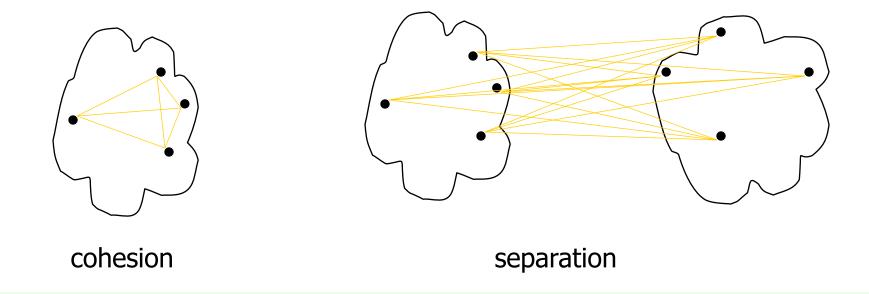
Where |C_i| is the size of cluster i



Internal Measures: Cohesion and Separation



- A proximity graph based approach can also be used for 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.





Internal measures: Silhouette



- A succinct measure to evaluate how well each object lies within its cluster
- It is defined for single points
- It considers both cohesion and separation
- Can be computed for
 - Individual points
 - Individual clusters
 - Clustering result



Internal measures: Silhouette



- For each object i
 - *a(i)*: the average dissimilarity of *i* with all other objects within the same cluster (the smaller the value, the better the assignment)
 - b(i): min(average dissimilarity of *i* to any other cluster, of which *i* is not a member)

$$s(i) = \frac{b(i) - a(i)}{\max\{a(i), b(i)\}}$$

- Ranges between -1 and +1
 - Typically between 0 and 1
 - The closer to 1, the better
- Silhouette for clusters and clusterings
 - The average s(i) over all data of a *cluster* measures how tightly grouped all the data in the cluster are
 - The average s(i) over all data of the dataset measures how appropriately the data has been clustered







Table 5.9. K-m	eans Clustering Results for LA Document Data S	et
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indicate the second sec											
Cluster	Entertainment	Financial	Foreign	Metro	National	Sports	Entropy	Purity			
1	3	5	40	506	96	27	1.2270	0.7474			
2	4	7	280	29	39	2	1.1472	0.7756			
3	1	1	1	7	4	671	0.1813	0.9796			
4	10	162	3	119	73	2	1.7487	0.4390			
5	331	22	5	70	13	23	1.3976	0.7134			
6	5	358	12	212	48	13	1.5523	0.5525			
Total	354	555	341	943	273	738	1.1450	0.7203			

entropy For each cluster, the class distribution of the data is calculated first, i.e., for cluster j we compute p_{ij} , the 'probability' that a member of cluster j belongs to class i as follows: $p_{ij} = m_{ij}/m_j$, where m_j is the number of values in cluster j and m_{ij} is the number of values of class i in cluster j. Then using this class distribution, the entropy of each cluster j is calculated using the standard formula $e_j = \sum_{i=1}^L p_{ij} \log_2 p_{ij}$, where the L is the number of classes. The total entropy for a set of clusters is calculated as the sum of the entropies of each cluster weighted by the size of each cluster, i.e., $e = \sum_{i=1}^K \frac{m_i}{m} e_j$, where m_j is the size of cluster j, K is the number of clusters, and m is the total number of data points.

purity Using the terminology derived for entropy, the purity of cluster j, is given by $purity_j = \max p_{ij}$ and the overall purity of a clustering by $purity = \sum_{i=1}^{K} \frac{m_i}{m} purity_j$.



Rand Index



Idea

Any two objects that are in the same cluster should be in the same class and vice versa

Given

- f_{00} = number of pairs of objects having a different class and a different cluster
- f_{01} = number of pairs of objects having a different class and the same cluster
- f_{10} = number of pairs of objects having the same class and a different cluster
- f_{11} = number of pairs of objects having the same class and the same cluster

Rand Index

Rand Index =
$$\frac{f_{00} + f_{11}}{f_{00} + f_{01} + f_{10} + f_{11}}$$



Final Comment on Cluster Validity



"The validation of clustering structures is the most difficult and frustrating part of cluster analysis.

Without a strong effort in this direction, cluster analysis will remain a black art accessible only to those true believers who have experience and great courage."

Algorithms for Clustering Data, Jain and Dubes

