

Clustering

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What is Clustering Analysis?

- **Aka binning/segmentation/hashing**
- **Finding similarities between data according to the characteristics found in the data and grouping similar data objects into clusters**
 - Number of clusters is not known ahead of time
- **Cluster: A collection of data objects**
 - similar (or related) to one another within the same group
 - dissimilar (or unrelated) to the objects in other groups
- **A type of Unsupervised Learning: no predefined classes**

Clustering Applications

- **Typical applications**
 - As a **stand-alone tool** to get insight into data distribution
 - As a **preprocessing step** for other algorithms
- **Biology:**
 - Taxonomy of living things: kingdom, phylum, class, order, family, genus and species
- **Information retrieval:**
 - Document clustering
- **Land use:**
 - Identification of areas of similar land use in an earth observation database

Clustering Applications

- **Marketing:**
 - Help marketers discover distinct groups in their customer bases, and then use this knowledge to develop targeted marketing programs
- **City-planning:**
 - Identifying groups of houses according to their house type, value, and geographical location
- **Climate:**
 - Understanding earth climate, find patterns of atmospheric and ocean
- **Economic Science:**
 - market research

Clustering as a Preprocessing Tool

- **Summarization of data**
- **Finding K-nearest Neighbors**
 - Localizing search to one or a small number of clusters
- **Outlier detection**
 - Outliers are often viewed as those “far away” from any cluster
- **Image Processing**
 - Compression: cluster similar colors -> replace all the colors within a cluster with one color

What Is Good Clustering?

- A good clustering method will produce **high quality clusters**
 - high intra-class similarity: **cohesive** within clusters
 - low inter-class similarity: **distinctive** between clusters

Clustering Types

- **Representative-based Clustering**
- **Hierarchical Clustering**
- **Density-based Clustering**
- **Spectral and Graph Clustering**

Representative-based Clustering

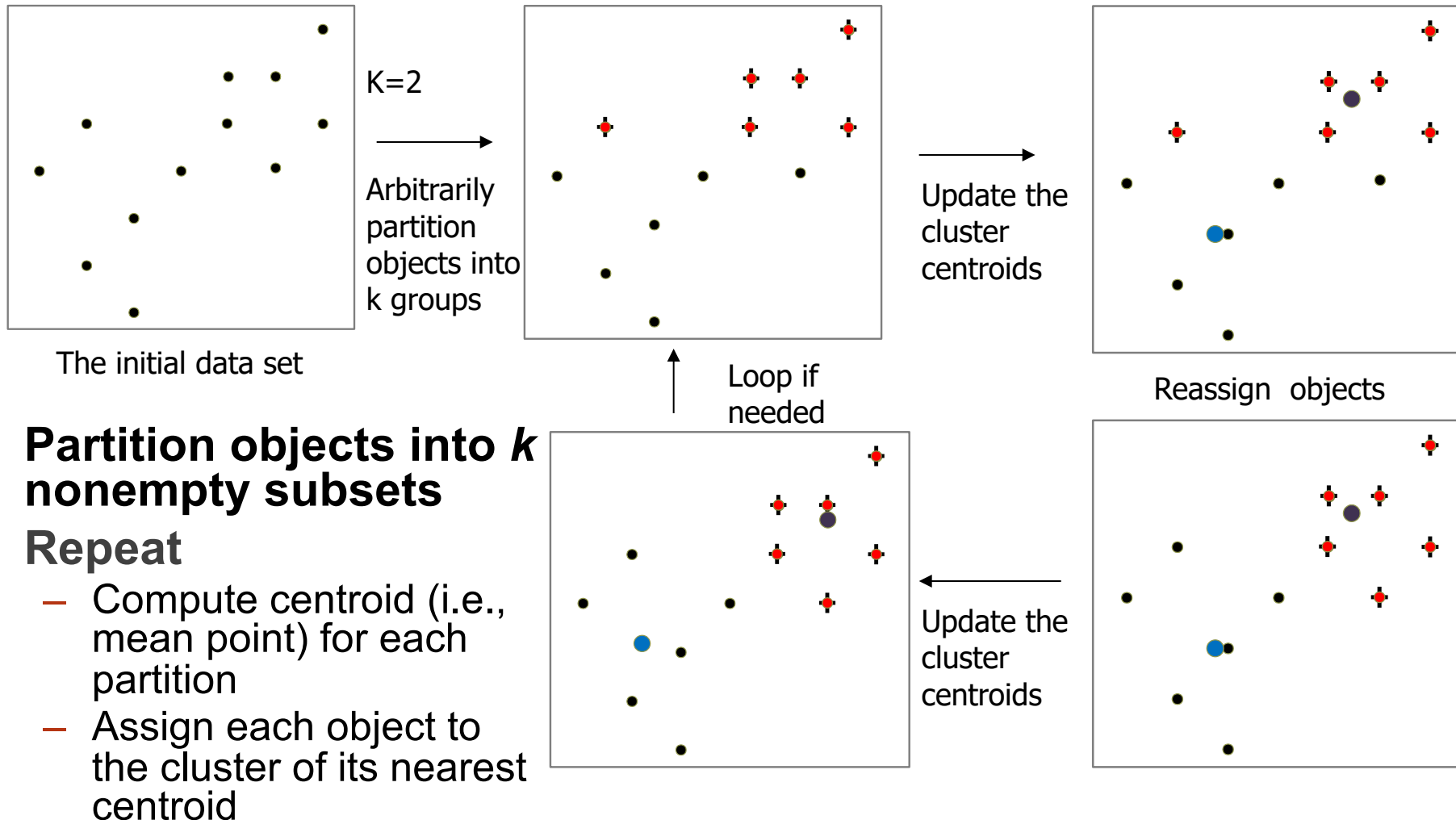
- **aka Prototype based clustering**
- **Given n data points and the number of desired cluster k**
 - partition the dataset into k groups or cluster
- **Data points in cluster are summarized with representative point**
 - Mean (aka centroid) of data points is popular
- **Brute-force/exhaustive approach**
 - generate all possible partitions of n points into k clusters:
 - $k^n/k!$: computationally infeasible with large n
 - evaluate some optimization score for each of them
 - retain the clustering that yields the best score

The *K-Means* Clustering Method

- Given k , the *k-means* algorithm is implemented in four steps:
 - Partition objects into k nonempty subsets
 - Compute seed points as the centroids of the clusters of the current partitioning (the centroid is the center, i.e., **mean point**, of the cluster)
 - Assign each object to the cluster with the nearest seed point
 - Go back to Step 2, stop when the assignment does not change

$$E = \sum_{i=1}^k \sum_{p \in C_i} (p - c_i)^2$$

An Example of K-Means Clustering

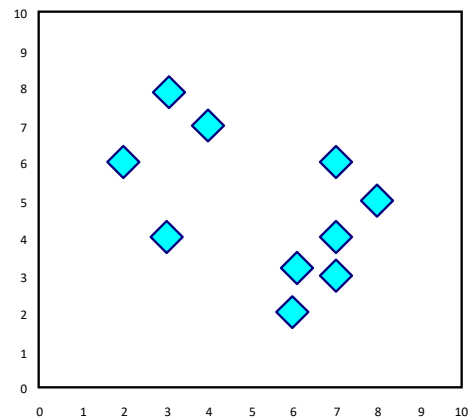


- **Partition objects into k nonempty subsets**
- **Repeat**
 - Compute centroid (i.e., mean point) for each partition
 - Assign each object to the cluster of its nearest centroid
- **Until no change**

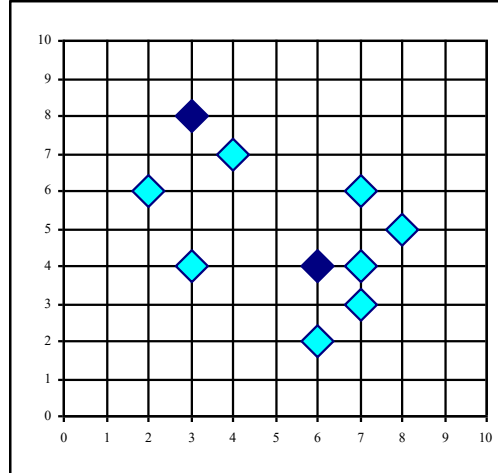
Comments on K-Means

- **Efficient algorithm: runs very fast**
- **Often terminates at a *local optimal***
- ***Cons:***
 - Applicable only to objects in a continuous n-dimensional space
 - Using the k-modes method for categorical data
 - In comparison, k-medoids can be applied to a wide range of data
 - Need to specify k , the *number* of clusters, in advance
 - there are ways to automatically determine the best k
 - Sensitive to noisy data and *outliers*
 - Not suitable to discover clusters with *non-convex shapes*

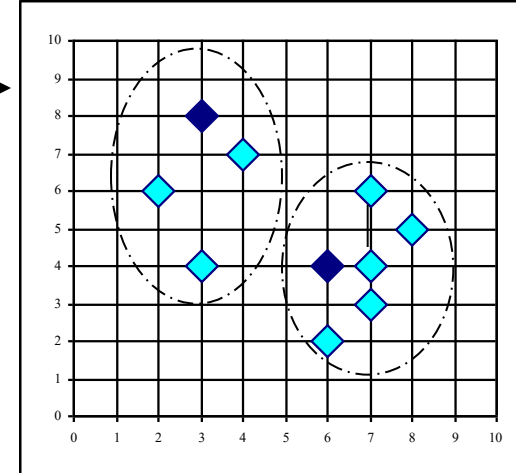
PAM: A Typical K-Medoids Algorithm



Arbitrary
choose k
object as
initial
medoids



Assign
each
remainin
g object
to
nearest
medoids



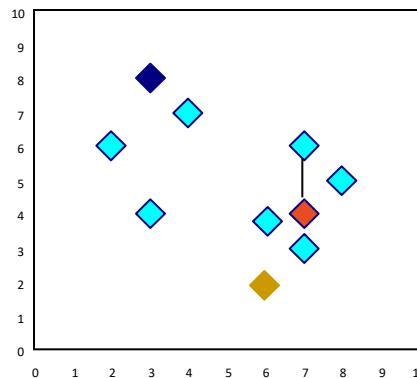
Total Cost = 20

$K=2$

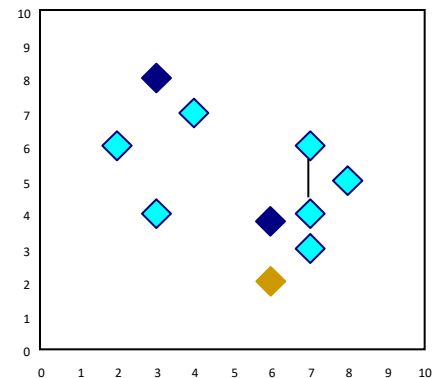
Do loop
Until no
change

Swapping O
and O_{random}
If quality is
improved.

Total Cost = 26



Compute
total cost of
swapping



Randomly select a
nonmedoid object, O_{random}

sklearn implementation

- **n_init = 10:**
 - run 10 times independently with different random centroids
- **max_iter = 300:**
 - max number of iteration for each run
 - stops if it converges early
- **tol=1e-04**
 - stop if change in center < tol
- **cluster_centers_:** a signature

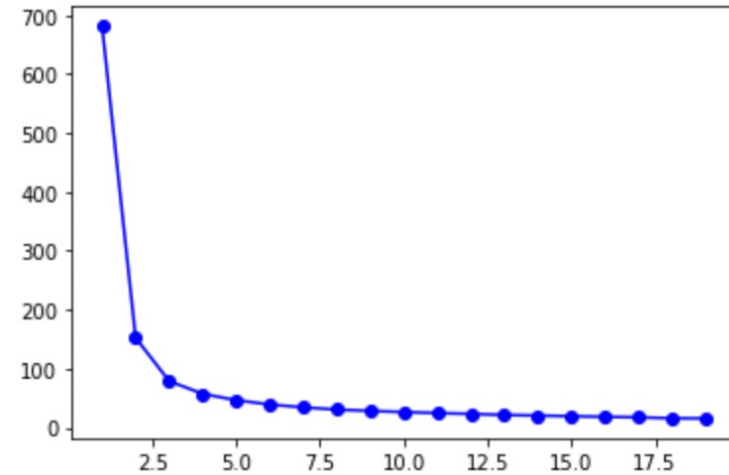
```
from sklearn.cluster import KMeans

km = KMeans(
    n_clusters=3, init='random',
    n_init=10, max_iter=300,
    tol=1e-04, random_state=0
)

y_km = km.fit_predict(X)
```

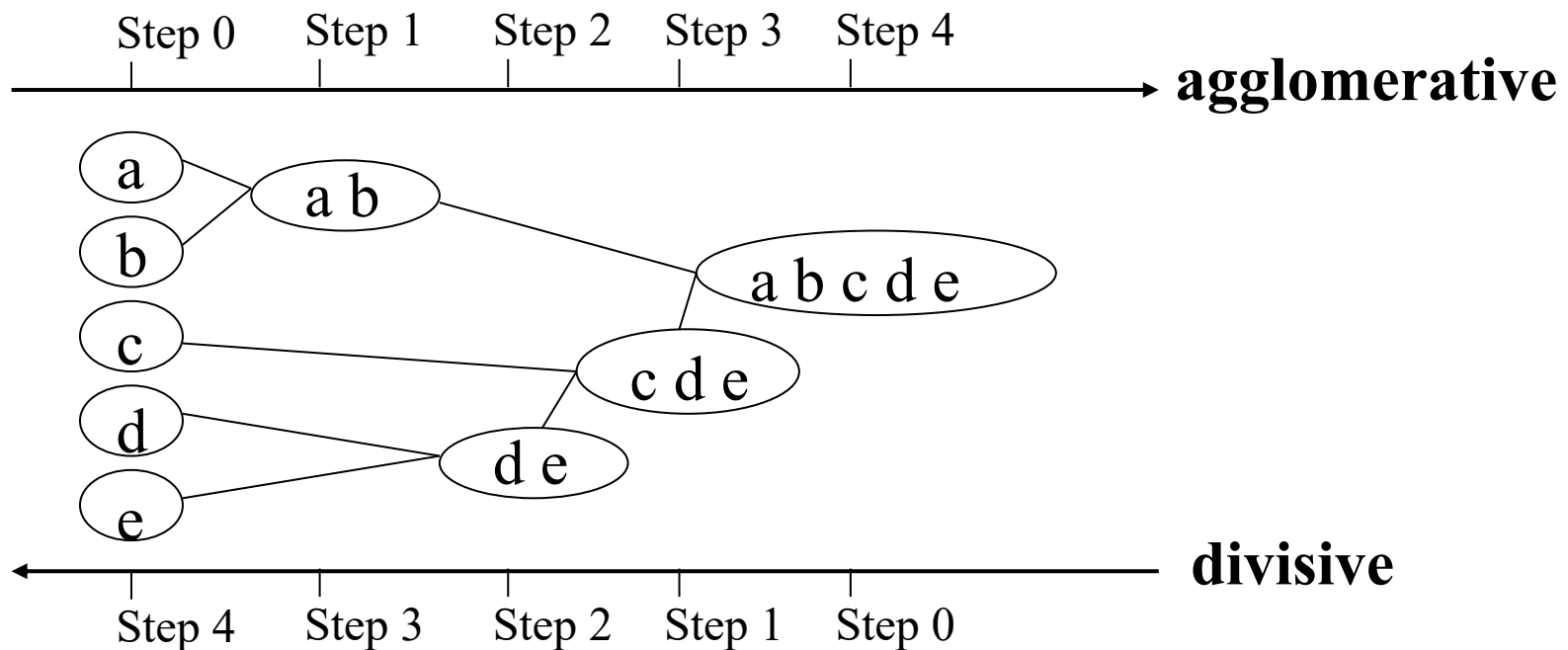
Choosing K

- **Elbow method**
 - Distortion/inertia vs K
 - Distortion: SSE $I = \sum_i (d(i, cr))^2$
- identify the value of k where the distortion begins to decrease most rapidly



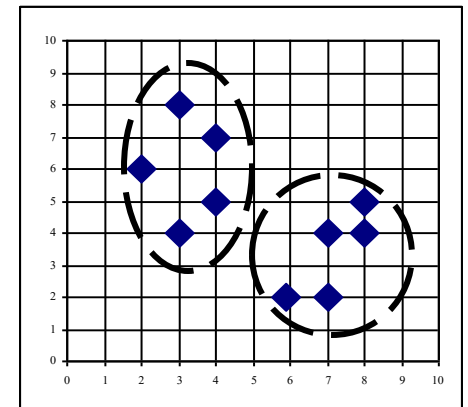
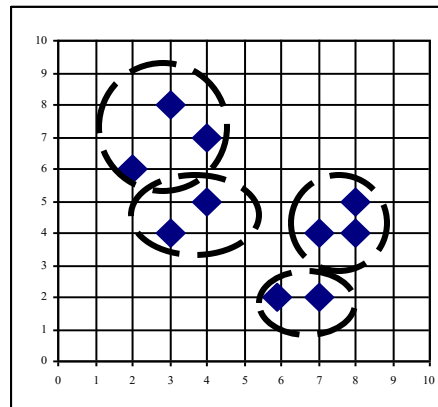
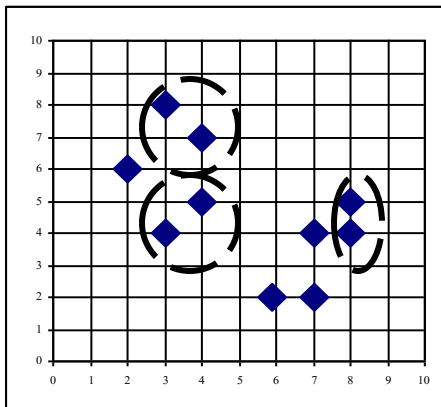
Hierarchical Clustering

- **Use distance matrix as clustering criteria**
 - No need to choose k
 - Need a terminating condition



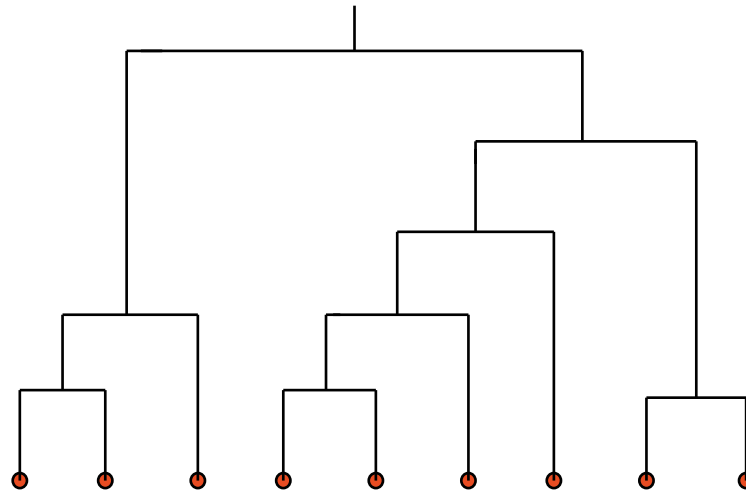
Agglomerative Clustering

- Use a link method and the dissimilarity matrix
- Merge nodes that have the least dissimilarity
- Eventually all nodes belong to the same cluster



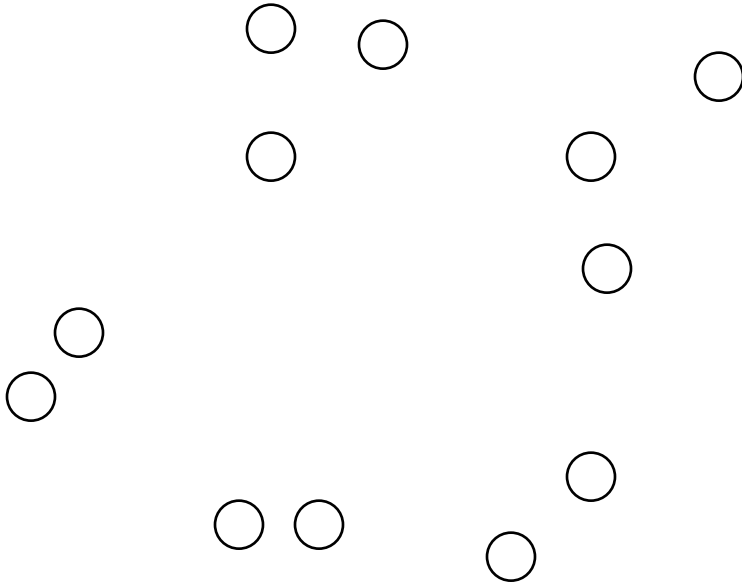
Dendrogram: Shows How Clusters are Merged

- **Decompose data points to several levels of nested partitioning**
 - Tree of clusters
- **A clustering is obtained by cutting the dendrogram at the desired level**



Steps 1 and 2

- Start with clusters of individual points and a proximity matrix



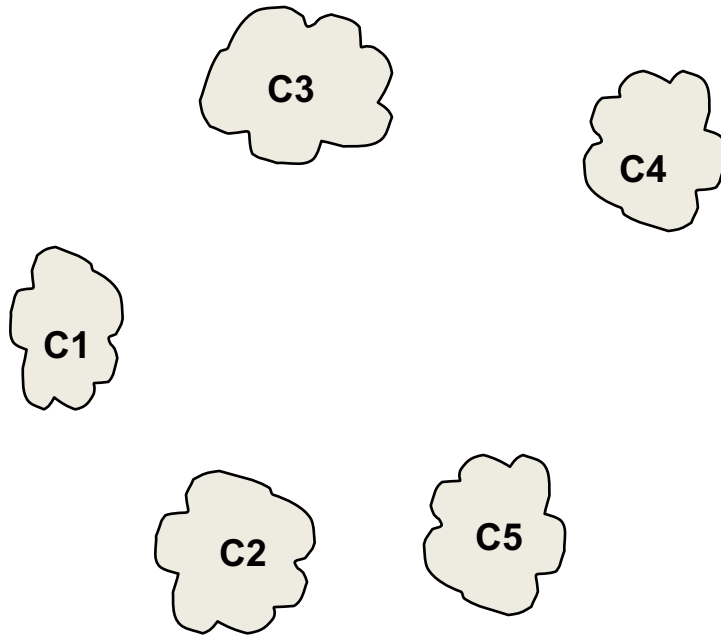
	p1	p2	p3	p4	p5	...
p1						
p2						
p3						
p4						
p5						
.						
.						
.						

Proximity Matrix



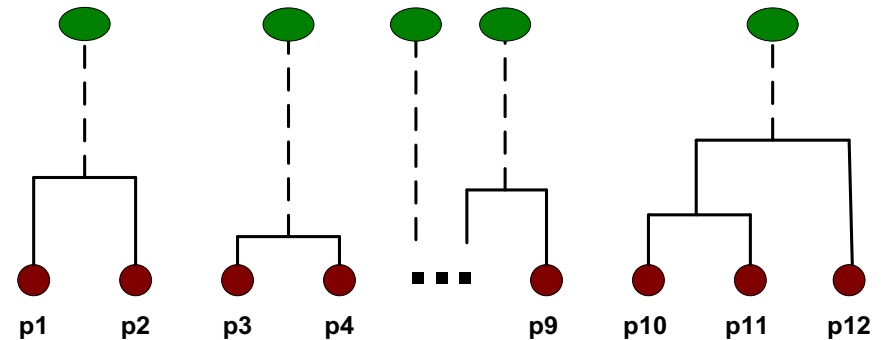
Intermediate Situation

- After some merging steps, we have some clusters



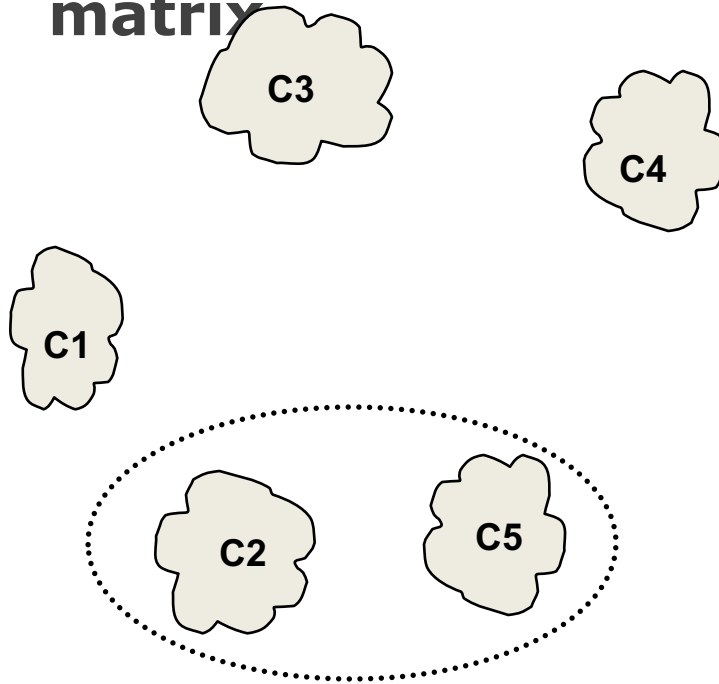
	C1	C2	C3	C4	C5
C1					
C2					
C3					
C4					
C5					

Proximity Matrix



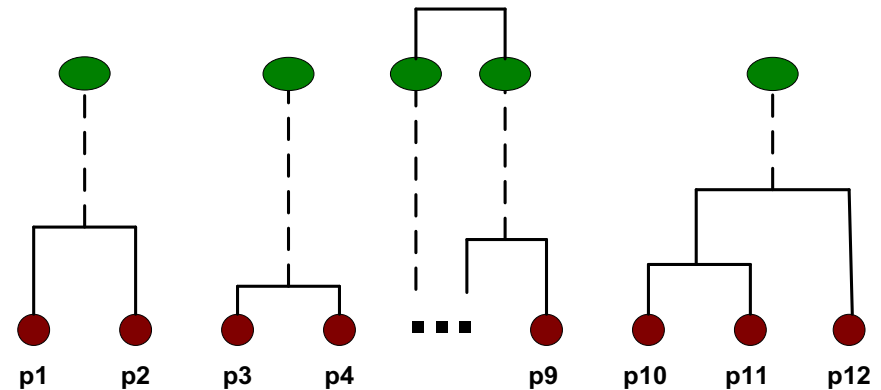
Step 4

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



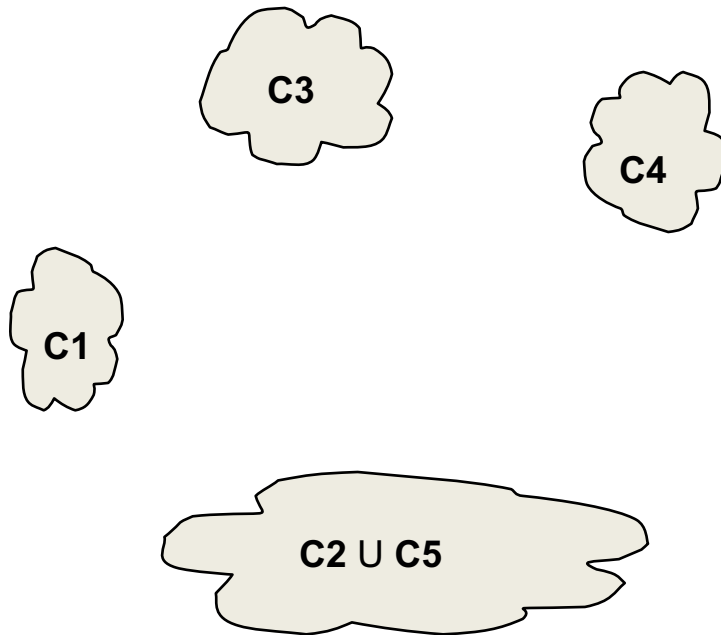
	C1	C2	C3	C4	C5
C1					
C2					
C3					
C4					
C5					

Proximity Matrix



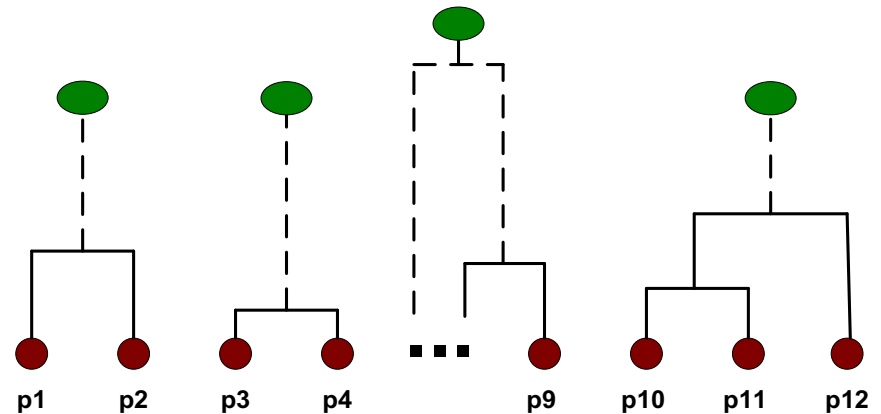
Step 5

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

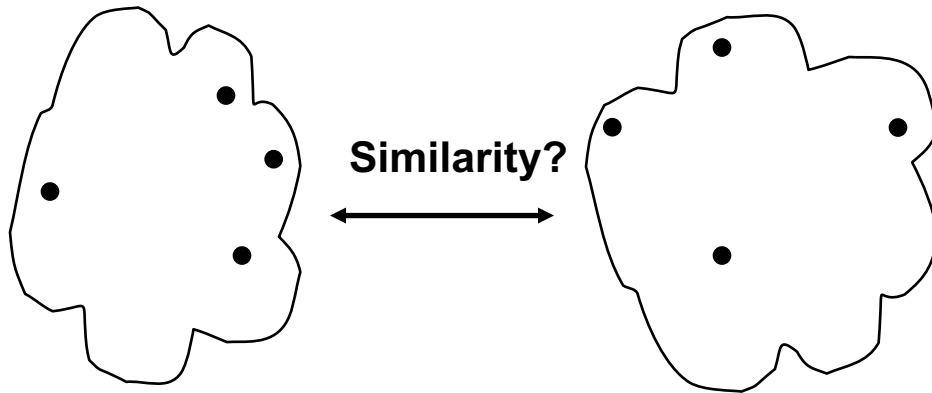


	C1	$\begin{matrix} C2 \\ \cup \\ C5 \end{matrix}$	C3	C4
C1		?		
$C2 \cup C5$?	?	?	?
C3		?		
C4		?		

Proximity Matrix



How to Define Inter-Cluster Distance

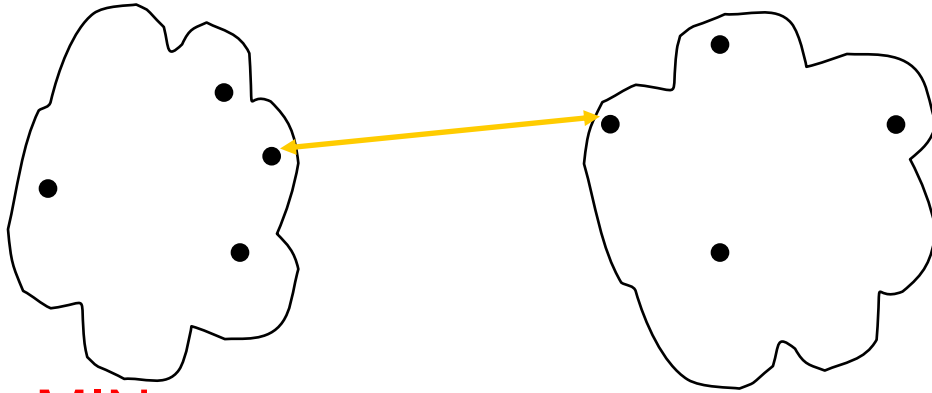


- MIN
- MAX
- Group Average
- Distance Between Centroids
- Other methods driven by an objective function
 - Ward's Method uses squared error

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

Proximity Matrix

How to Define Inter-Cluster Similarity

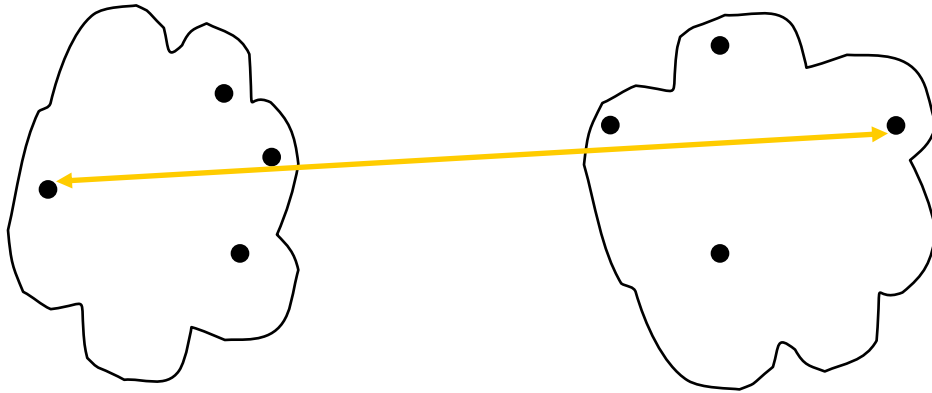


- MIN
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p2						
p3						
p4						
p5						
.						
.						
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Proximity Matrix

How to Define Inter-Cluster Similarity

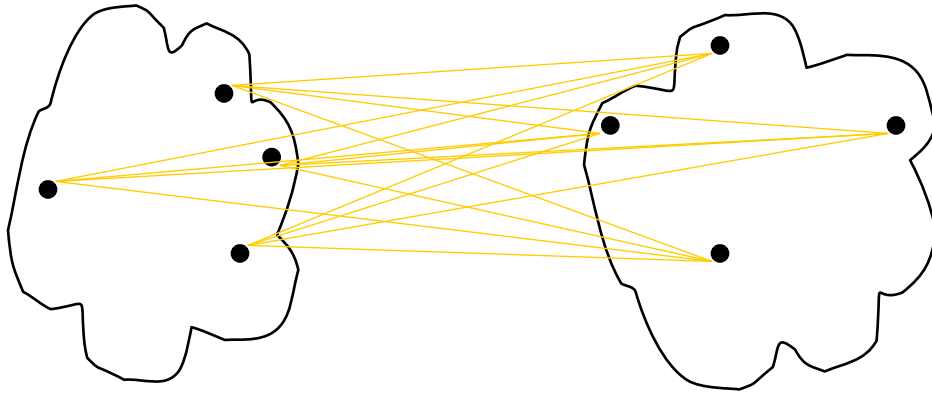


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Proximity Matrix

How to Define Inter-Cluster Similarity

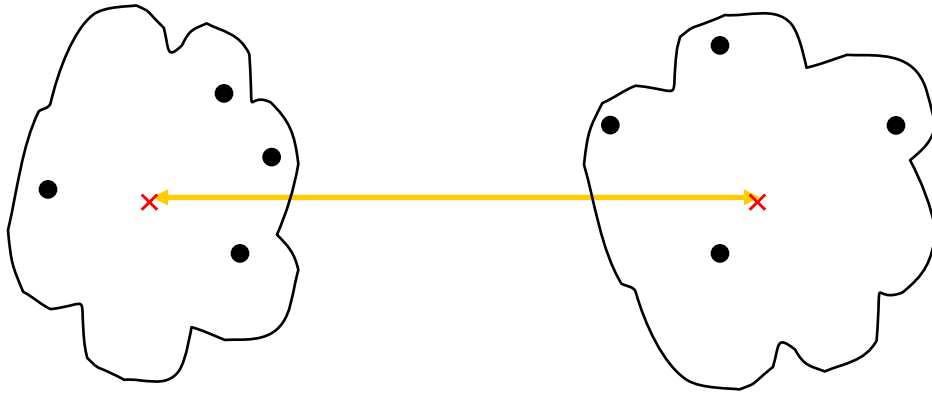


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Proximity Matrix

How to Define Inter-Cluster Similarity



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p1						
p2						
p3						
p4						
p5						
.						
.						
.						

Proximity Matrix

Distance between Clusters

- **Clusters are merge based on distance**
- **Single link:**
 - smallest distance between an element in one cluster and an element in the other, i.e., $\text{dist}(K_i, K_j) = \min \text{dist}(t_{ip}, t_{jq})$
- **Complete link:**
 - largest distance between an element in one cluster and an element in the other, i.e., $\text{dist}(K_i, K_j) = \max \text{dist}(t_{ip}, t_{jq})$
- **Average:**
 - avg distance between an element in one cluster and an element in the other, i.e., $\text{dist}(K_i, K_j) = \text{avg dist}(t_{ip}, t_{jq})$
- **Ward:**
 - based on minimizing the variance between clusters (SSE)

Issues with Hierarchical Clustering

- **Can never undo what was done previously**
 - Compare with k-means
- **Do not scale well**
 - time complexity $O(n^2)$

Density-Based Clustering

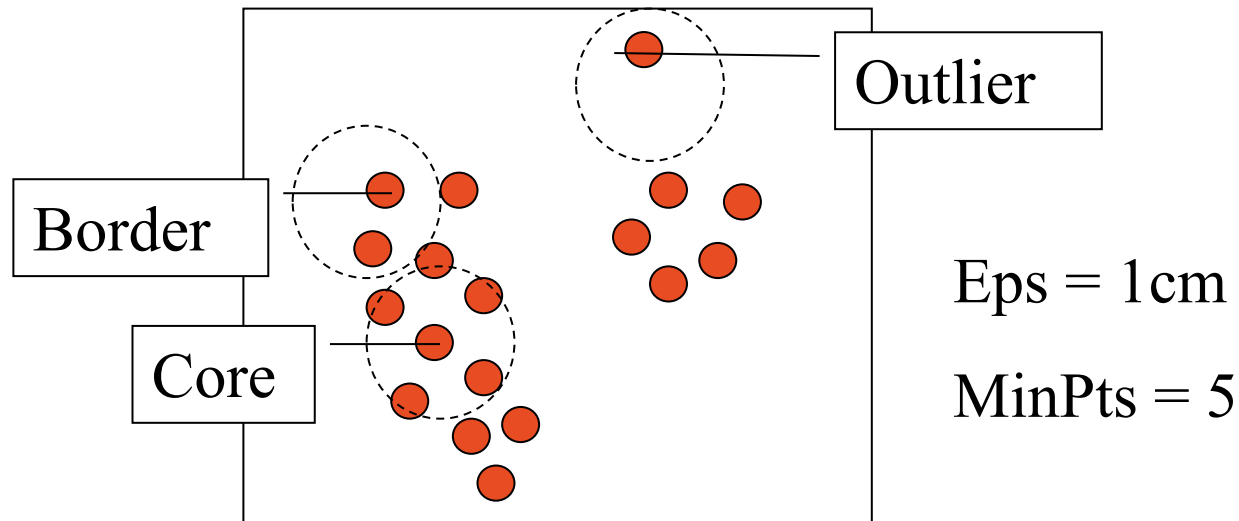
- **Clustering based on density (local cluster criterion), such as density-connected points**
- **Major features:**
 - Discover clusters of arbitrary shape
 - Handle noise
 - One scan
 - Need density parameters as termination condition
- **Example**
 - DBSCAN, OPTICS, DENCLUE

Density-Based Clustering: Basic Concepts

- Classifying points based on the characteristic of their local neighborhood
- **Two parameters:**
 - **Eps**: Maximum radius of the neighborhood
 - **MinPts**: Minimum number of points in an Eps-neighborhood of that point
- **$N_{Eps}(p)$: $\{q \text{ belongs to } D \mid \text{dist}(p,q) \leq Eps\}$**

DBSCAN: Density-Based Spatial Clustering of Applications with Noise

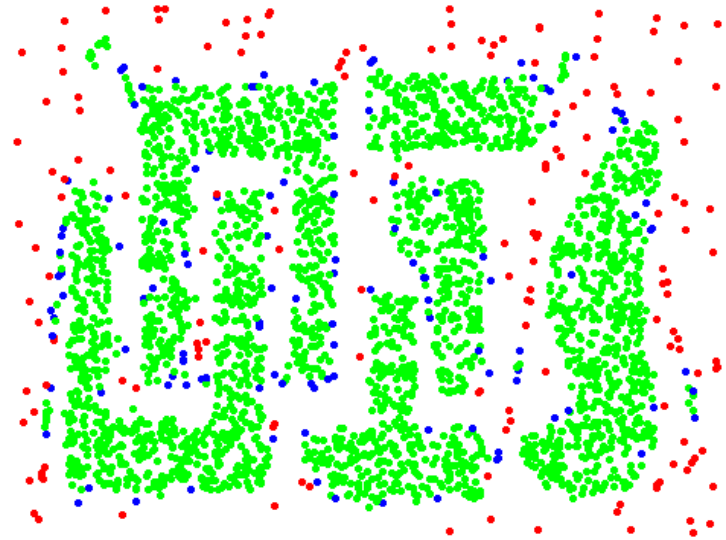
- **Relies on a *density-based* notion of cluster:**
 - A *cluster* is defined as a maximal set of density-connected points
- **Discovers clusters of arbitrary shape in spatial databases with noise**



DBSCAN: Core, Border and Noise Points



Original Points

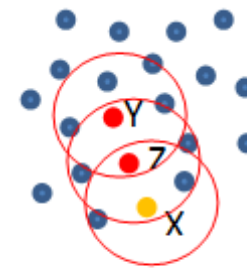


Point types: **core**,
border and **noise**

Eps = 10, MinPts = 4

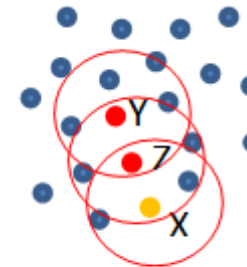
DBSCAN: The Algorithm

- A point is considered reachable from another point if there is a path consisting of core points between the starting and ending point
- Any point that is not reachable is considered an outlier



X is density reachable from Y,
but Y is not density reachable
from X

a. Density-reachability of points



X and Y are density
connected by Z.

b. Density connectivity of points

DBSCAN: The Algorithm

- **Arbitrary select a point p**
- **Retrieve all points density-reachable from p w.r.t. 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**

Measures of Cluster Validity

- **Numerical measures that are applied to judge various aspects of cluster validity, are classified into the following two types.**
 - **Supervised:** Used to measure the extent to which cluster labels match externally supplied class labels.
 - Entropy
 - Often called *external indices* because they use information external to the data
 - **Unsupervised:** Used to measure the goodness of a clustering structure *without* respect to external information.
 - Sum of Squared Error (SSE)
 - Often called *internal indices* because they only use information in the data
- **You can use supervised or unsupervised measures to compare clusters or clusterings**

Unsupervised 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
- **Example:**
 - Silhouette score
 - Dunn Index