Week 9 Clustering

Seokho Chi Associate Professor I Ph.D. SNU Construction Innovation Lab

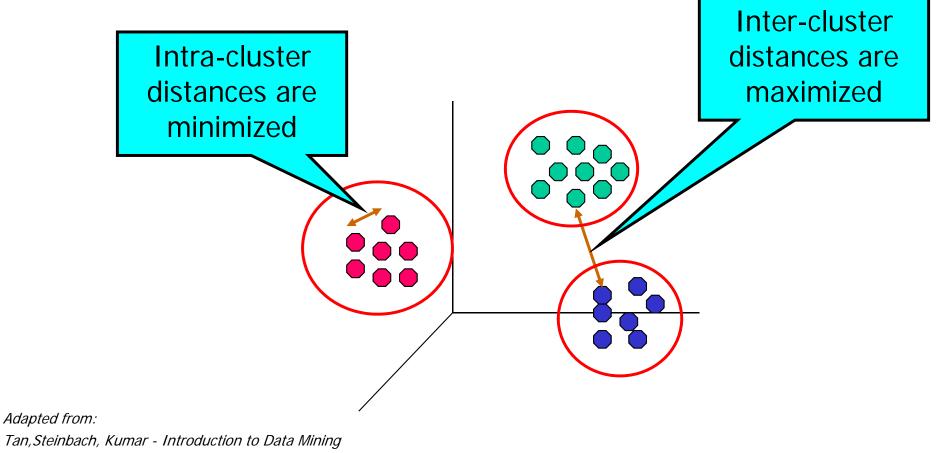


Source: Tan, Kumar, Steinback (2006)



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



Han, Kamber - Data Mining: Concepts and Techniques

General Applications of Clustering

- Pattern Recognition
- Spatial Data Analysis
 - create thematic maps in GIS by clustering feature spaces
 - detect spatial clusters and explain them in spatial data mining
- Image Processing
- Economic Science (especially market research)
- - Cluster Weblog data to discover groups of similar access patterns

Adapted from:

Examples of Clustering Applications

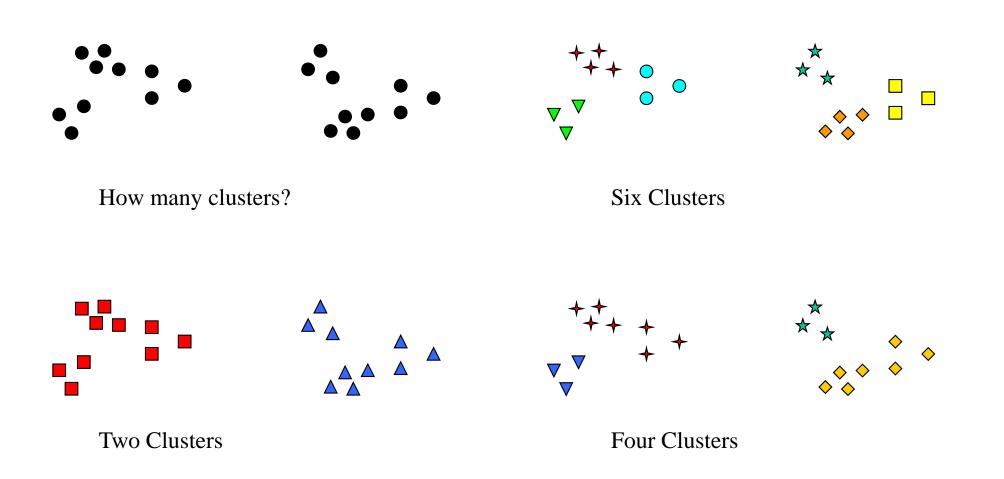
- <u>Marketing</u>: Help marketers discover distinct groups in their customer bases, and then use this knowledge to develop targeted marketing programs
- Land use: Identification of areas of similar land use in an earth observation database
- <u>Insurance</u>: Identifying groups of motor insurance policy holders with a high average claim cost
- <u>City-planning</u>: Identifying groups of houses according to their house type, value, and geographical location
- <u>Earthquake studies</u>: Observed earthquake epicenters should be clustered along continent faults

Adapted from:

What Is Good Clustering?

- A <u>good clustering</u> method will produce high quality clusters with
 - high <u>intra-class</u> similarity
 - low <u>inter-class</u> similarity
- The <u>quality</u> of a clustering result depends on both the similarity measure used by the method and its implementation.
- The <u>quality</u> of a clustering method is also measured by its ability to discover some or all of the <u>hidden</u> patterns.

Notion of a Cluster can be Ambiguous



Major Clustering Approaches

- <u>Partitioning algorithms</u>: A division data objects into nonoverlapping subsets (clusters) such that each data object is in exactly one subset
- <u>Hierarchy algorithms</u>: A set of nested clusters organized as a hierarchical tree
- Density-based: based on connectivity and density functions
- <u>Grid-based</u>: based on a multiple-level granularity structure
- <u>Model-based</u>: A model is hypothesized for each of the clusters and the idea is to find the best fit of that model to each other

Characteristics of the Input Data Are Important

- Type of distribution
 - Type of proximity or density measure
 - Sparseness
- Type of data
 - Attribute type
 - Dimensionality
- Noise and outliers

K-means Clustering

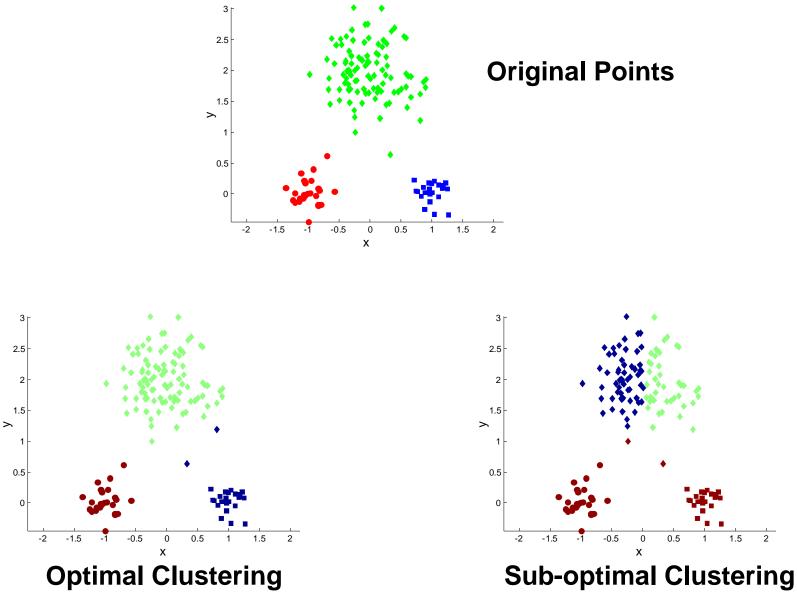
- Partitional clustering approach
- Each cluster is associated with a centroid (center point)
- Each point is assigned to the cluster with the closest centroid
- Number of clusters, K, must be specified
- The basic algorithm is very simple
 - 1: Select K points as the initial centroids.
 - 2: repeat
 - 3: 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

Adapted from:

K-means Clustering – Details

- Initial centroids are often chosen randomly.
 - Clusters produced may 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.

Two different K-means Clusterings

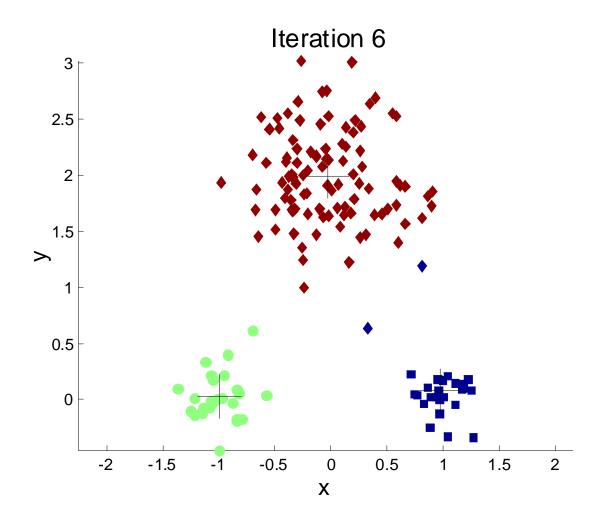


1.5

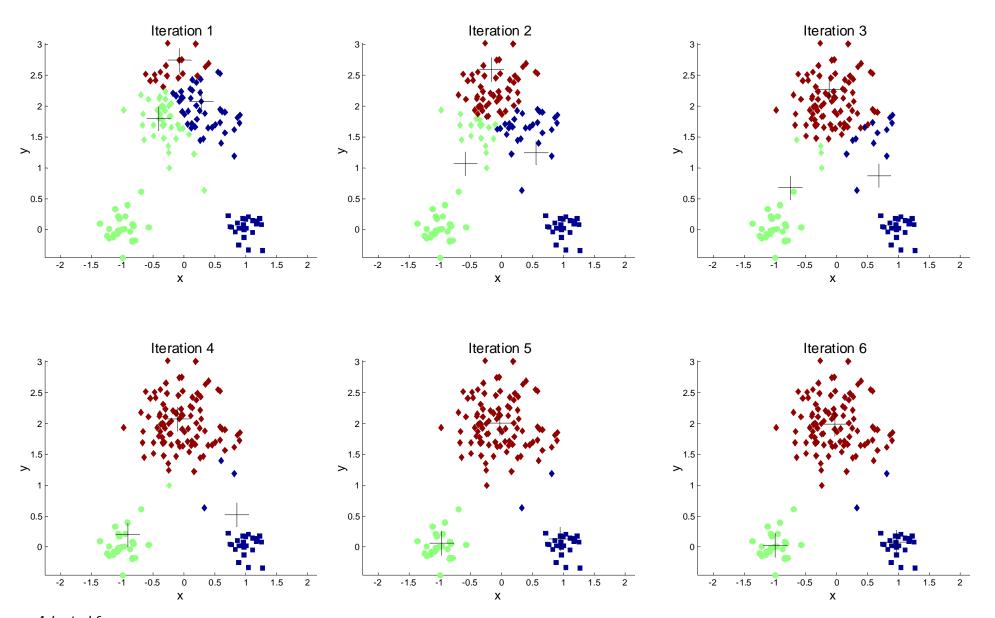
2

Adapted from:

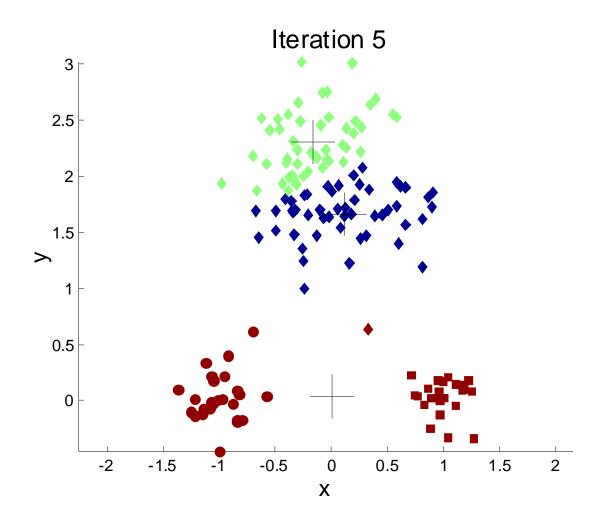
Importance of Choosing Initial Centroids (A)



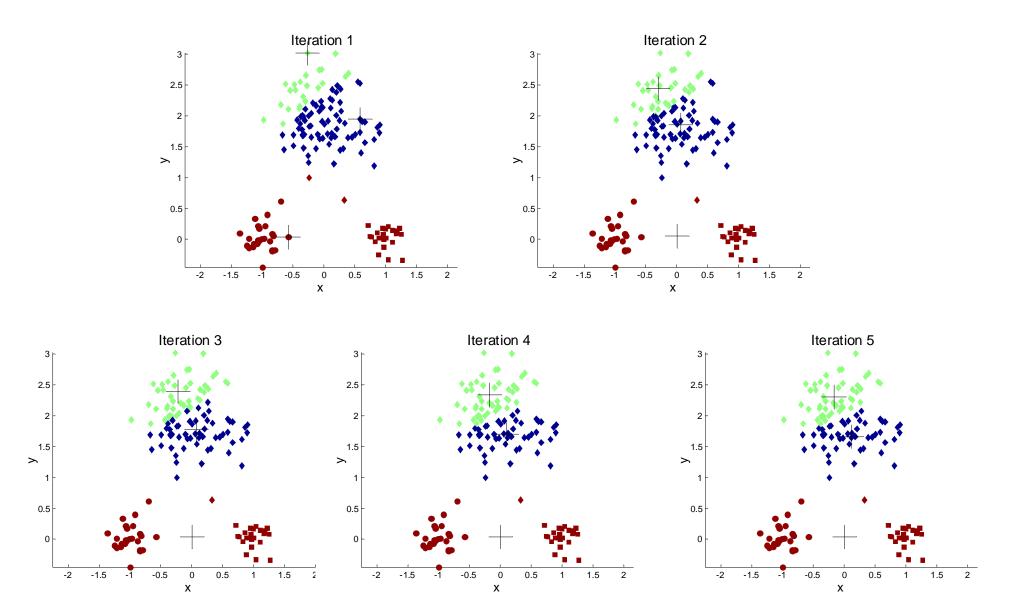
Importance of Choosing Initial Centroids (A)



Importance of Choosing Initial Centroids (B)



Importance of Choosing Initial Centroids (B)



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.

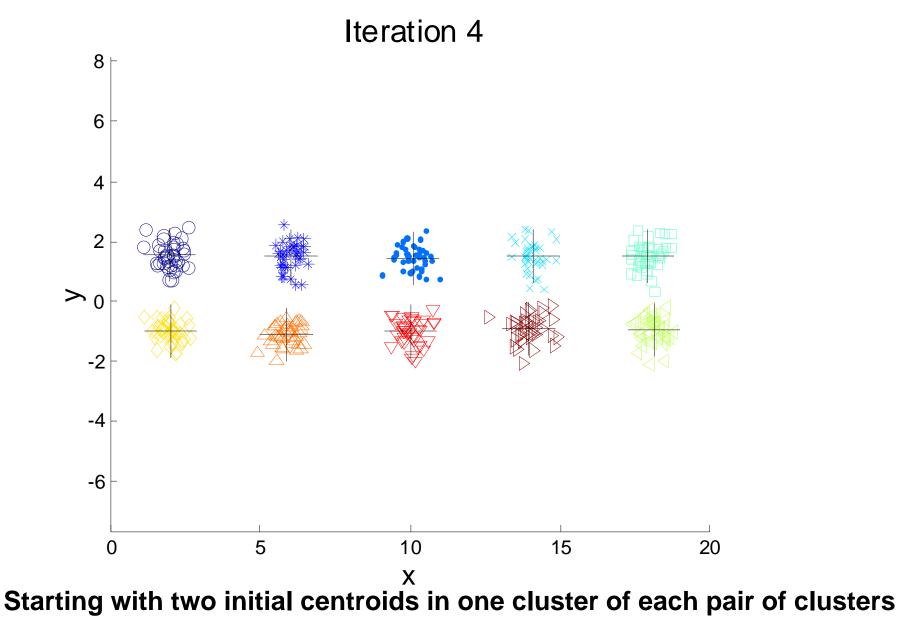
$$SSE = \sum_{i=1}^{K} \sum_{x \in C_i} dist^{-2}(m_i, x)$$

- *x* is a data point in cluster C_i and m_i is the representative (center) point for cluster C_i
- 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

Problems with Selecting Initial Points

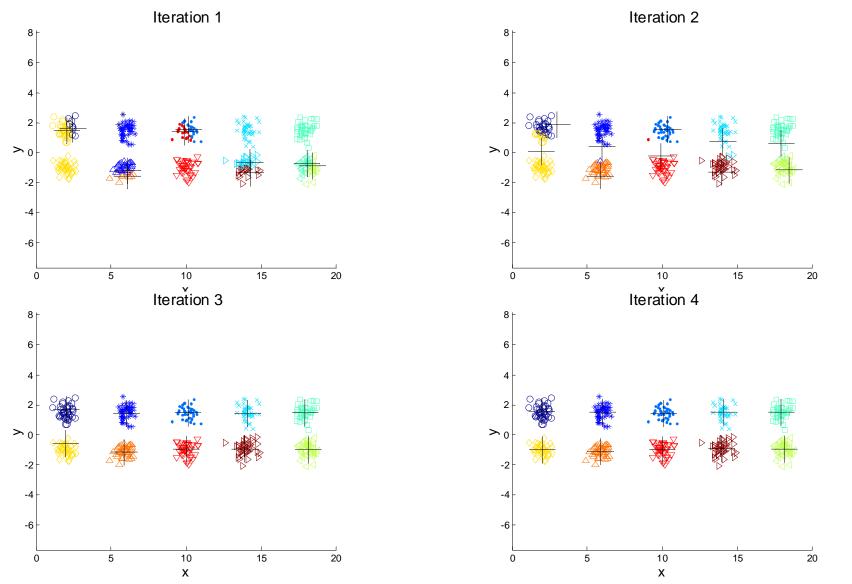
- If there are K 'real' clusters then the chance of selecting one centroid from each cluster is small.
 - Especially when K is large
 - Sometimes the initial centroids will readjust themselves in 'right' way, and sometimes they don't
 - Consider an example of five pairs of clusters

10 Clusters Example (A)



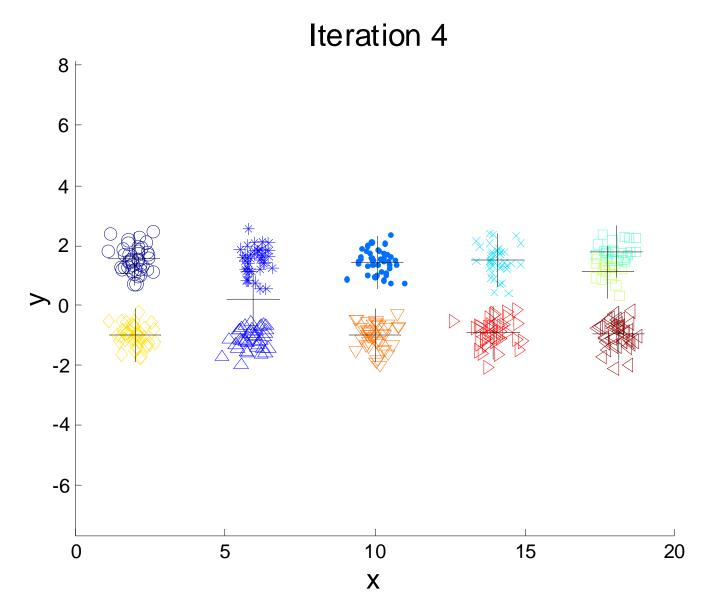
Adapted from:

10 Clusters Example (A)



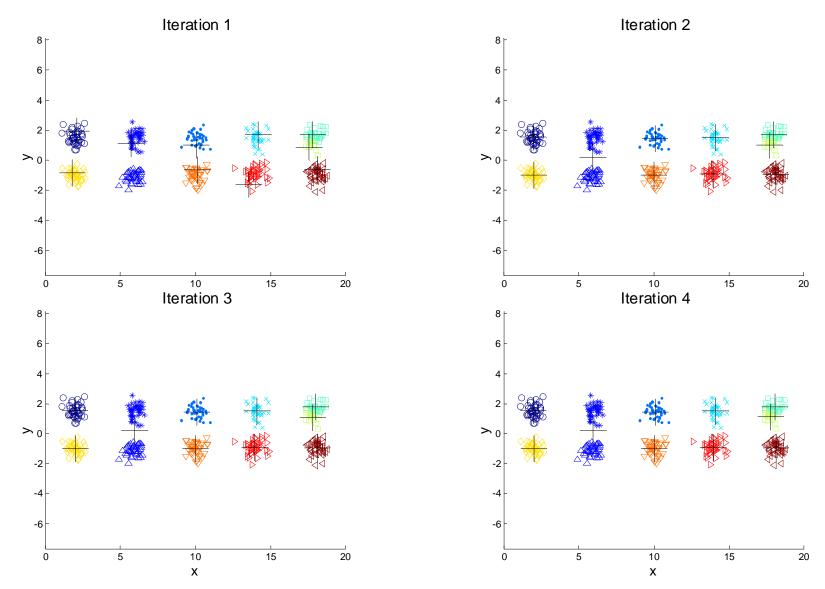
Starting with two initial centroids in one cluster of each pair of clusters

10 Clusters Example (B)



Starting with some pairs of clusters having three initial centroids, while other have only one or two. *Adapted from:*

10 Clusters Example (B)



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

Adapted from:

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
- Post-processing
 - Merge similar clusters after the processing
- Bisecting K-means
 - Start with one cluster, and then continuously split
 - Not as susceptible to initialization issues

Adapted from:

Pre-processing and Post-processing

- Pre-processing
 - Normalize the data (prevent from being dominated by big attribute values)
 - 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

Adapted from:

Bisecting K-means

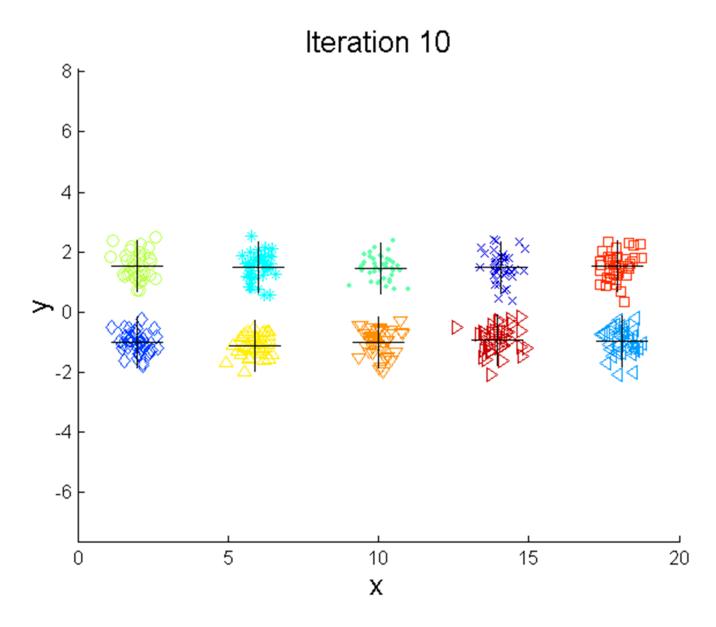
- Bisecting K-means algorithm
 - Start with one cluster and split continuously
 - No need to determine the number of clusters
 - Variant of K-means that can produce a partitional or a hierarchical clustering
- 1: Initialize the list of clusters to contain the cluster containing all points.

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

Adapted from:

Bisecting K-means Example

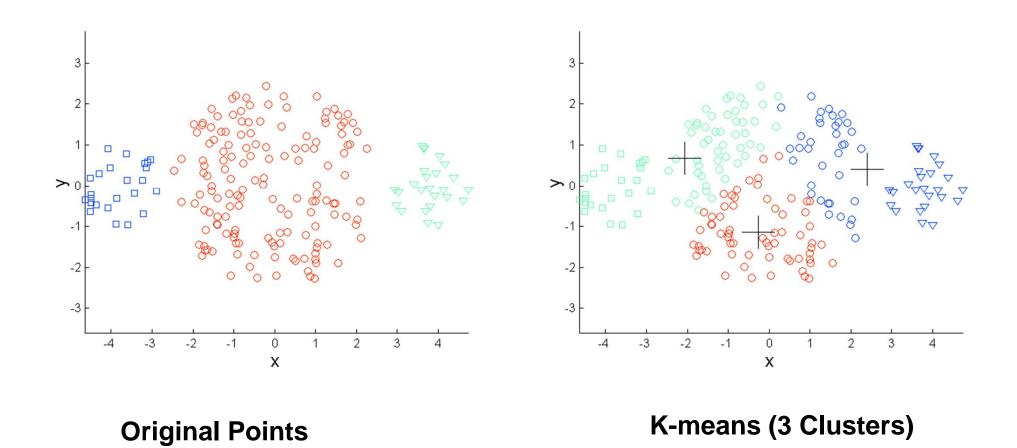


Adapted from:

Limitations of K-means

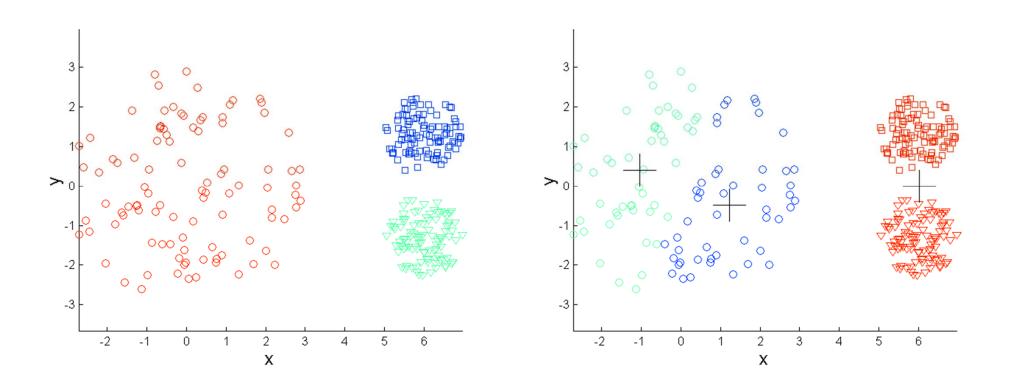
- K-means has problems when clusters are of differing
 - Sizes
 - Densities
 - Non-globular shapes
- K-means has problems when the data contains outliers.

Limitations of K-means: Differing Sizes



Adapted from:

Limitations of K-means: Differing Density

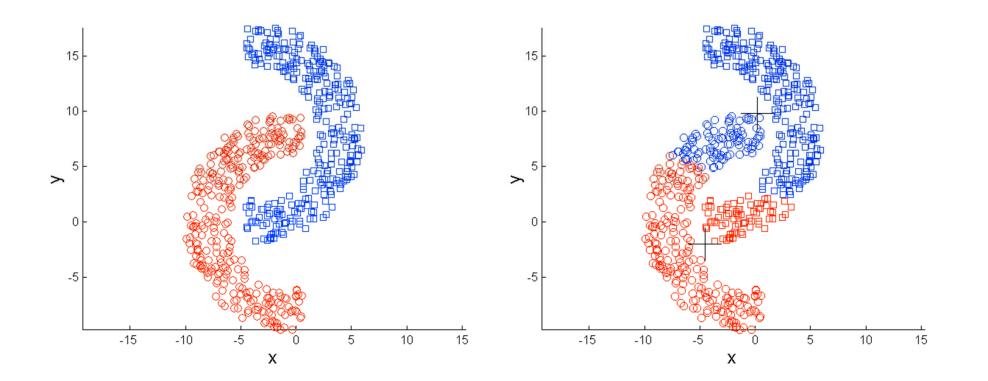


Original Points

K-means (3 Clusters)

Adapted from:

Limitations of K-means: Non-globular Shapes

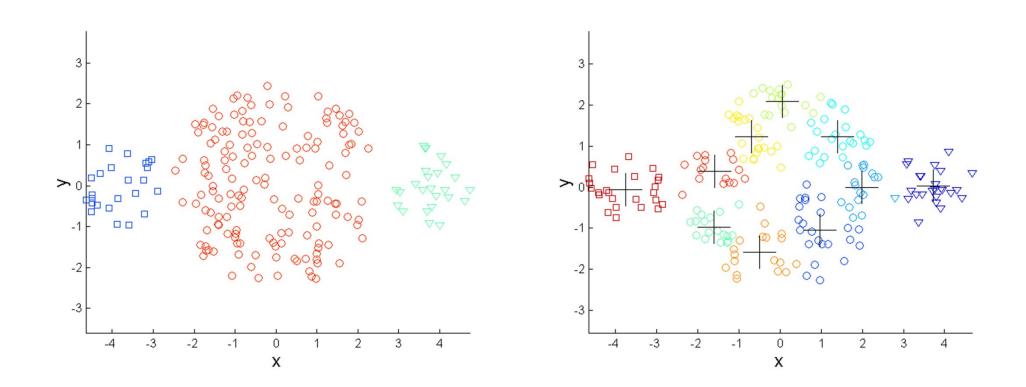


Original Points

K-means (2 Clusters)

Adapted from:

Overcoming K-means Limitations



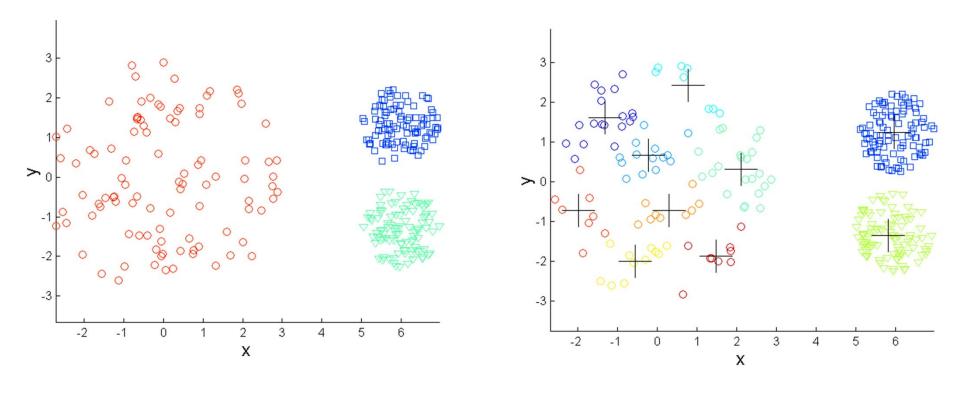
Original Points

K-means Clusters

One solution is to use many clusters. Find parts of clusters, but need to put together.

Adapted from:

Overcoming K-means Limitations

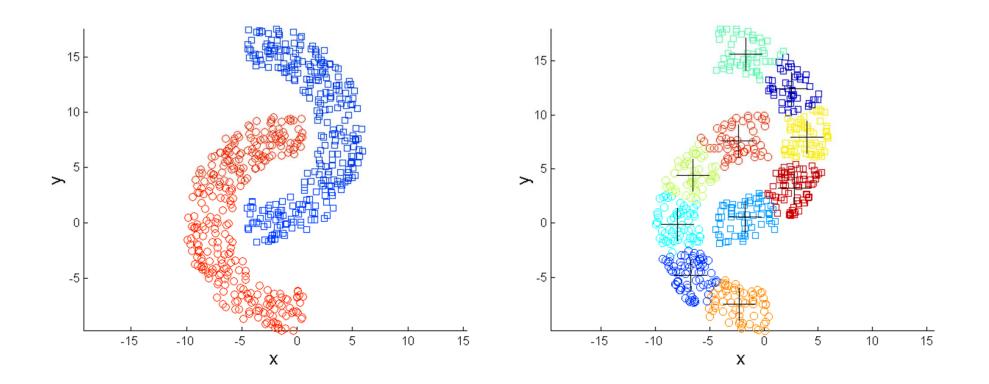


Original Points

K-means Clusters

Adapted from:

Overcoming K-means Limitations

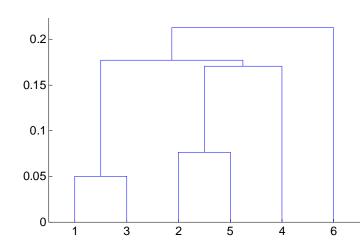


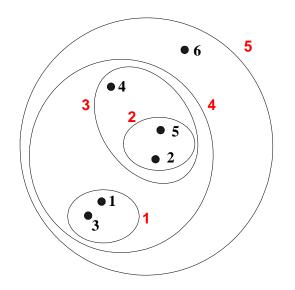
Original Points

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

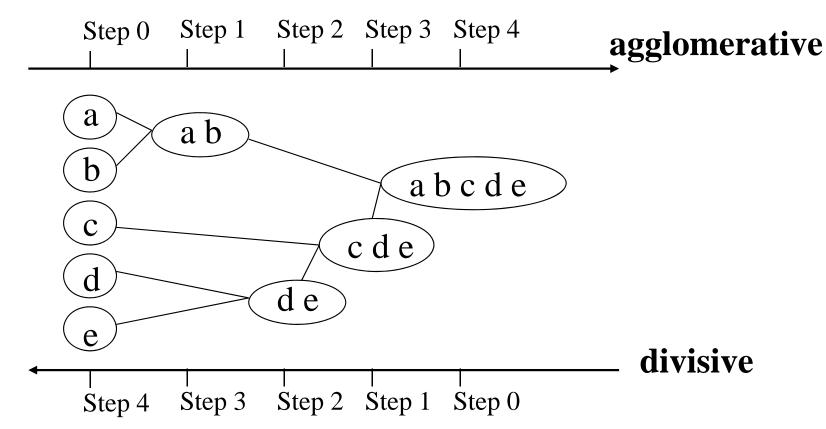
- Do not have to assume any particular number of clusters
 - Any desired number of clusters can be obtained by 'cutting' the dendogram at the proper level

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

Hierarchical Clustering

 This method does not require the number of clusters k as an input, but needs a termination condition



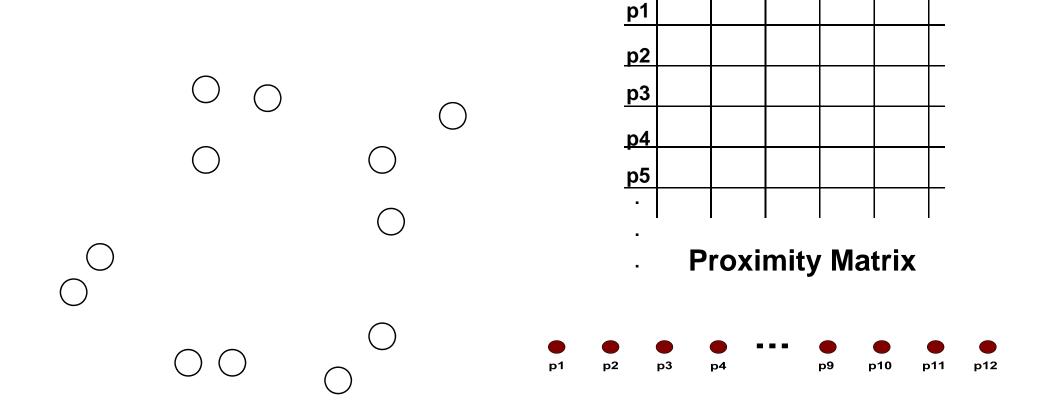
Adapted from:

Agglomerative Clustering Algorithm

- More 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

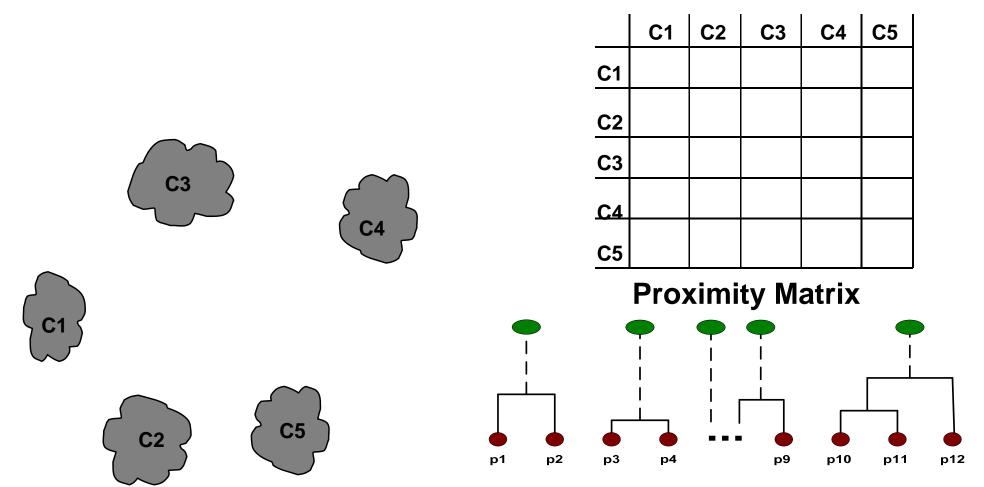
 Start with clusters of individual points and a proximity matrix
 p1 | p2 | p3 | p4 | p5



Adapted from:

Intermediate Situation

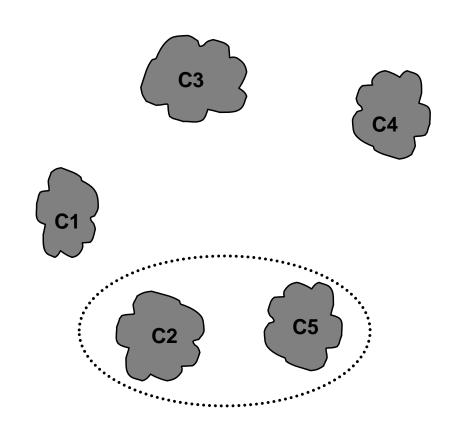
• After some merging steps, we have some clusters

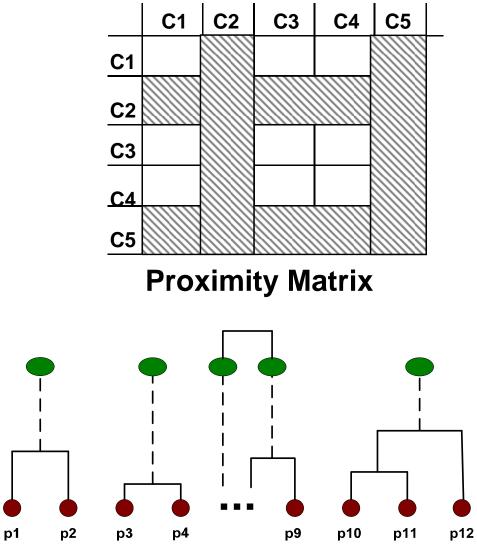


Adapted from:

Intermediate Situation

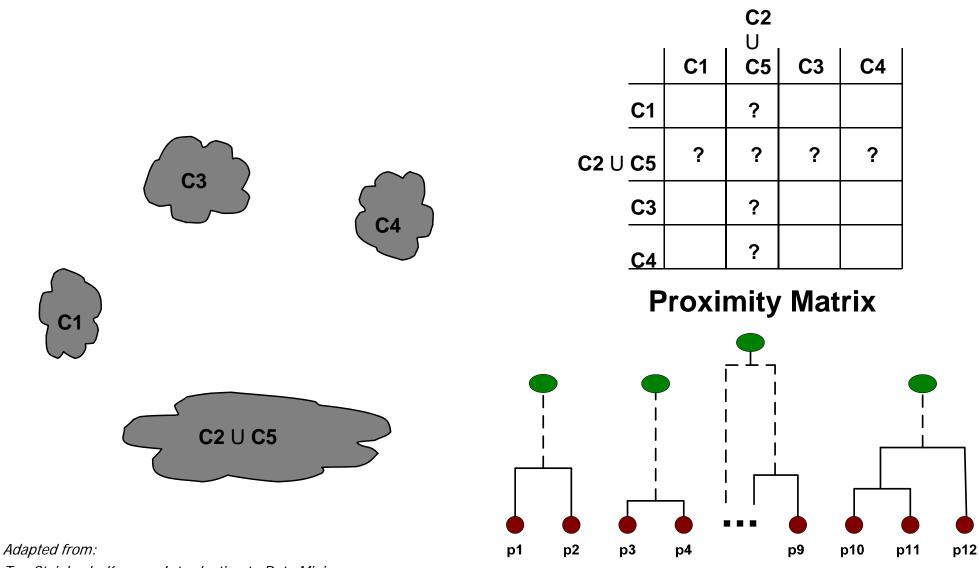
 We want to merge the two closest clusters (C2 and C5) and update the proximity matrix.

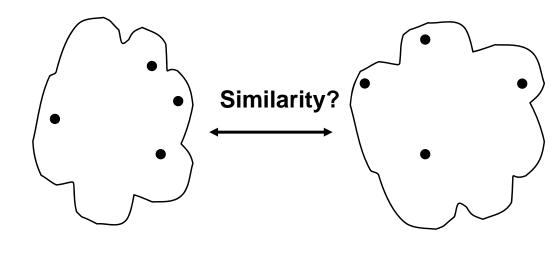


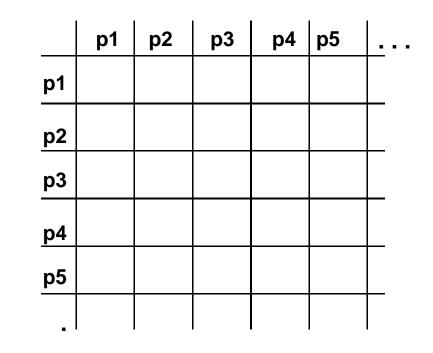


After Merging

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





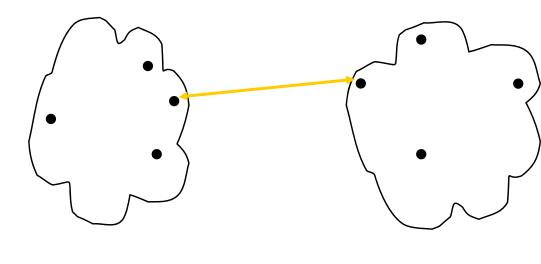


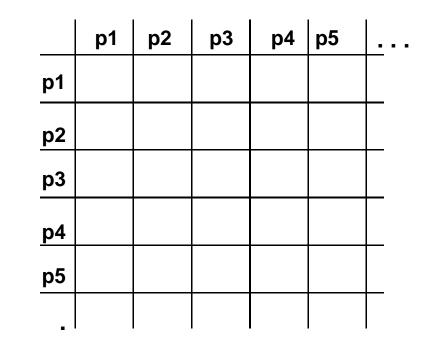
- MIN
- MAX
- Group Average

Proximity Matrix

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

Adapted from:

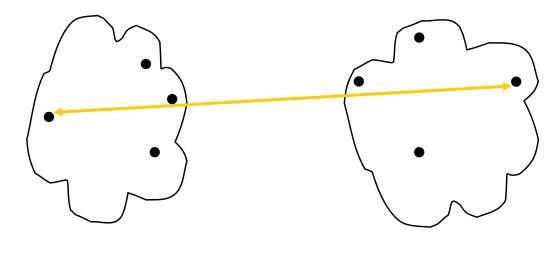


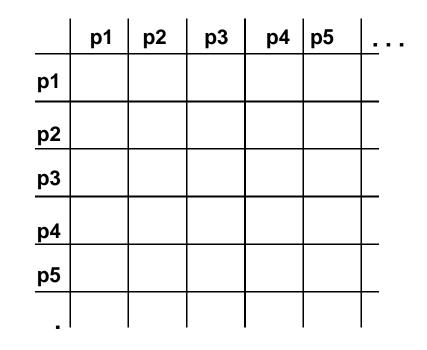


- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
 - Ward's Method uses squared error

Adapted from:

Tan, Steinbach, Kumar - Introduction to Data Mining Han, Kamber - Data Mining: Concepts and Techniques

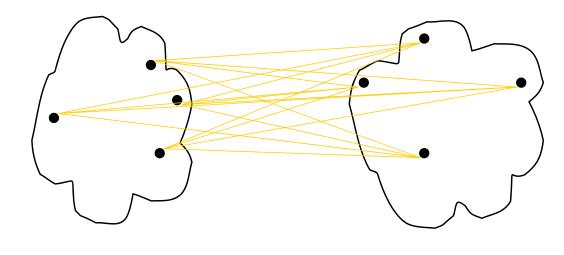


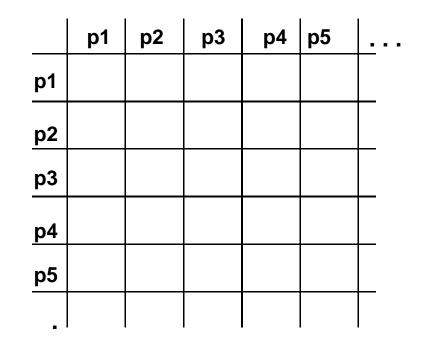


- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
 - Ward's Method uses squared error

Adapted from:

Tan, Steinbach, Kumar - Introduction to Data Mining Han, Kamber - Data Mining: Concepts and Techniques

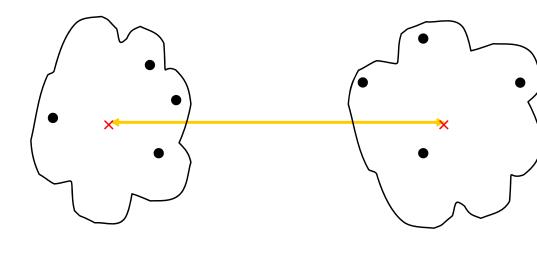


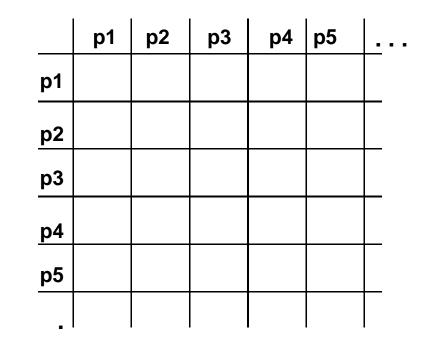


- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
 - Ward's Method uses squared error

Adapted from:

Tan, Steinbach, Kumar - Introduction to Data Mining Han, Kamber - Data Mining: Concepts and Techniques





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

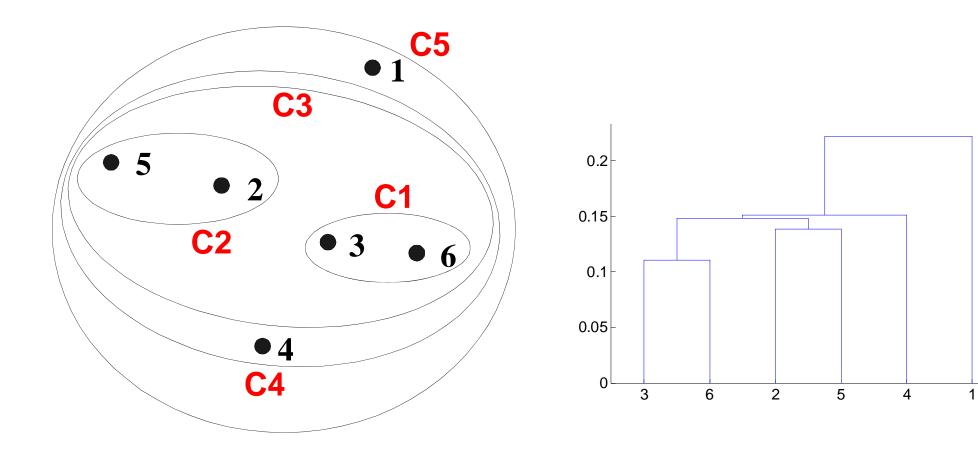
Adapted from:

Tan, Steinbach, Kumar - Introduction to Data Mining Han, Kamber - Data Mining: Concepts and Techniques

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.

Hierarchical Clustering: MIN

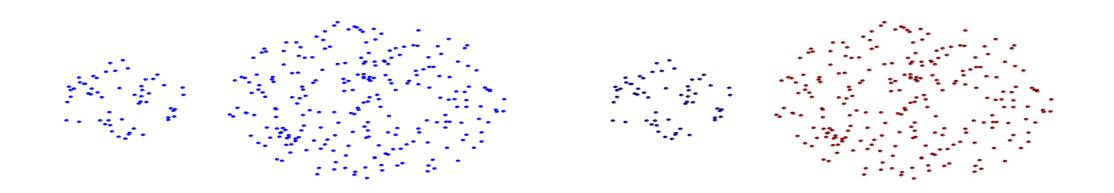


Nested Clusters

Adapted from:

Tan,Steinbach, Kumar - Introduction to Data Mining Han, Kamber - Data Mining: Concepts and Techniques Dendrogram

Strength of MIN

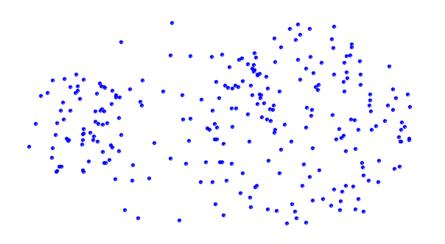


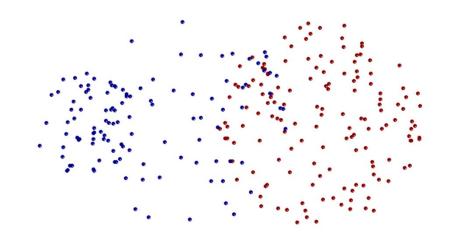
Original Points

Two Clusters

• Can handle non-elliptical shapes

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

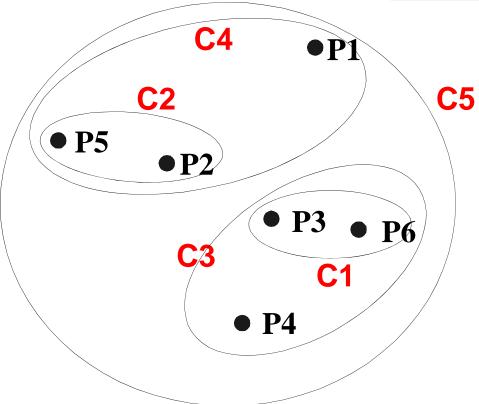
least similar (most distant) points in the different

clusters

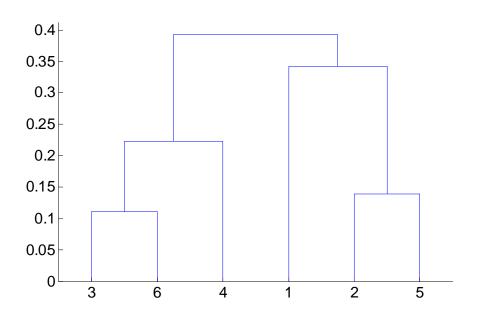
– Determined by all pairs of points in the two clusters

Hierarchical Clustering: MAX

	C1 or C2	P1	P4
C1(P3, P6)	P5 & P6	P1 & P6	P4 & P6
C2(P2, P5)	P5 & P6	P1 & P5	P4 & P5



*MAX \rightarrow then, FIND MIN among MAX



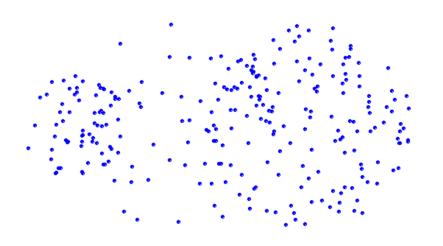
Nested Clusters

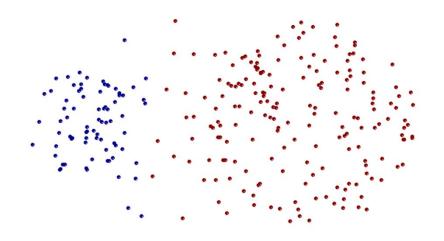
Adapted from:

Tan, Steinbach, Kumar - Introduction to Data Mining Han, Kamber - Data Mining: Concepts and Techniques

Dendrogram

Strength of MAX



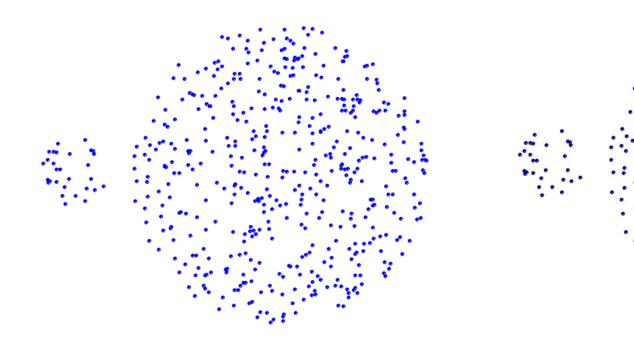


Original Points

Two Clusters

• Less susceptible to noise and outliers

Limitations of MAX



Original Points

Two Clusters

•Tends to break large clusters

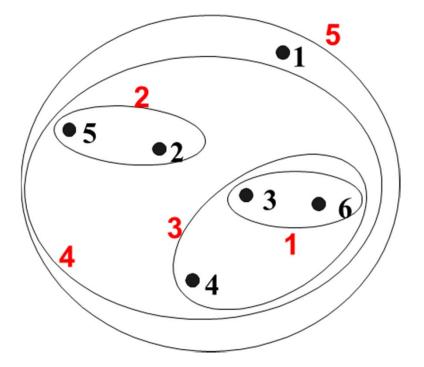
•Biased towards globular clusters

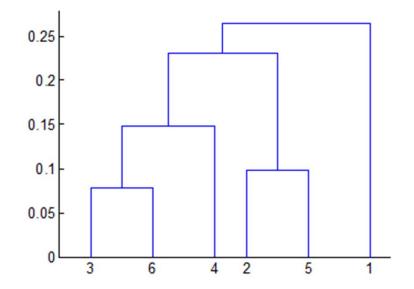
Cluster Similarity: Group Average

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

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

Hierarchical Clustering: Group Average





Nested Clusters

Dendrogram

Adapted from:

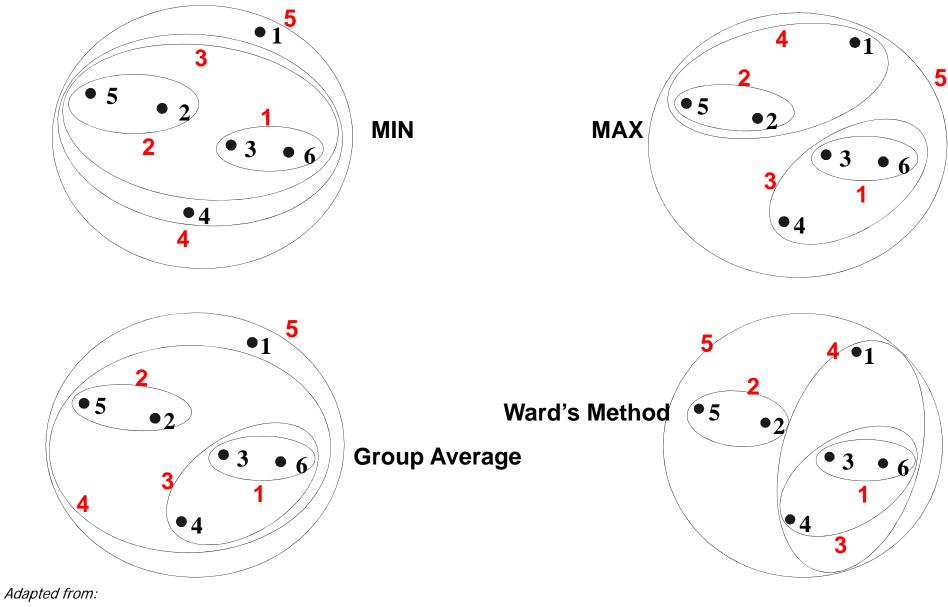
Hierarchical Clustering: Group Average

- 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: Limitations

- Once a decision is made to combine two clusters, it cannot be undone
- Sensitivity to noise and outliers
- Difficulty handling different sized clusters and convex shapes
- Large computation time and storage (compute all the matrix first)
- No objective function: change depends on different measurement scheme

Density-Based Clustering Methods

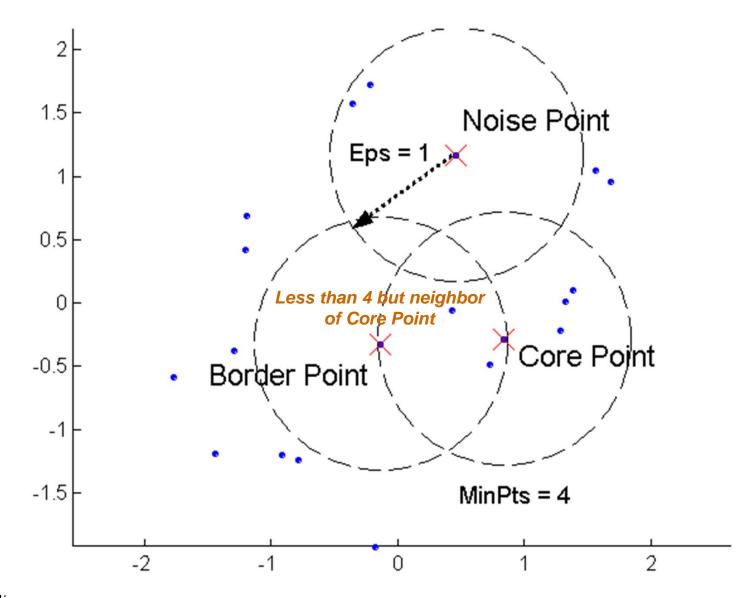
- Clustering based on density (local cluster criterion), such as density-connected points
- Major features
 - Discover clusters of arbitrary shape
 - Handle noise
 - One scan
 - Need density parameters as termination condition
- Several interesting studies
 - DBSCAN: Ester, et al. (KDD'96)
 - OPTICS: Ankerst, et al (SIGMOD'99).
 - <u>DENCLUE</u>: Hinneburg & D. Keim (KDD'98)
 - <u>CLIQUE</u>: Agrawal, et al. (SIGMOD'98)

Adapted from:

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 the 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.

Core, Border, and Noise Points

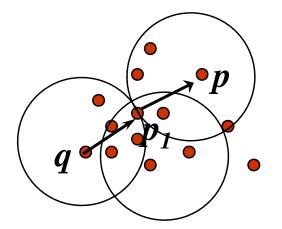


Adapted from:

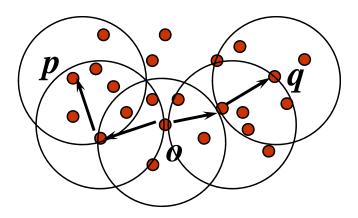
Density-Based Clustering Concepts

Density-reachable:

- A point *p* is density-reachable from a point *q* with respect to *Eps*, *MinPts* if there is a chain of points $p_1, ..., p_n, p_1 =$ $q, p_n = p$ such that p_{i+1} is directly density-reachable from p_i



- Density-connected
 - A point *p* is density-connected to a point *q* with *Eps*, *MinPts* if there is a point *o* such that both, *p* and *q* are density-reachable from *o* with respect to *Eps* and *MinPts*.



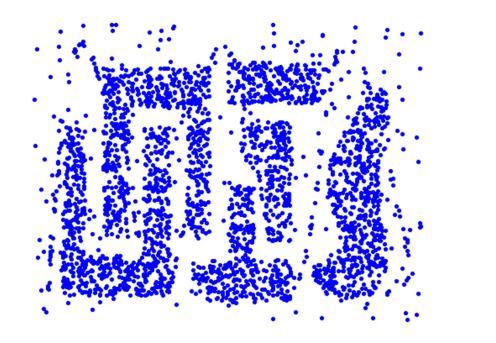
Adapted from:

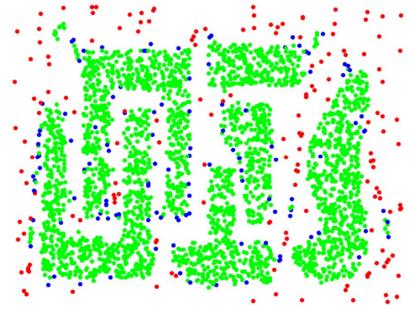
DBSCAN Algorithm

- Arbitrary select a point *p*
- Retrieve all points density-reachable from *p* with respect to *Eps* and *MinPts*.
- If **p** is a core point, a cluster is formed.
- If *p* is a border point, no points are density-reachable from *p* and DBSCAN visits the next point of the database.
- Continue the process until all of the points have been processed.
 1. Label all points as core, border, or noise
 - 2. Eliminate noise points
 - 3. Put an edge
 - 4. Make each group of connected cores
 - 5. Border points

Adapted from:

DBSCAN: Core, Border and Noise Points





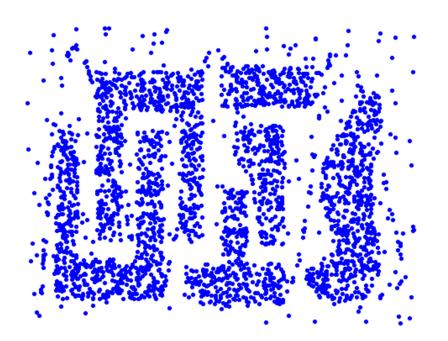
Original Points

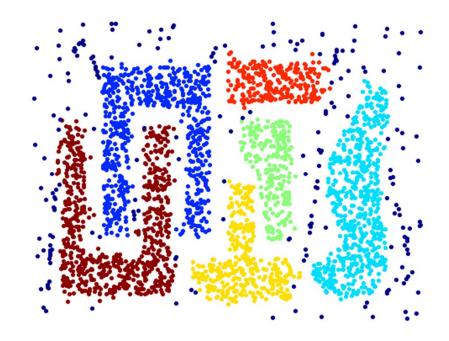
Point types: core, border and noise

Eps = 10, MinPts = 4

Adapted from:

When DBSCAN Works Well





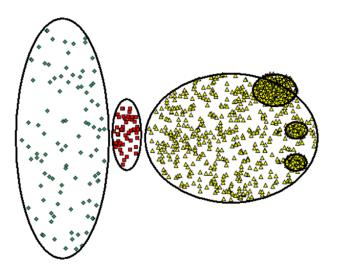
Original Points

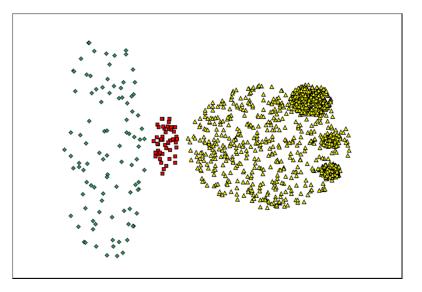


• Resistant to Noise

• Can handle clusters of different shapes and sizes

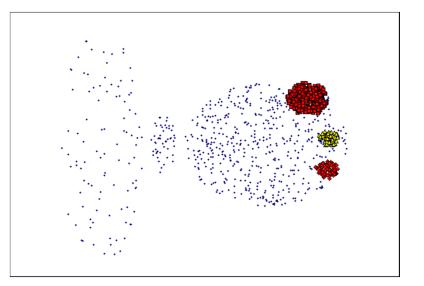
When DBSCAN Does NOT Work Well





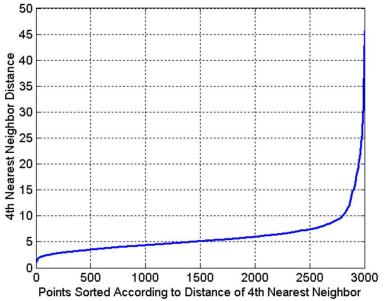
Original Points

- Varying densities
- High-dimensional data
- Eps = ?, MinPts = ?



DBSCAN: Determining Eps and MinPts

- Idea: Look at the behavior of the distance from a point to its kth nearest neighbor (k-dist)
- For points that belong to some cluster, the value of k-dist will be small since they have many neighboring points.
- For points that are not in a cluster, such as noise points, the k-dist will be relatively large.
- Thus, compute the k-dist for all the data points → sort them in increasing order → plot the sorted value → find a sharp change for Eps
 - \rightarrow take the value of k as MinPts



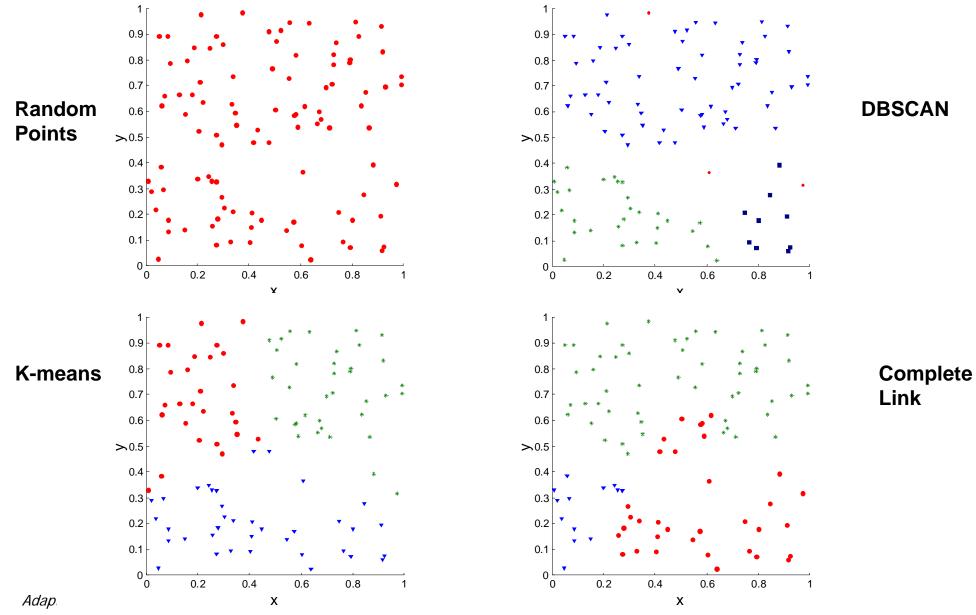
Adapted from:

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

Adapted from:

Clusters found in Random Data



Different Aspects of Cluster Validation

- Determining the clustering tendency of a set of data, i.e., distinguishing whether non-random 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.
- 3. 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.

Adapted from:

Measures of Cluster Validity

- Numerical measures that are applied to judge various aspects of cluster validity, are classified into the following three types.
 - External Index: Used to measure the extent to which cluster labels match externally supplied class labels.
 e.g.: Entropy
 - Internal Index: Used to measure the goodness of a clustering structure *without* respect to external information.

– e.g.: 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

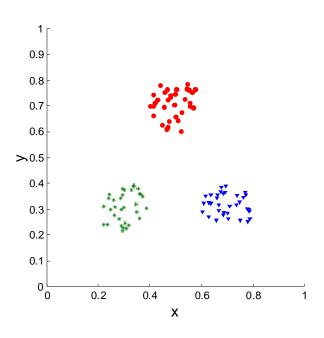
Adapted from:

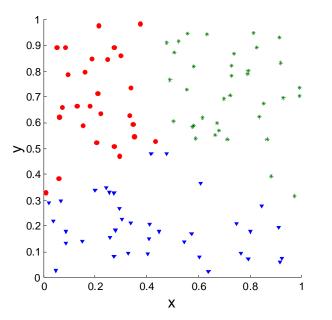
Measuring Cluster Validity Via Correlation

- Two matrices:
 - Proximity (similarity) Matrix
 - "Incidence" Matrix
 - 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
- Compute the correlation between the two matrices
- 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.

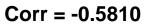
Measuring Cluster Validity Via Correlation

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



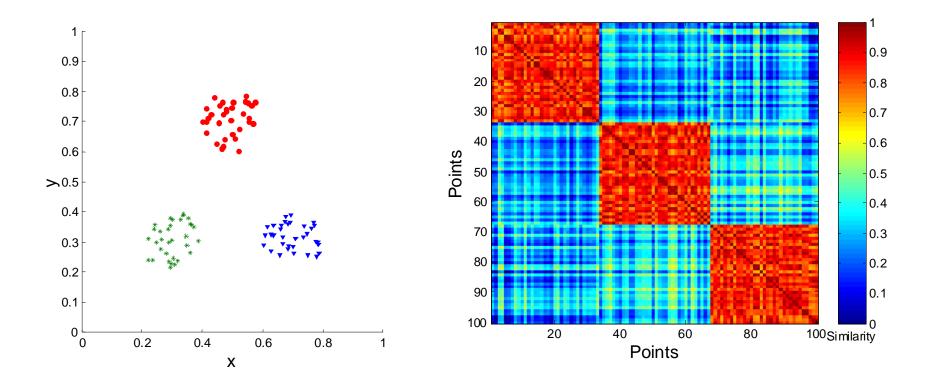


Corr = -0.9235



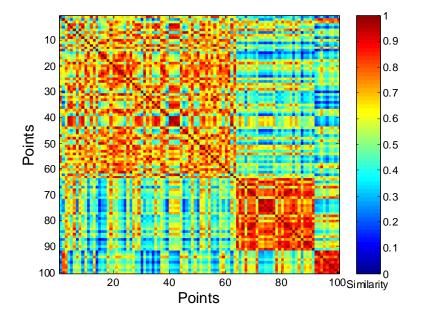
Adapted from:

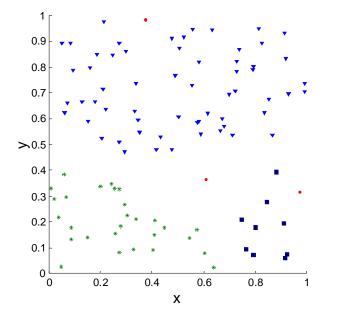
 Order the similarity matrix with respect to cluster labels and inspect visually.



Adapted from: Tan,Steinbach, Kumar - Introduction to Data Mining Han, Kamber - Data Mining: Concepts and Techniques

Clusters in random data are not so crisp

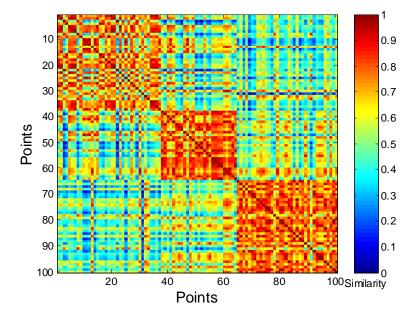


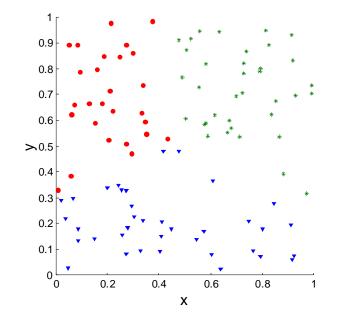


DBSCAN

Adapted from:

Clusters in random data are not so crisp

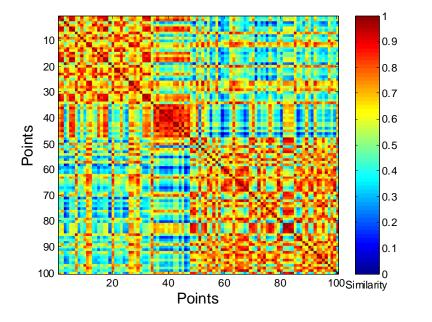


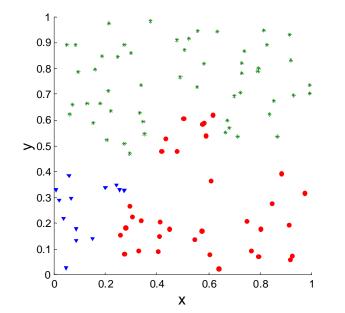


K-means

Adapted from:

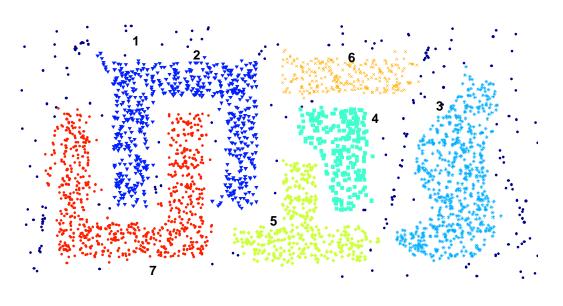
Clusters in random data are not so crisp

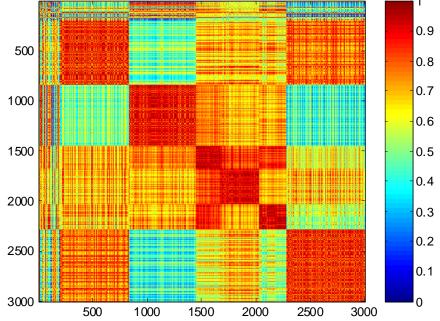




Complete Link

Adapted from:





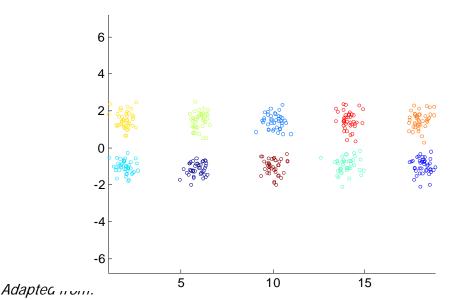
DBSCAN

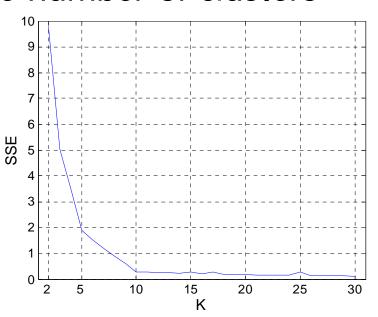
Internal Measures: SSE

- Clusters in more complicated figures aren't well separated
- Internal Index: Used to measure the goodness of a clustering structure without respect to external information

– SSE

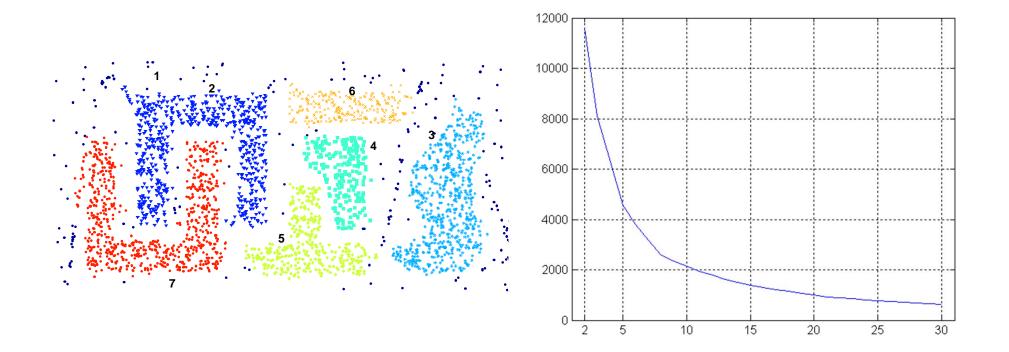
- SSE is good for comparing two clusterings or two clusters (average SSE).
- Can also be used to estimate the number of clusters





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
 - Example: SSE
- Cluster Separation: Measure how distinct or wellseparated a cluster is from other clusters
- Example: Squared Error
 - Cohesion is measured by the within cluster sum of squares (SSE)

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

– Separation is measured by the between cluster sum of squares

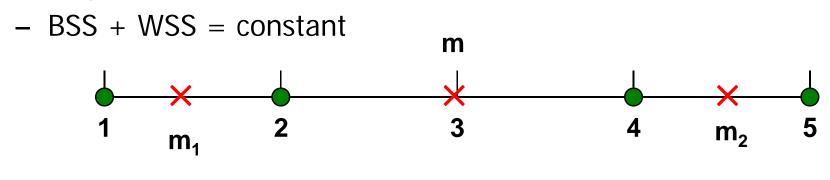
$$BSS = \sum_{i} |C_{i}| (m - m_{i})^{2}$$

where $|C_{i}|$ is the size of cluster is

Adapted from:

Internal Measures: Cohesion and Separation

• Example: SSE



K=1 cluster:

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

BSS = 4 × (3-3)^{2} = 0
Total = 10 + 0 = 10

K=2 clusters:

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

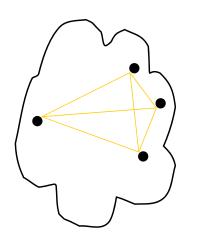
$$BSS = 2 \times (3 - 1.5)^{2} + 2 \times (4.5 - 3)^{2} = 9$$

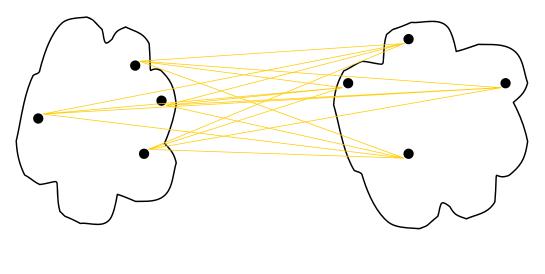
$$Total = 1 + 9 = 10$$

Adapted from:

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.





cohesion

separation

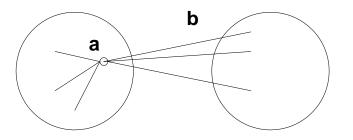
Adapted from:

Internal Measures: Silhouette Coefficient

- Silhouette Coefficient combine ideas of both cohesion and separation, but for individual points, as well as clusters
- 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 then given by

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

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



External Measures of Cluster Validity: Entropy and Purity

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

Table 5.9. K-means Clustering Results for LA Document Data Set

- 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 veighted 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 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 $purity = \sum_{i=1}^{K} \frac{m_i}{m} purity_j$.

Adapted from:

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

Grid-Based Clustering

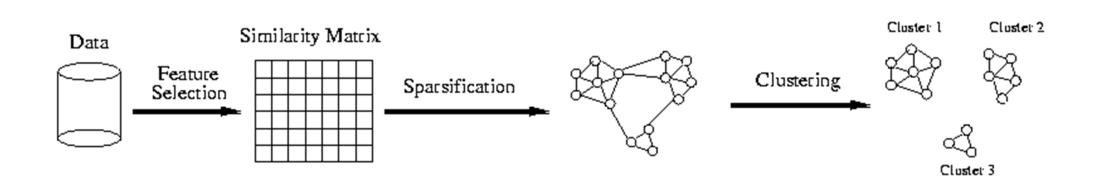
- Efficient way to organize a set of data, at least in low dimensions
- Algorithm:
 - 1. Define a set of grid cells
 - 2. Assign objects to the appropriate cells and compute the density of each cell
 - 3. Eliminate cells having a density below a specified threshold
 - 4. Form clusters from contiguous (adjacent groups of dense cells

Adapted from:

Graph-Based Clustering

- Graph-Based clustering uses the proximity graph
 - Start with the proximity matrix
 - Consider each point as a node in a graph
 - Each edge between two nodes has a weight which is the proximity between the two points
 - Initially the proximity graph is fully connected
- In the simplest case, clusters are connected components in the graph.

Sparsification in the Clustering Process



Graph-Based Clustering: Sparsification

- The amount of data that needs to be processed is drastically reduced
 - Setting many of low similarity values to 0
 - Break all links that have a similarity below a specified threshold or keep only links to the k nearest neighbors of point
 - Sparsification can eliminate more than 99% of the entries in a proximity matrix
 - The amount of time required to cluster the data is drastically reduced
 - The size of the problems that can be handled is increased

Adapted from:

Graph-Based Clustering: Sparsification ...

- Clustering may work better
 - Sparsification techniques keep the connections to the most similar (nearest) neighbors of a point while breaking the connections to less similar points.
 - The nearest neighbors of a point tend to belong to the same class as the point itself.
 - This reduces the impact of noise and outliers and sharpens the distinction between clusters.
- Sparsification facilitates the use of graph partitioning algorithms (or algorithms based on graph partitioning algorithms).
 - Chameleon and Hypergraph-based Clustering

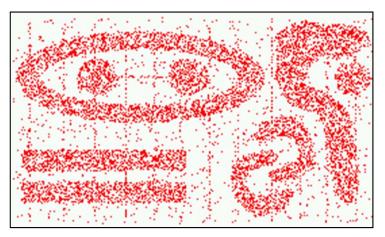
Adapted from:

Characteristics of Spatial Data Sets

- Clusters are defined as densely populated regions of the space
- Clusters have arbitrary shapes, orientation, and non-uniform sizes
- Difference in densities across clusters and variation in density within clusters
- Existence of special artifacts (*streaks*) and noise

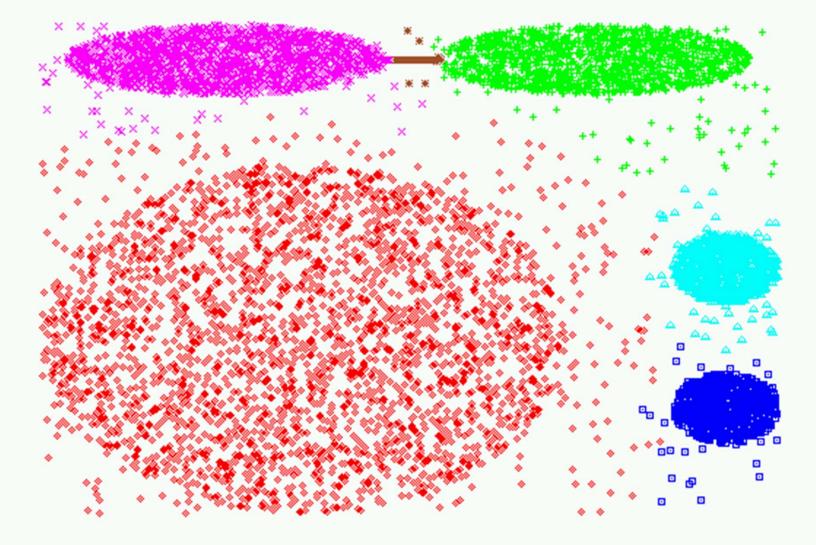
The clustering algorithm must address the above characteristics and also require minimal supervision.



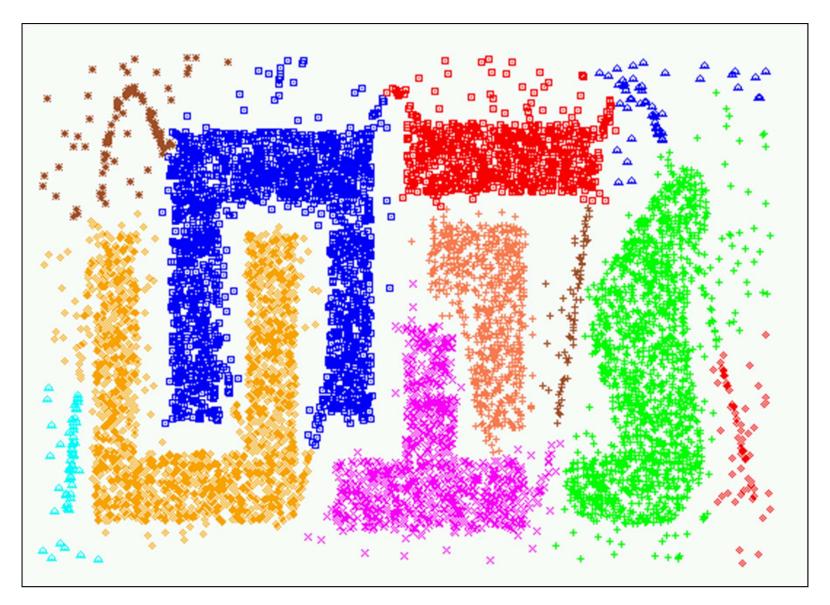


Experimental Results: CHAMELEON

*An agglomerative clustering algorithm with dynamic modeling using an efficient graph partitioning algorithm (considers density, shapes, closeness, interconnectivity, etc.)

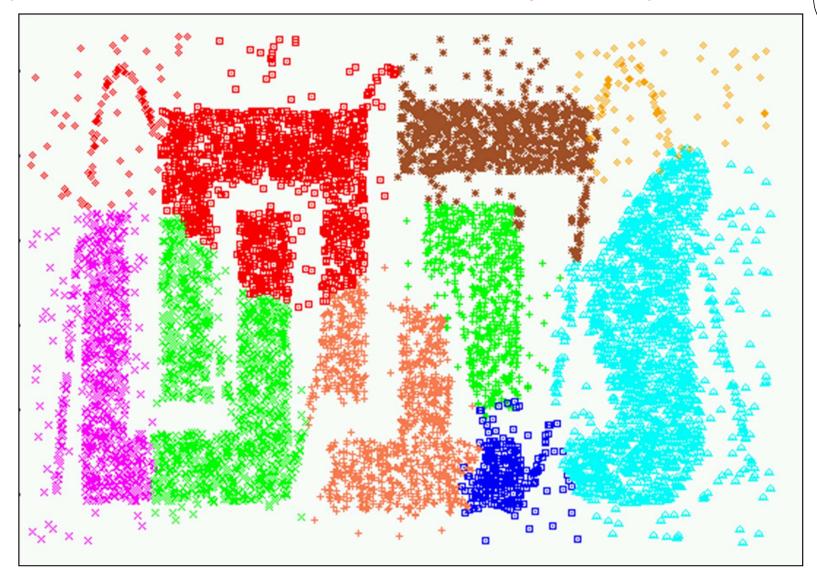


Experimental Results: CHAMELEON



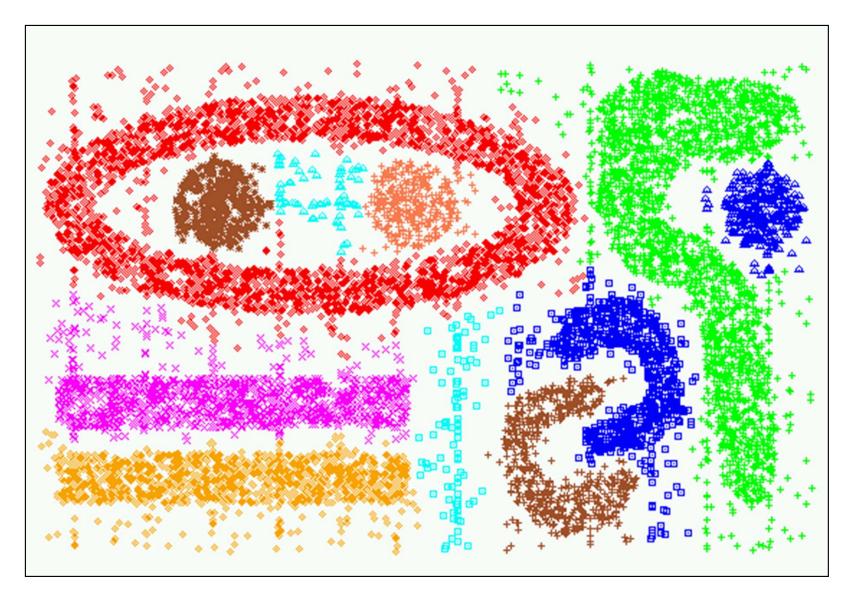
Experimental Results: CURE (10 clusters)

*Clustering Using REpresentatives: (1) Equally partitions, (2) uses a constant number of points to represent a cluster which capture the geometry and shape of the cluster and (3) then shrinks them toward the center of the cluster by a factor, alpha

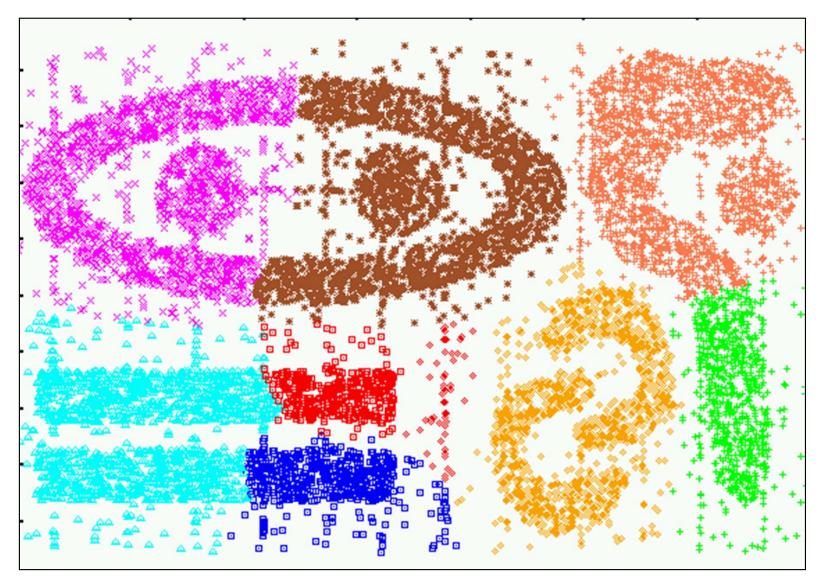


Adapted from:

Experimental Results: CHAMELEON



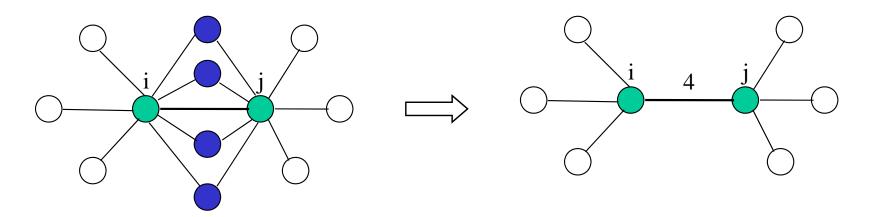
Experimental Results: CURE (9 clusters)



Shared Nearest Neighbor (SNN) Similarity

If two points are similar to many of the same points, then they are similar to one another, even if a direct measurement of similarity does not indicate this.

SNN graph: the weight of an edge is the number of shared neighbors between vertices given that the vertices are connected



SNN Clustering Algorithm

1. Compute the similarity matrix

This corresponds to a similarity graph with data points for nodes and edges whose weights are the similarities between data points

2. Sparsify the similarity matrix by keeping only the *k* most similar neighbors

This corresponds to only keeping the *k* strongest links of the similarity graph

3. Construct the shared nearest neighbor graph from the sparsified similarity matrix.

At this point, we could apply a similarity threshold and find the connected components to obtain the clusters (Jarvis-Patrick algorithm)

4. Find the SNN density of each Point.

Using a user specified parameters, *Eps*, find the number points that have an SNN similarity of *Eps* or greater to each point. This is the SNN density of the point

SNN Clustering Algorithm ...

5. Find the core points

Using a user specified parameter, *MinPts*, find the core points, i.e., all points that have an SNN density greater than *MinPts*

6. Form clusters from the core points

If two core points are within a radius, *Eps*, of each other they are place in the same cluster

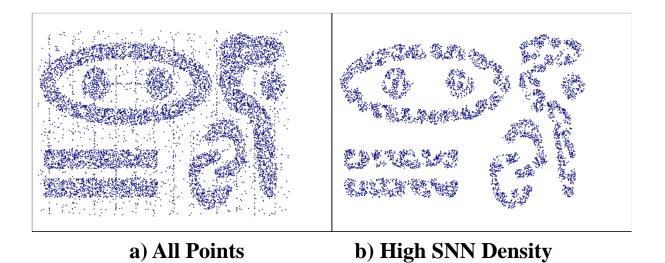
7. Discard all noise points

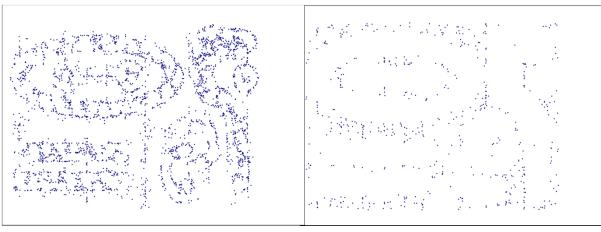
All non-core points that are not within a radius of *Eps* of a core point are discarded

8. Assign all non-noise, non-core points to clusters This can be done by assigning such points to the nearest core point

(Note that steps 4-8 are DBSCAN)

SNN Density



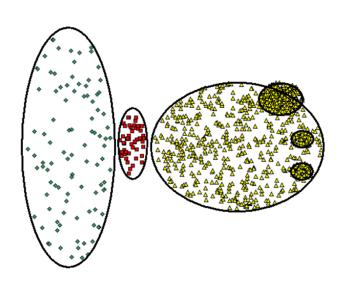


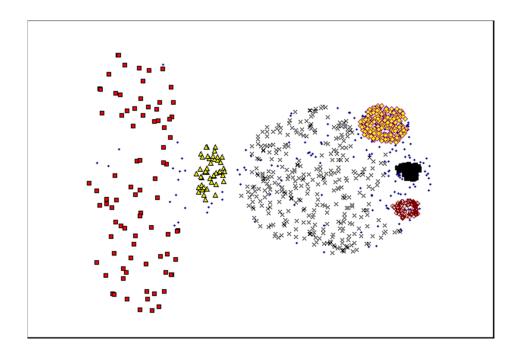
c) Medium SNN Density

d) Low SNN Density

Adapted from:

SNN Clustering Can Handle Differing Densities

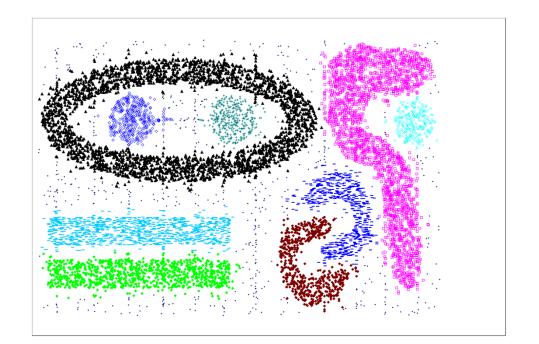


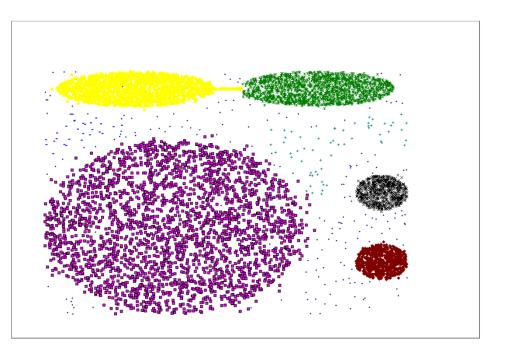


Original Points

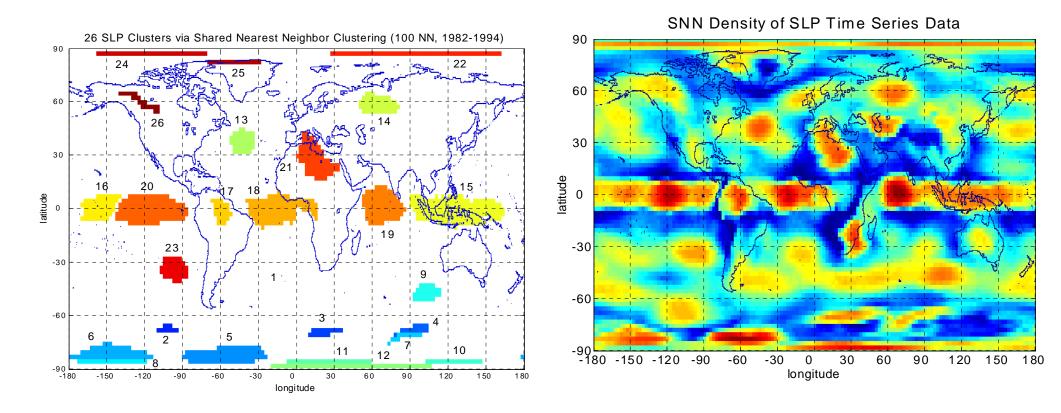
SNN Clustering

SNN Clustering Can Handle Other Difficult Situations





Finding Clusters of Time Series In Spatio-Temporal Data



SNN Clusters of Pressure

SNN Density of Pressure