



3D geoinformation

Department of Urbanism  
Faculty of Architecture and the Built Environment  
Delft University of Technology

GEO5017

Machine Learning for the Built Environment

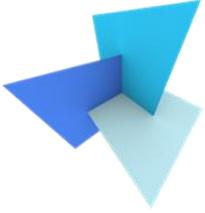
<https://3d.bk.tudelft.nl/courses/geo5017/>

# Clustering & Nearest Neighbor Classification

Liangliang Nan

<https://3d.bk.tudelft.nl/liangliang/>

# Agenda

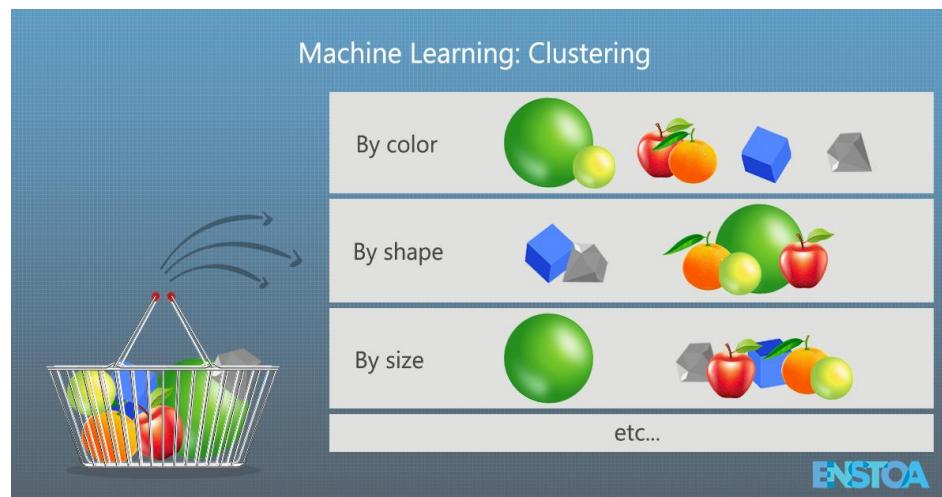


- Overview
  - What is clustering?
  - Distance measure (similarity measure)
  - Types of clustering algorithms
- Clustering algorithms
  - K-means clustering
  - Hierarchical clustering
  - Density-based clustering
- Nearest neighbor classification
- Features



# What is clustering?

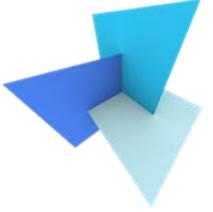
- Clustering
  - A process that **partitions** a given dataset into homogeneous groups based on given features such that **similar** objects are kept in a group whereas **dissimilar** objects are in different groups.



What is a cluster?

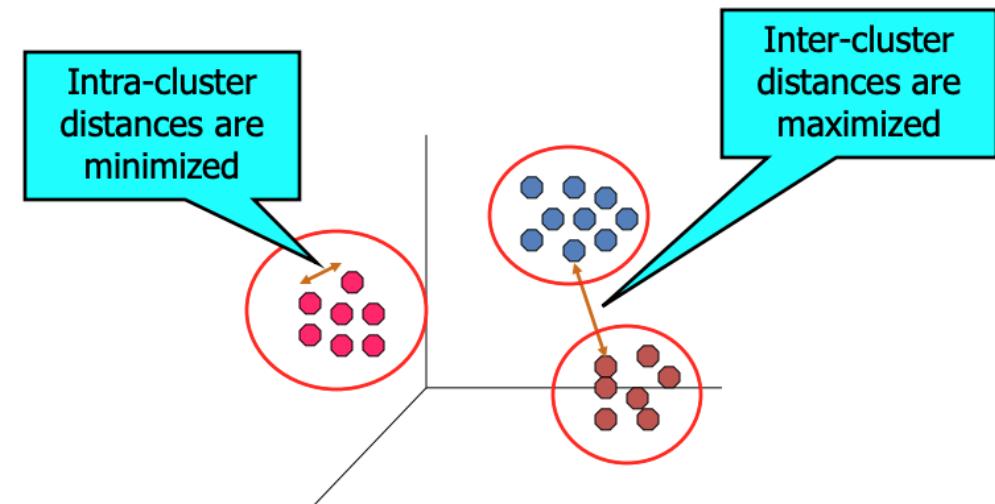
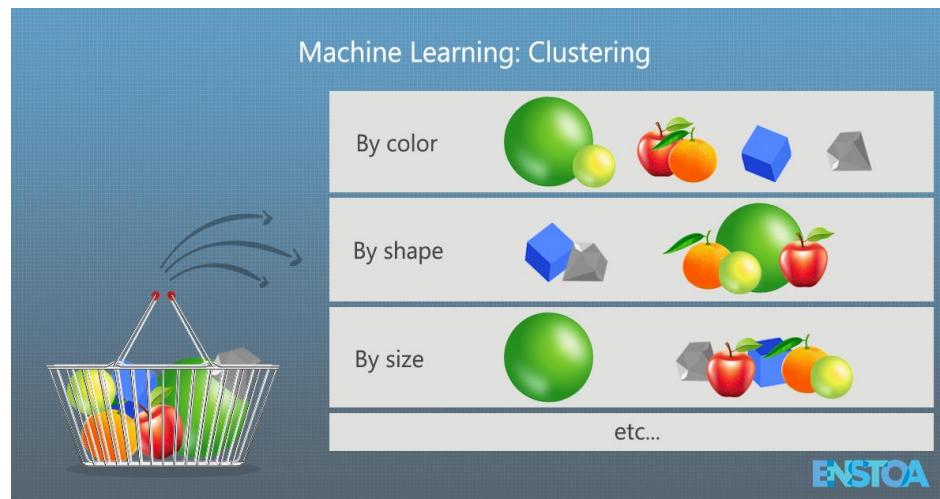
What constitutes a good cluster?

What is the “best” criterion for clustering?



# What is clustering?

- Clustering
  - A process that **partitions** a given dataset into homogeneous groups based on given features such that **similar** objects are kept in a group whereas **dissimilar** objects are in different groups.





# What is clustering?

- Clustering: two components in an algorithm
  - Distance measure → defines similarities
  - Clustering algorithm → partitions the dataset



Different distance measures lead to different clustering results



# Distance measure

- Problem dependent
  - Minkowski distance/metric is often used
    - Generalization of Euclidean distance ( $L^2$ ) and Manhattan distance ( $L^1$ )

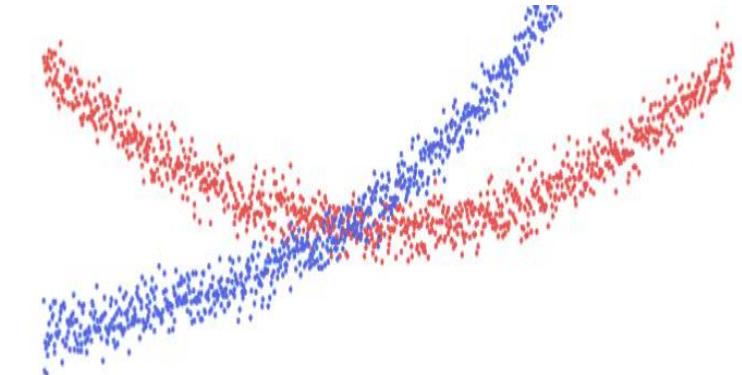
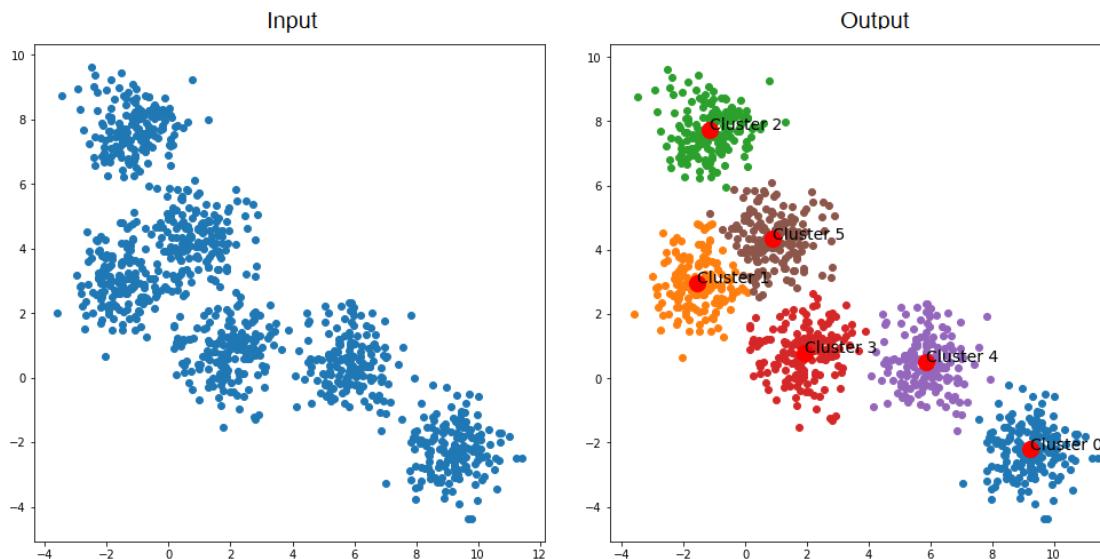
$$D(x_i, x_j) = \left( \sum_{k=1}^d |x_{i,k} - x_{j,k}|^p \right)^{\frac{1}{p}}$$

- Domain knowledge is required
  - When components of data feature vectors not immediately comparable, e.g.,
    - color vs size
    - distance to city center vs energy label

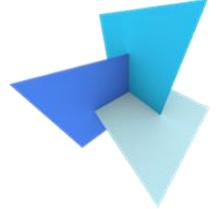


# Types of clustering algorithms

- Different criteria
  - Exclusive vs overlapping
    - Whether a data point can belong to two or more clusters



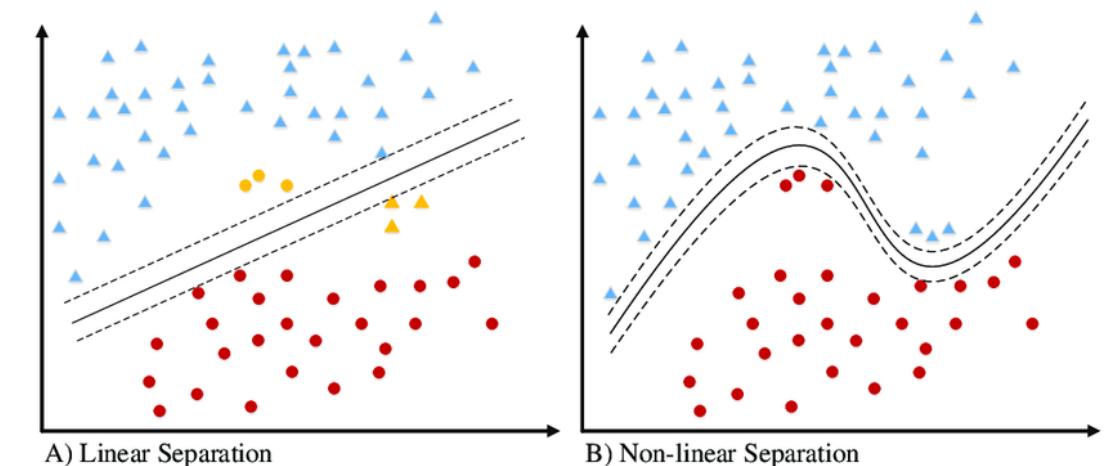
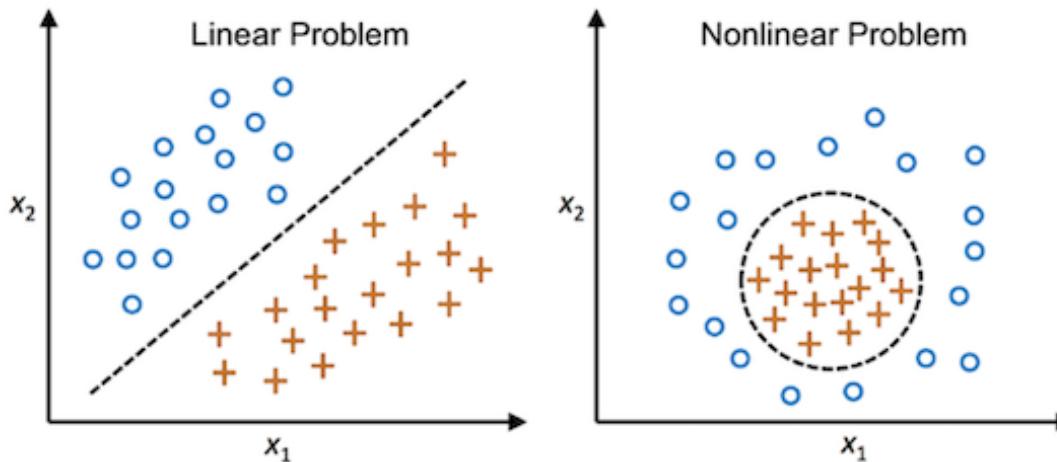
# Types of clustering algorithms



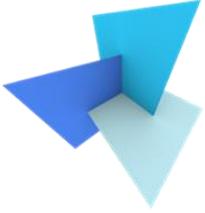
- Different criteria
  - Exclusive vs overlapping
    - Whether a data point can belong to two or more clusters
  - Linear vs non-linear
    - The applicability to different types of data

We will learn:

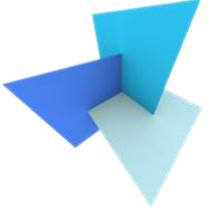
- Linear: K-means, hierarchical clustering
- Non-linear: density-based clustering



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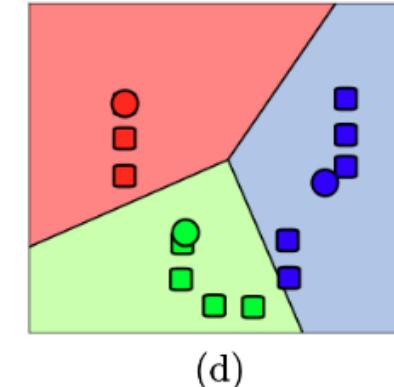
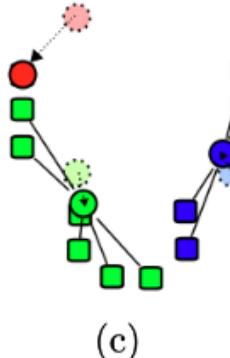
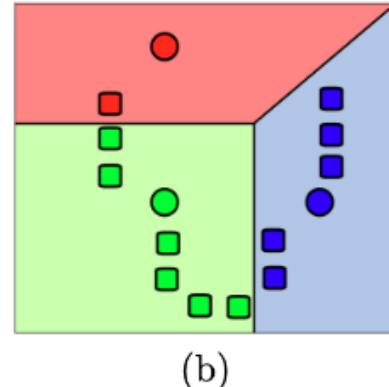
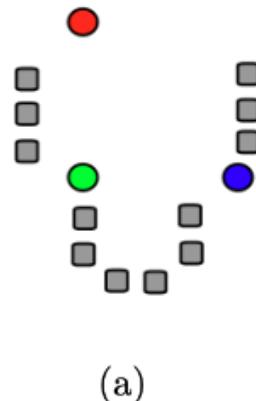


# k-means

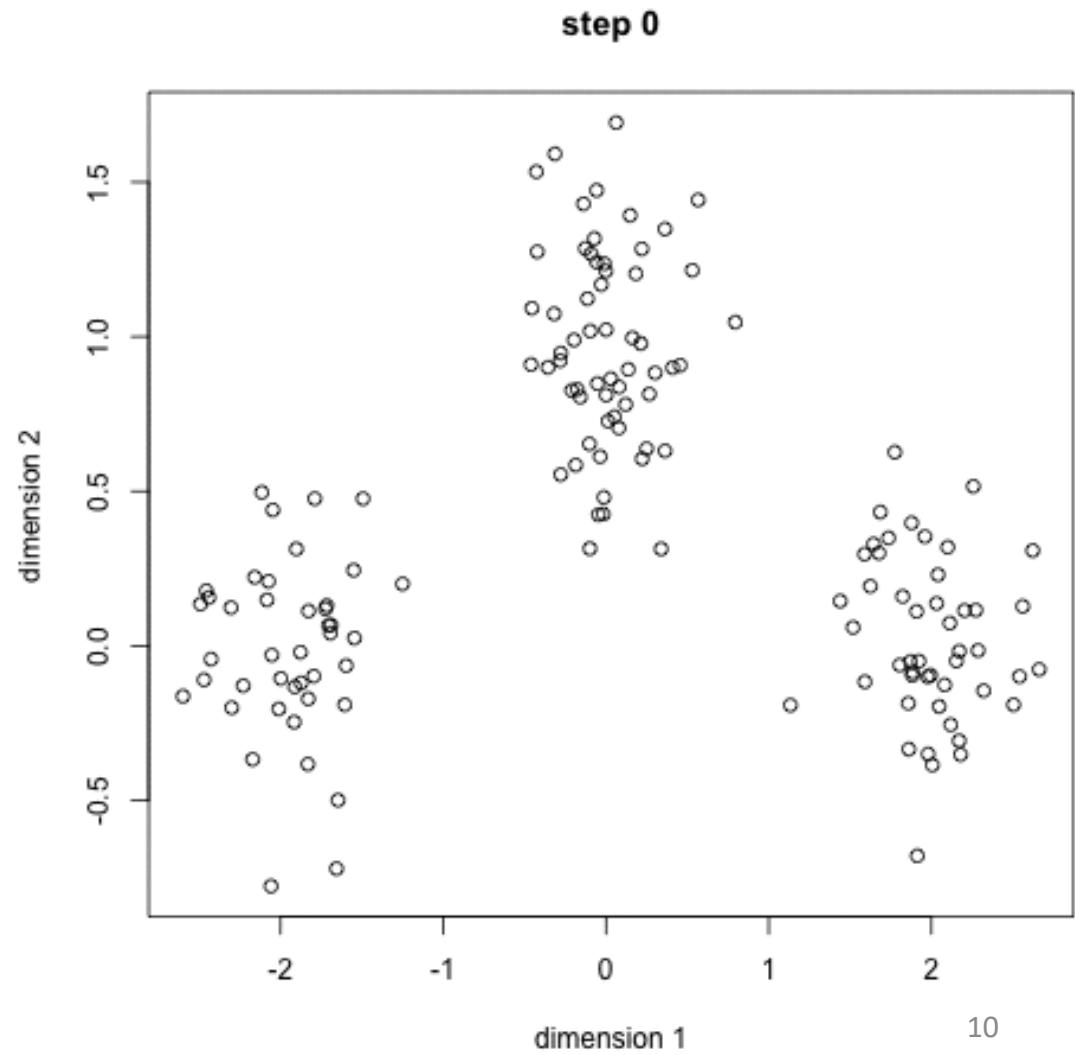
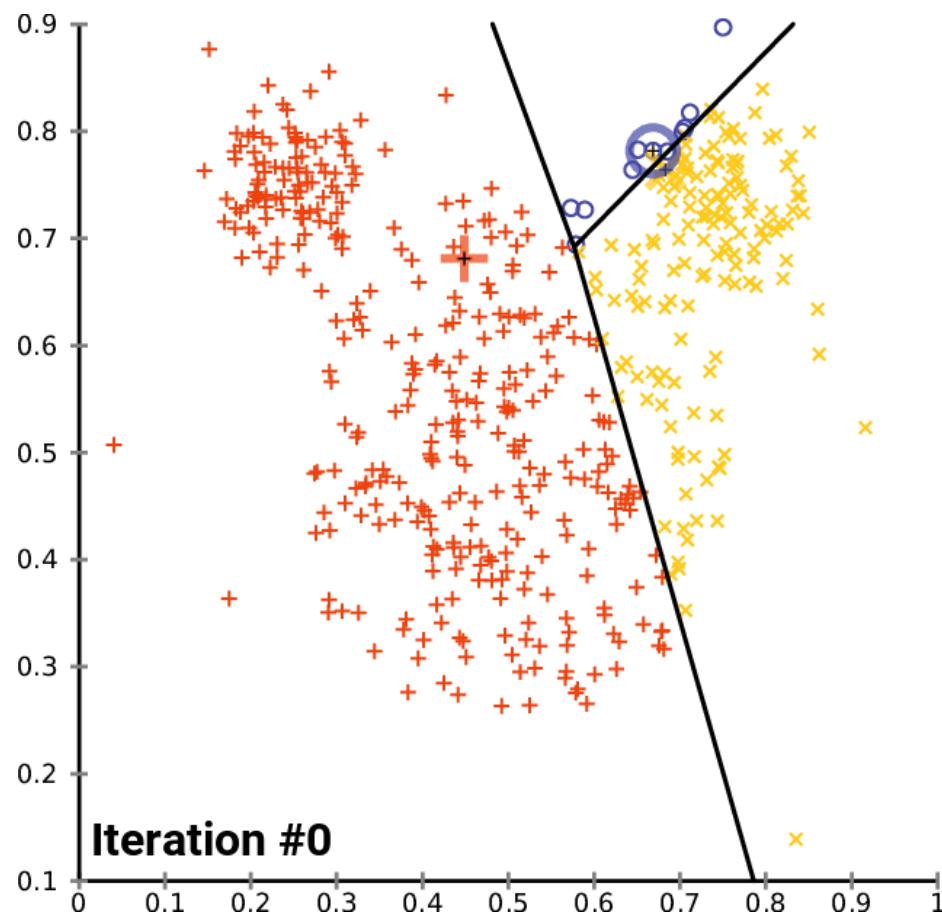
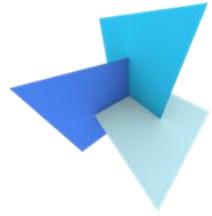
- 1) Initialize the  $k$  clusters  $\ell^0 = \{c_1^0, c_2^0 \dots c_k^0\}$  in a way such that the initial centroids are placed as far as possible from each other.
- 2) Calculate the centroids of the clusters:  $u_j^i = \frac{1}{|c_j^i|} \sum_{x \in c_j^i} x$ , where  $j = 1, \dots, k$  and  $i$  denotes the  $i$ -th iteration.
- 3) Take each point belonging to a given data set and associate it to the nearest centroid:

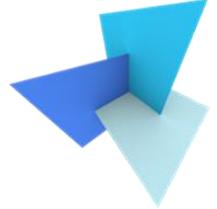
$$\begin{aligned} c_j^{i+1} &= \{x \mid d(x, u_j^i) \leq d(x, u_{j'}^i), \forall j', 1 \leq j' \leq k\} \\ \ell^{i+1} &= \{c_j^{i+1} \mid 1 \leq j \leq k\} \end{aligned} \quad (2)$$

- 4) Repeat steps 2 and 3 until no more changes can be made to the clusters, i.e.,  $\ell^{i+1} = \ell^i$ . In other words, centroids do not move any more.



# k-means





# k-means: an optimization perspective

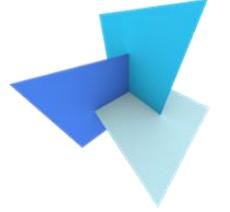
- Objective function

- SSE (Sum of Squared Error)

$$J = \sum_{i=1}^k \sum_{x \in c_i} \|x - u_i\|^2$$



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



# k-means: an optimization perspective

- Objective function

- SSE (Sum of Squared Error)

$$J = \sum_{i=1}^k \sum_{x \in c_i} \|x - u_i\|^2$$

- Convergence

- K-means is exactly coordinate descent on  $J$

- Step 2: fixed cluster assignment—compute cluster centroids that minimize the current error
    - Step 3: fixed cluster centroids—find cluster assignment that minimizes the current error

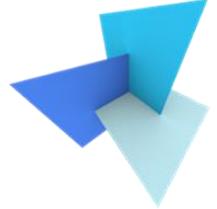
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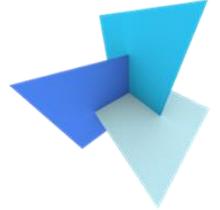
# k-means: an optimization perspective

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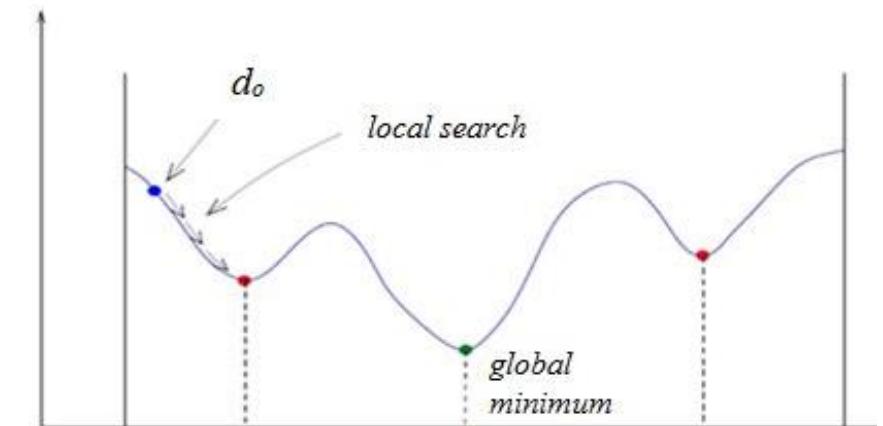
$J$  converges a global minimum?



# k-means: an optimization perspective



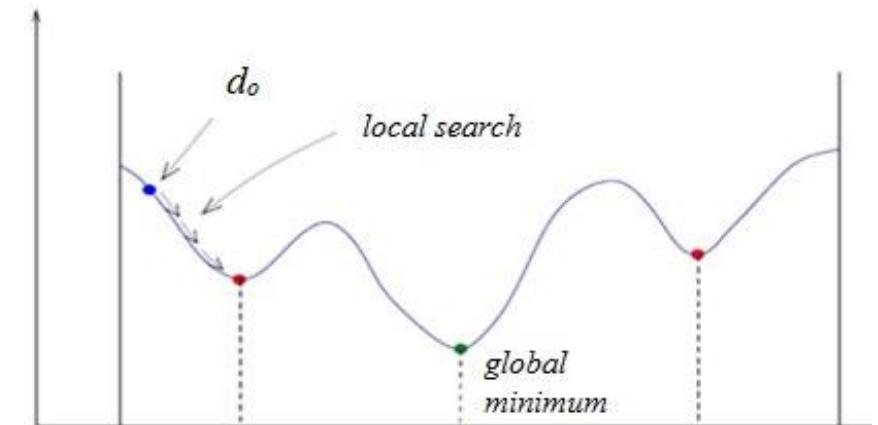
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- Not necessarily the optimal configuration
  - i.e., local minimum of the objective function



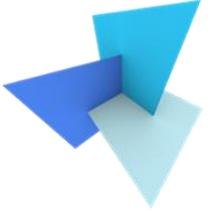


# k-means: an optimization perspective

- Objective function
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  - K-means is exactly coordinate descent on  $J$ 
    - Step 2: fixed cluster assignment—compute cluster centroids that minimize the current error
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- Not necessarily the optimal configuration
  - i.e., local minimum of the objective function
  - Solution: repeat many times and pick the best
  - Best configuration not guaranteed



# k-means



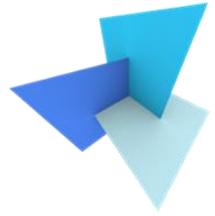
- Advantages
  - Fast and efficient
  - Given good results when groups are distinct or well separated from each other
  - Easy to implement



# k-means

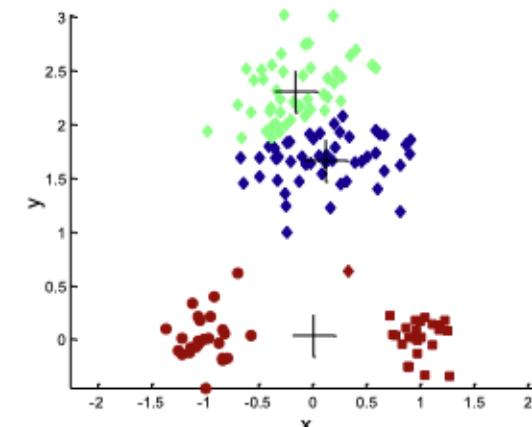
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- Limitations

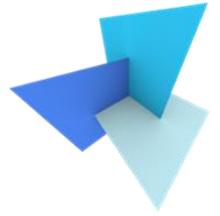




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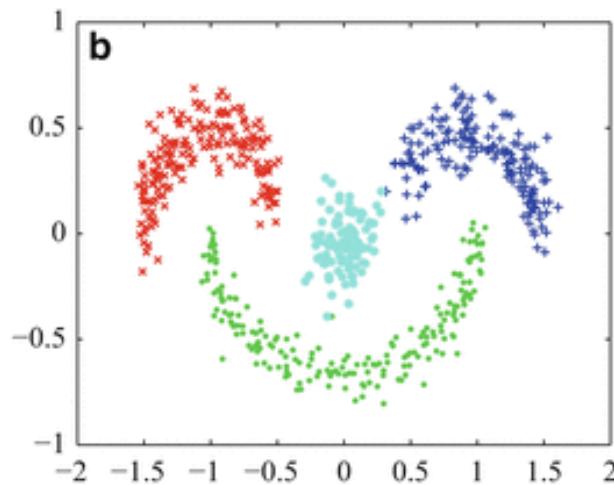
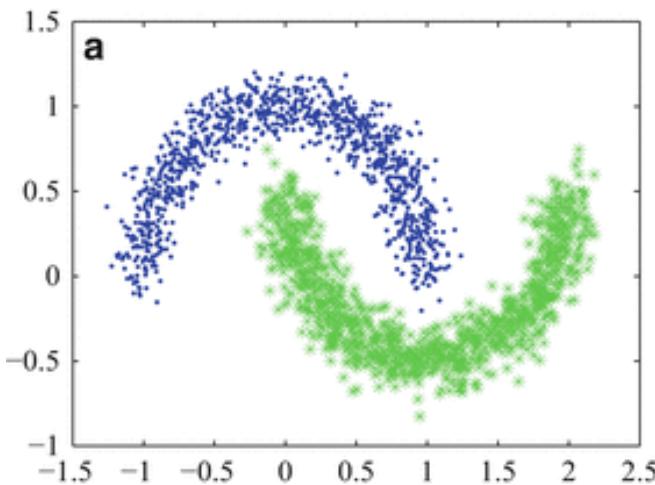
- Advantages
  - Fast and efficient
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  - Easy to implement
- Limitations
  - Requires a priori specification of the number (i.e.,  $k$ ) of clusters
  - Local minima
    - Sensitive to initialization
    - Cannot guarantee optimal clusters
  - Not invariant to non-linear transformations
    - e.g., cartesian coordinates vs polar coordinates



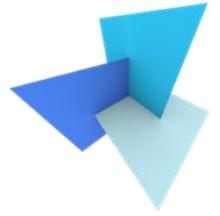


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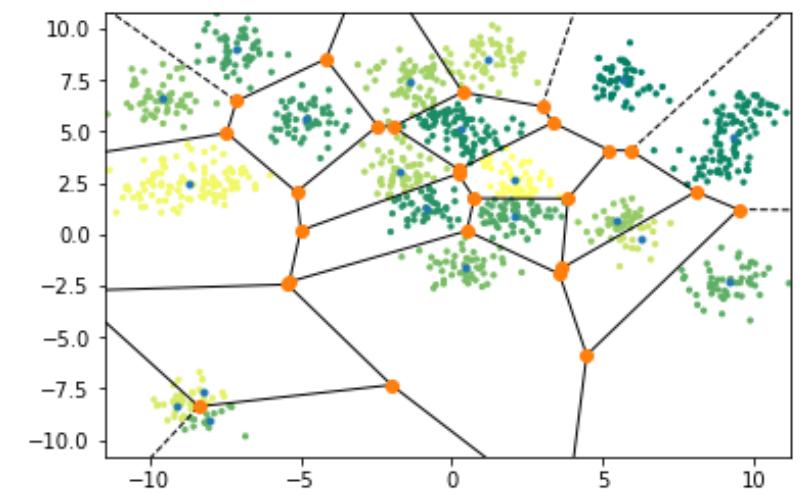
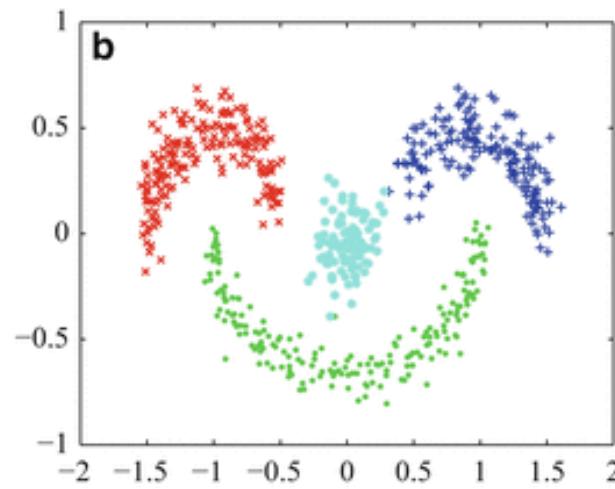
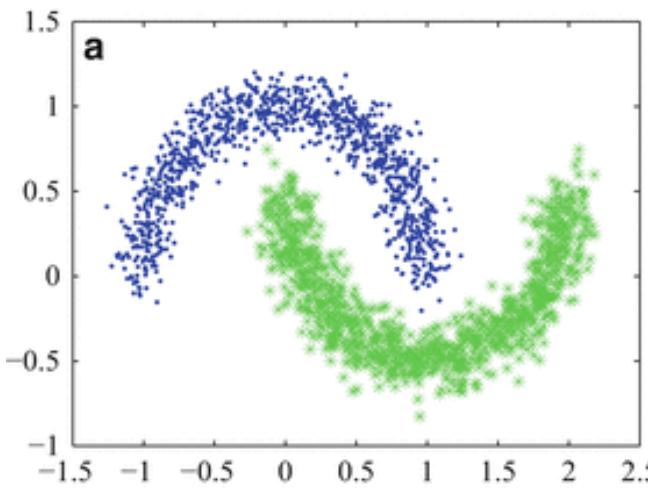
- Can k-means handle?

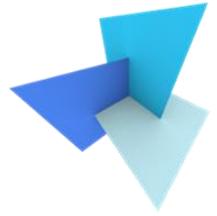


# k-means



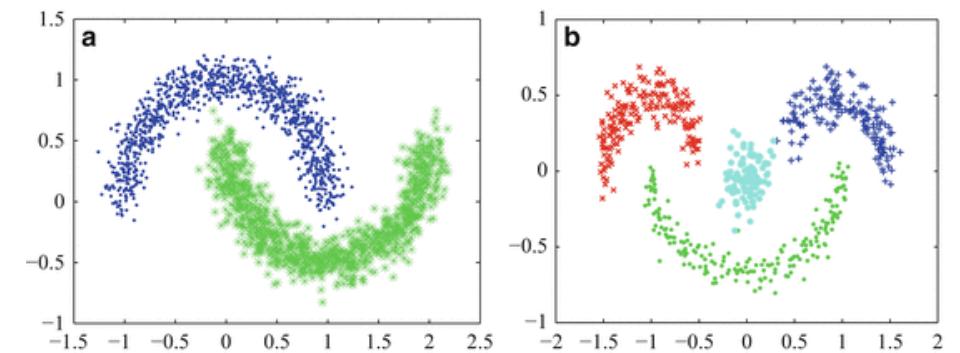
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# k-means

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- Limitations
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    - Cannot guarantee optimal clusters
  - Not invariant to non-linear transformations
    - e.g., cartesian coordinates vs polar coordinates
  - Cannot process non-linear datasets

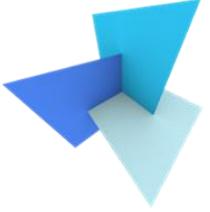




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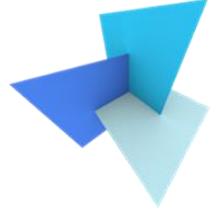




# Hierarchical clustering

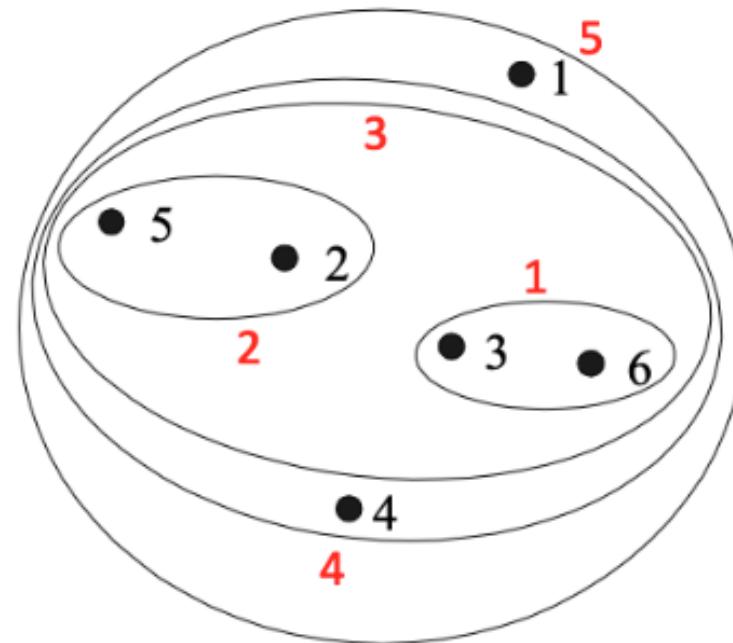
Given a set of  $N$  objects  $S = \{s_1, s_2, \dots, s_N\}$  to be clustered and a function of distance between two clusters  $c_i$  and  $c_j$ , build a hierarchy tree on  $S$  such that for every  $c_i, c_j \in S$ ,  $c_i \cap c_j = \emptyset$ . The basic process of hierarchical clustering is as follows:

- 1) Start by assigning each object to a cluster  $c_i = s_i (i = 1, \dots, N)$ , so that if you have  $N$  objects, you have  $N$  clusters  $\ell = \{c_1, c_2, \dots, c_N\}$ , each containing just one item.
- 2) Find the pair of clusters  $(c_i, c_j)$  such that  $D(c_i, c_j) \leq D(c_{i'}, c_{j'})$ ,  $\forall c_{i'} \neq c_{j'} \in \ell$  and merge them into a single cluster  $c_k = c_i \cup c_j$ . Delete  $c_i$  and  $c_j$  from  $\ell$  and insert  $c_k$  into  $\ell$  so that now you have one cluster less.
- 3) Compute distances (similarities) between the new cluster and each of the old clusters.
- 4) Repeat steps 2) and 3) until all items are clustered into a single cluster of size  $N$ .

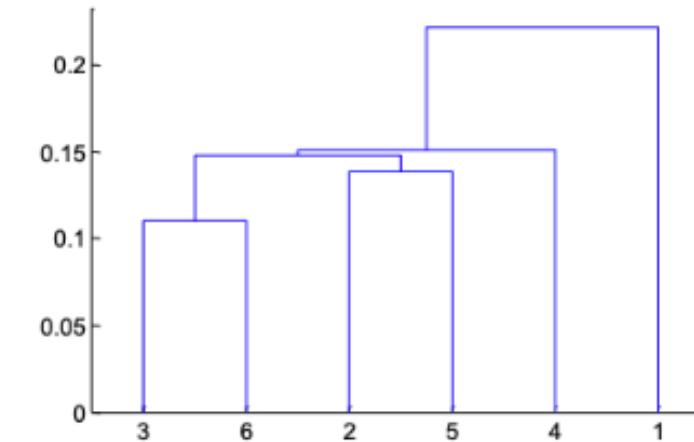


# Hierarchical clustering

- Example



Nested Clusters



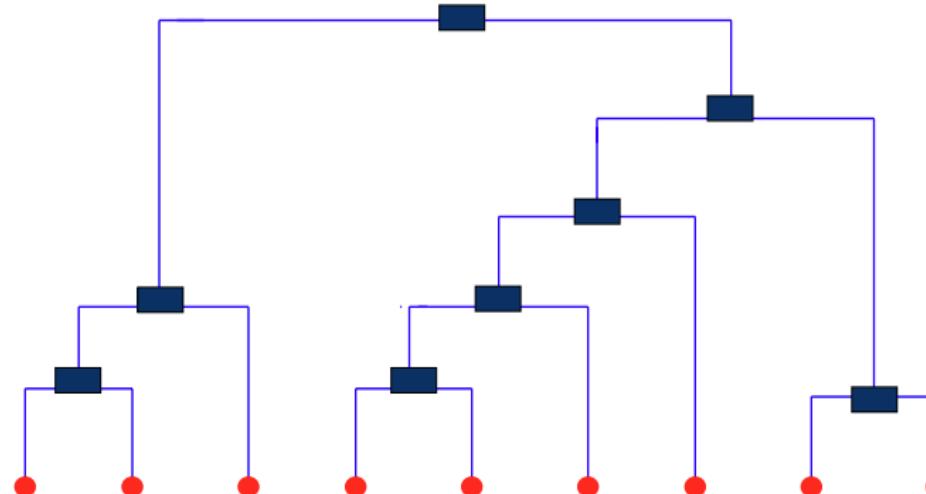
Dendrogram

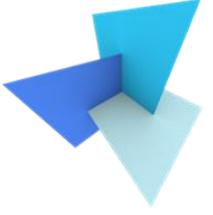
An example of hierarchical clustering



# Hierarchical clustering

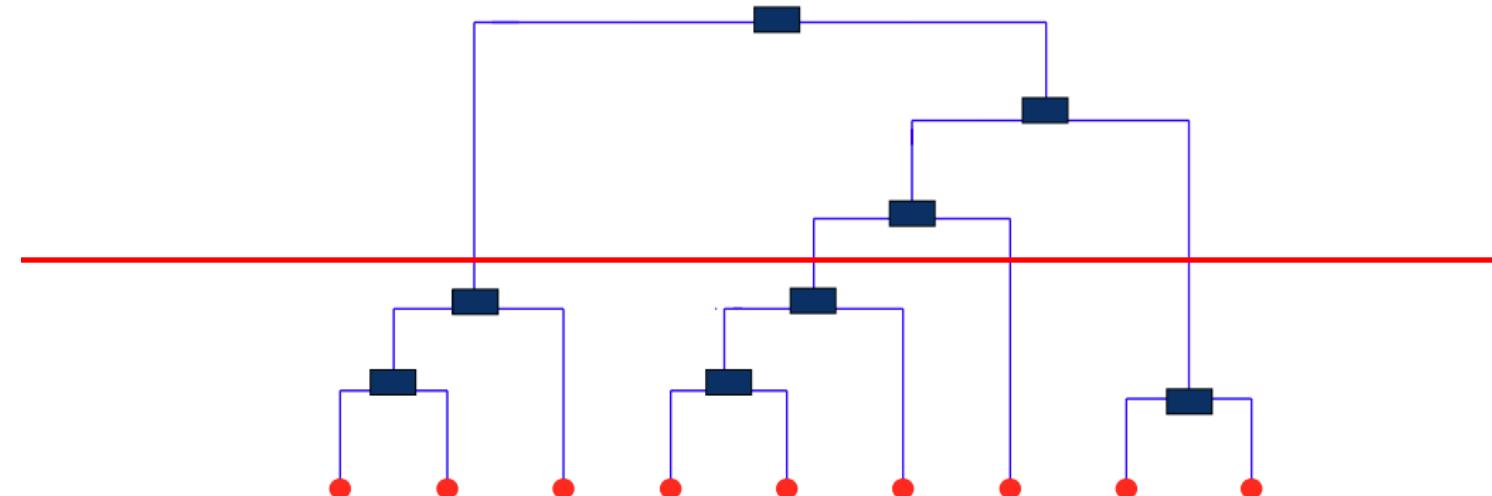
- Dendrogram
  - A tree that shows how clusters are merged/split hierarchically
  - Each node on the tree is a cluster; each leaf node is a singleton cluster



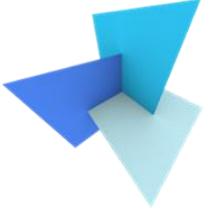


# Hierarchical clustering

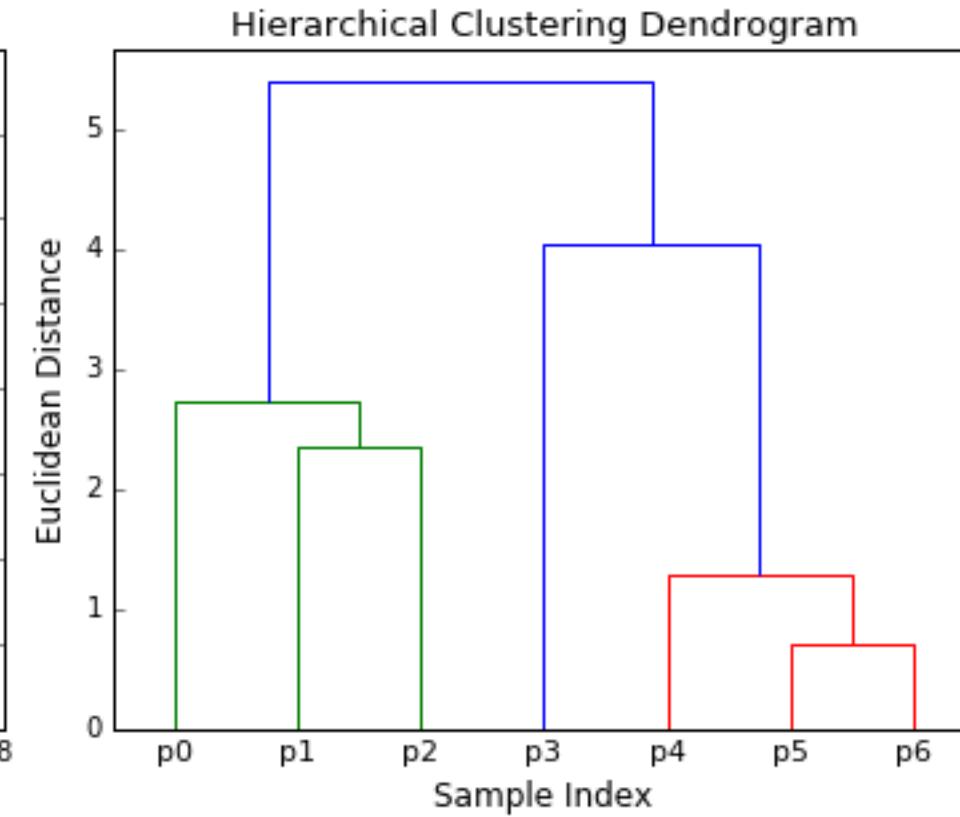
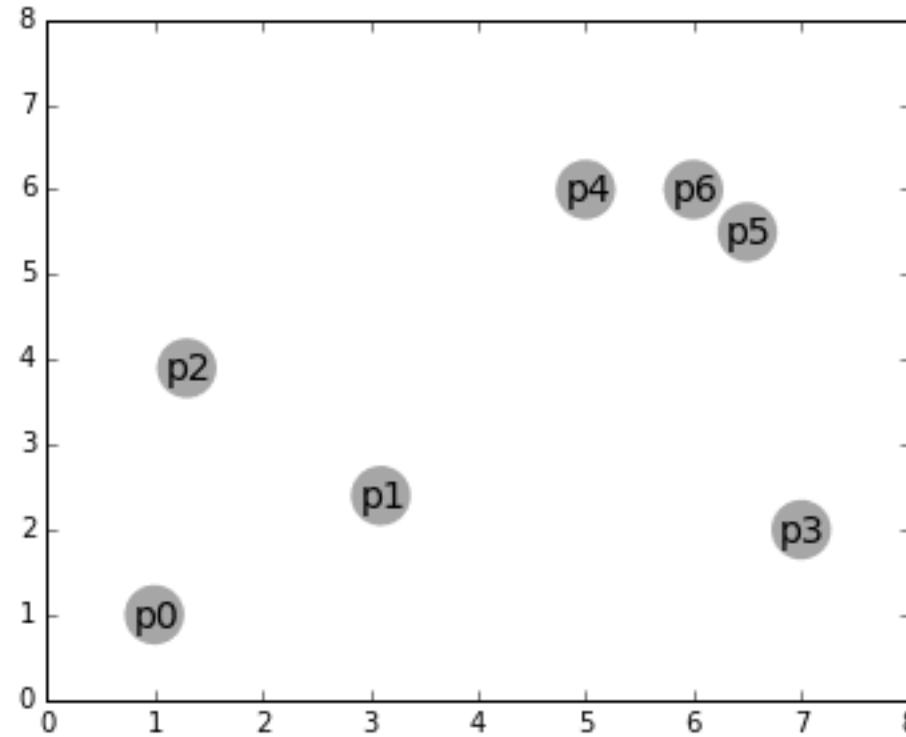
- Dendrogram
  - A tree that shows how clusters are merged/split hierarchically
  - Each node on the tree is a cluster; each leaf node is a singleton cluster
  - A clustering is obtained by cutting the dendrogram at the desired level (then each connected component forms a cluster)



# Hierarchical clustering



- Example





# Hierarchical clustering

- Three different distance measures
  - Single-nearest distance: single linkage
  - Complete-farthest distance: complete linkage
  - Average distance: average linkage

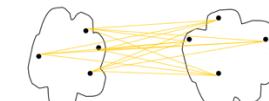
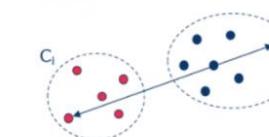
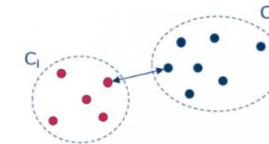
  

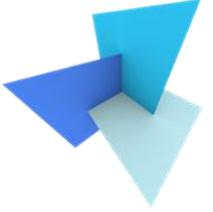
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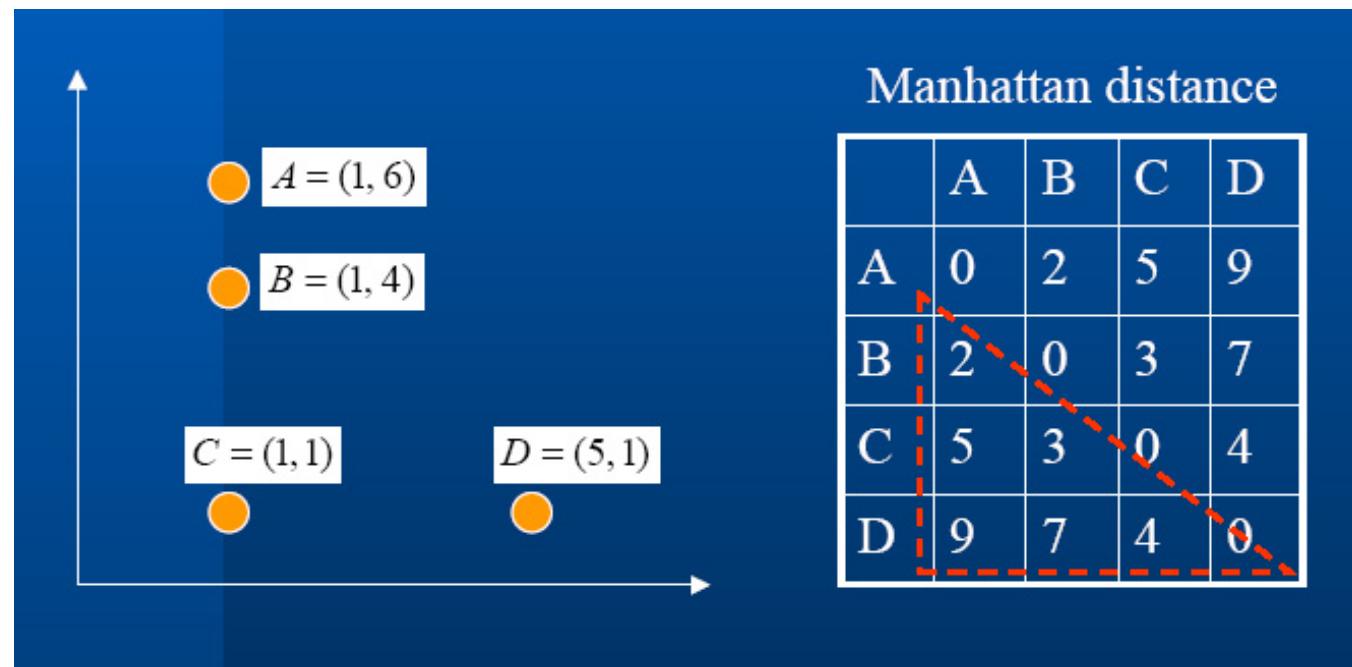
- Three different distance measures
  - Single-nearest distance (single linkage): **shortest** distance between any pair
$$D(c_i, c_j) = \min d(a, b), \forall a \in c_i \text{ and } b \in c_j$$
  - Complete-farthest distance (complete linkage): **greatest** distance between any pair
$$D(c_i, c_j) = \max d(a, b), \forall a \in c_i \text{ and } b \in c_j$$
  - Average distance or average linkage: **average** distance between all pairs

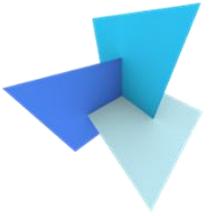




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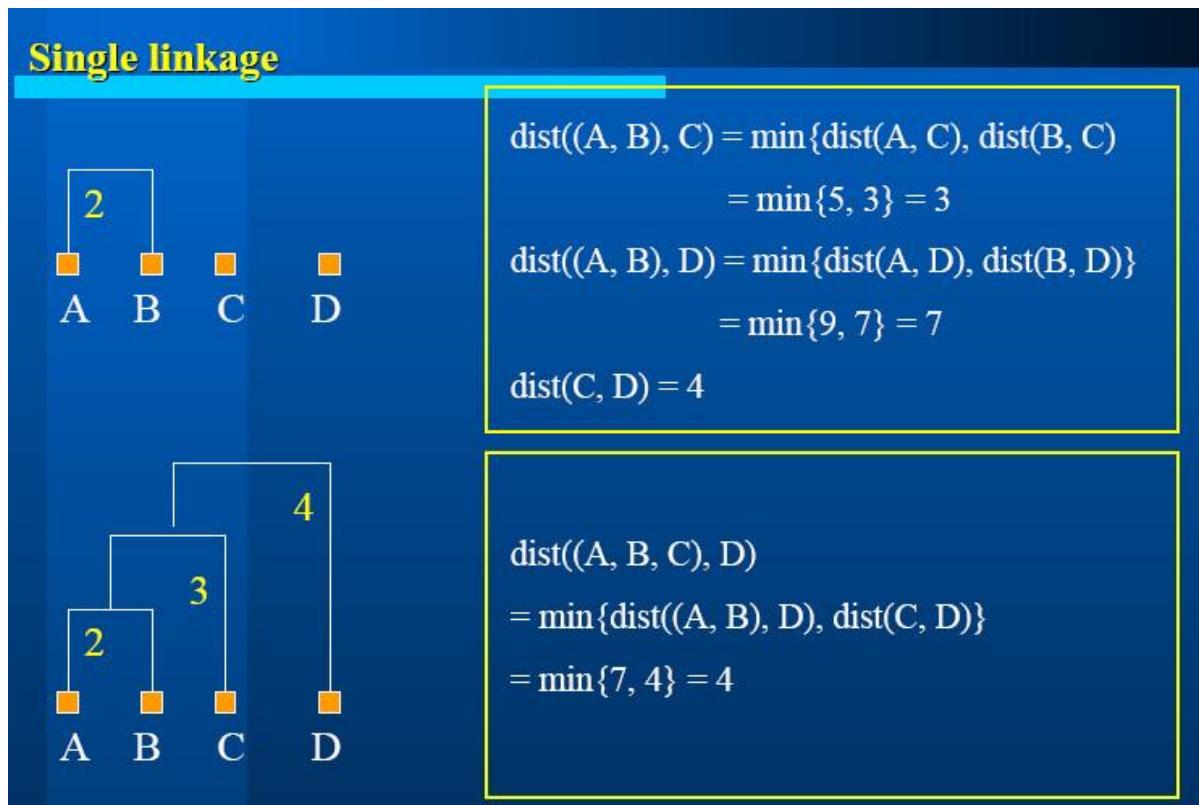
- Example: clustering 4 data items in 2D space



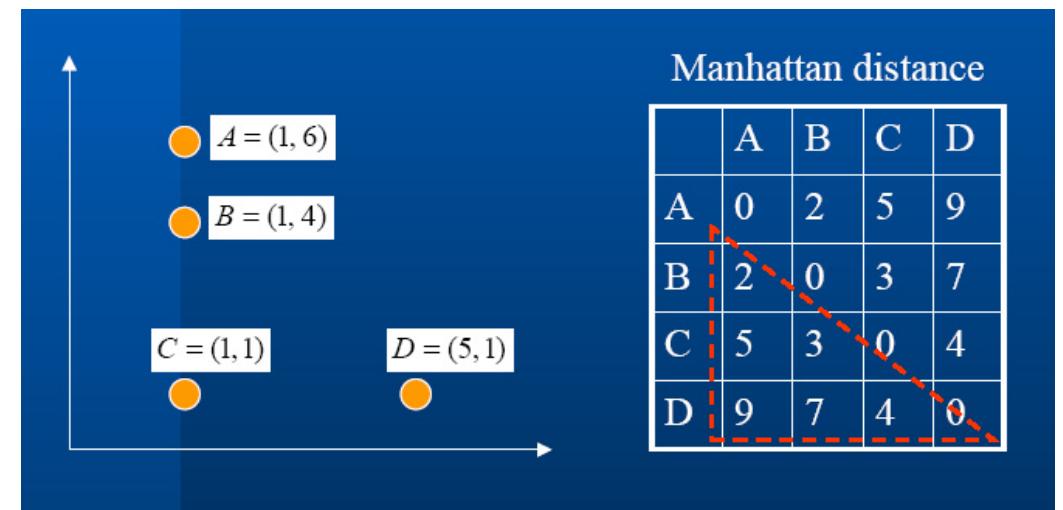


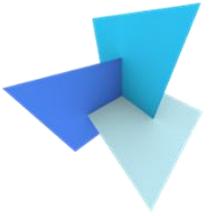
# Hierarchical clustering

- Method: *single-linkage* clustering



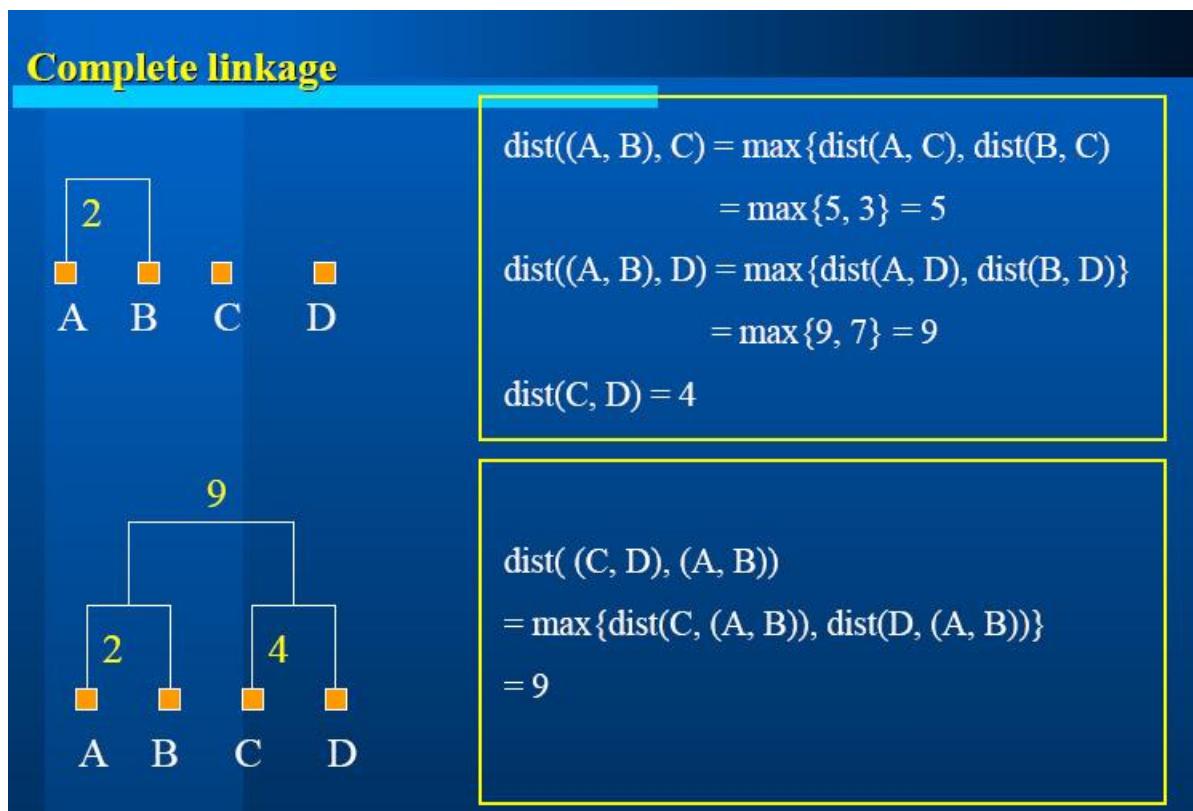
$$D(c_i, c_j) = \min d(a, b), \forall a \in c_i \text{ and } b \in c_j$$



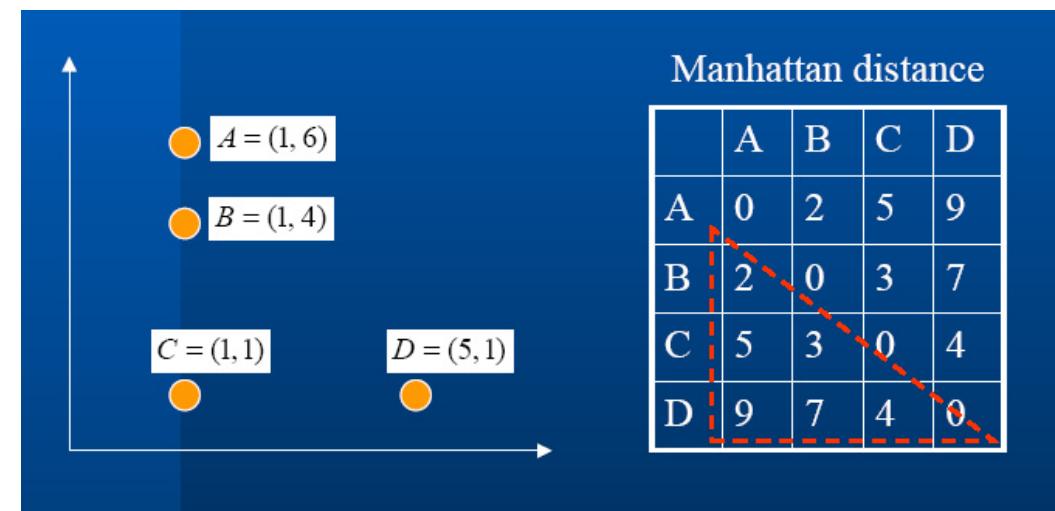


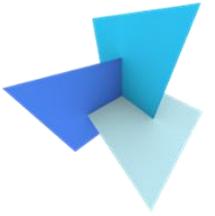
# Hierarchical clustering

- Method: *complete-linkage* clustering



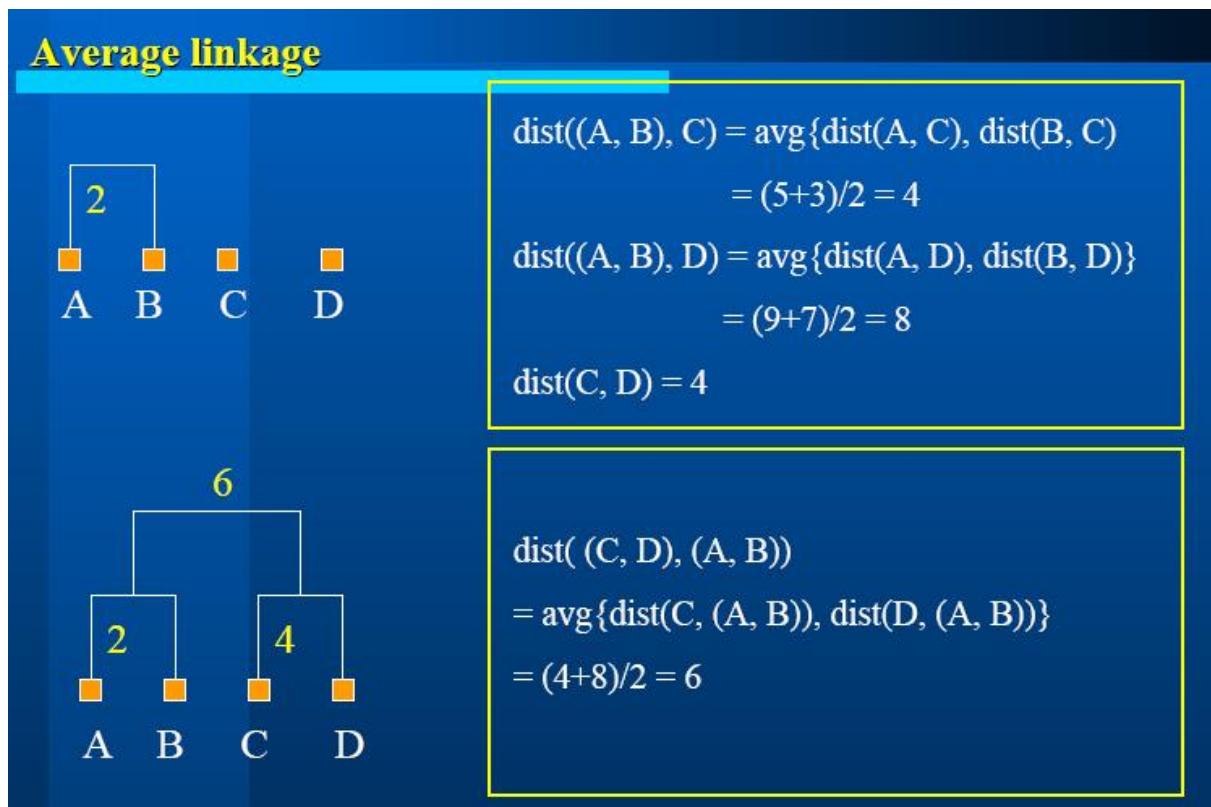
$$D(c_i, c_j) = \max d(a, b), \forall a \in c_i \text{ and } b \in c_j$$



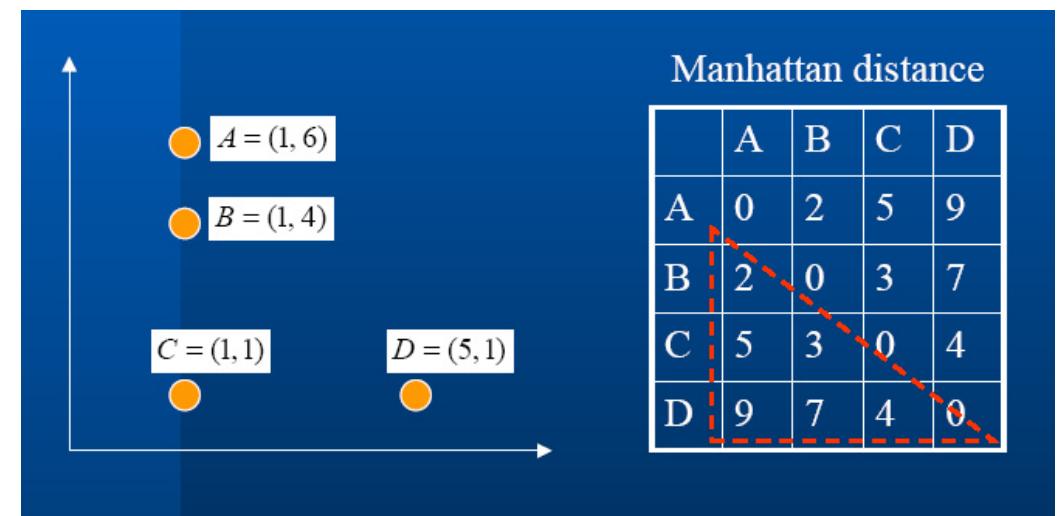


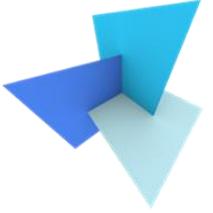
# Hierarchical clustering

- Method: *average-linkage* clustering



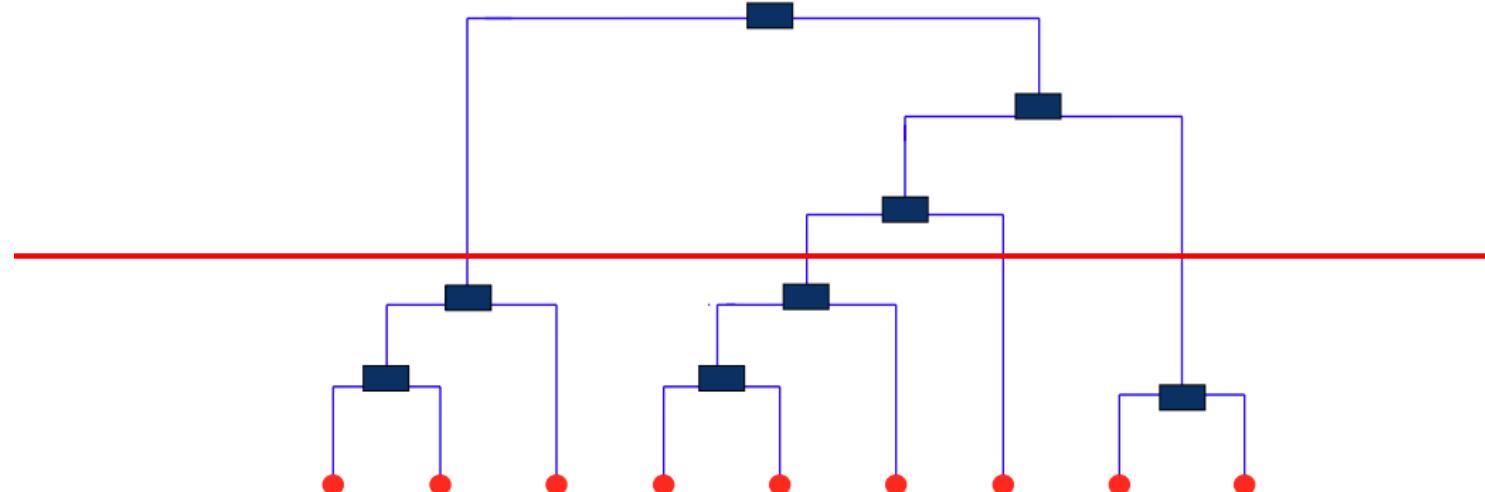
$$D(c_i, c_j) = \frac{1}{|c_i||c_j|} \sum_{a \in c_i, b \in c_j} d(a, b)$$





# Hierarchical clustering

- Advantages
  - No a priori information about the number of clusters required
    - Any desired number of clusters can be obtained by 'cutting' the dendrogram at the proper level
  - Easy to implement and gives best result in some cases





# Hierarchical clustering

- Advantages
  - No a priori information about the number of clusters required
  - Easy to implement and gives best result in some cases
- Limitations
  - Can never undo what (i.e., merging two clusters) was done previously
  - Can be slow if a large number data points (due to pairwise distance computation)
  - It may not be easy to choose a proper distance measure



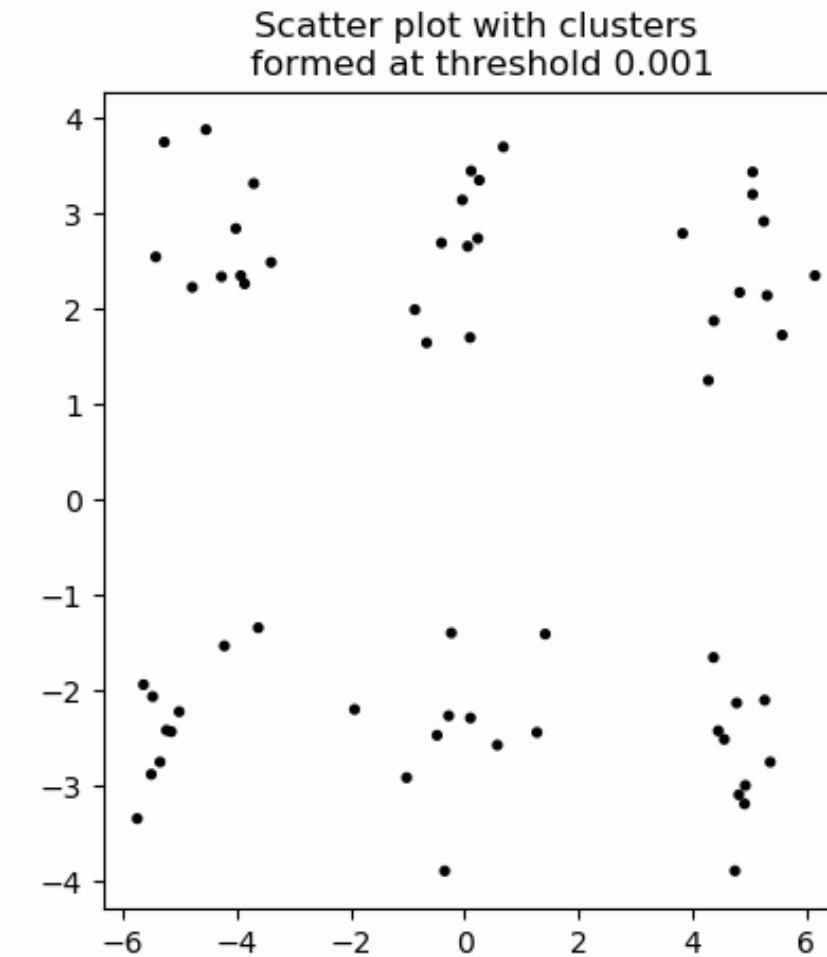
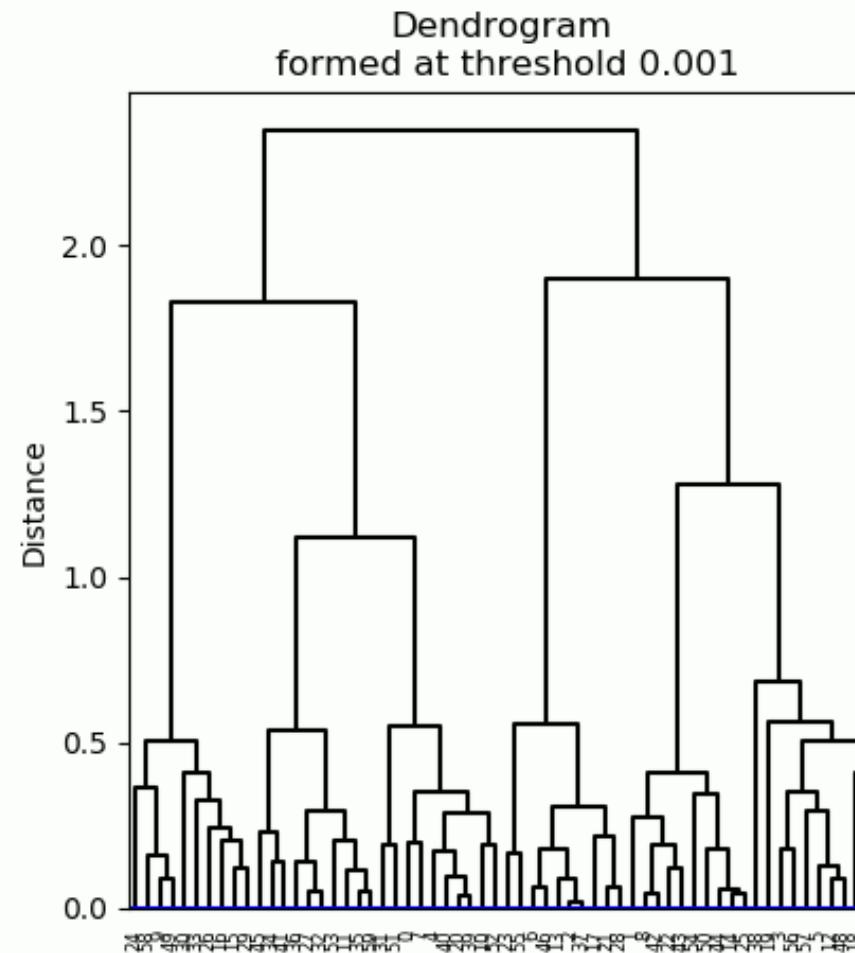
Single linkage: well-separated clusters



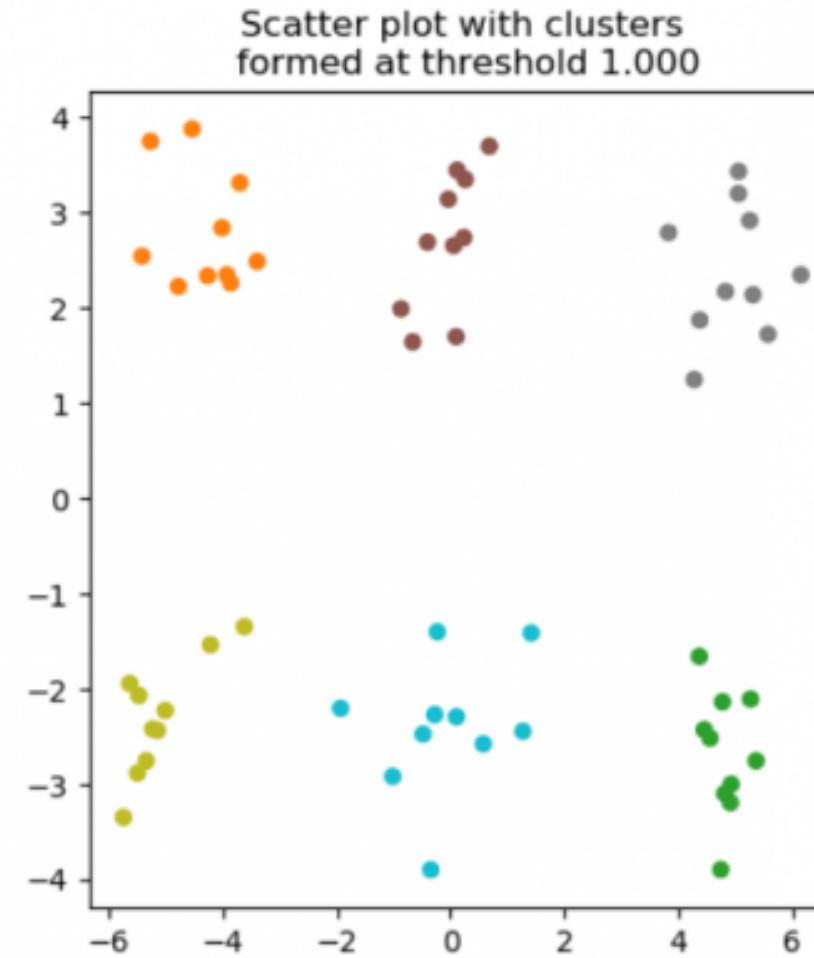
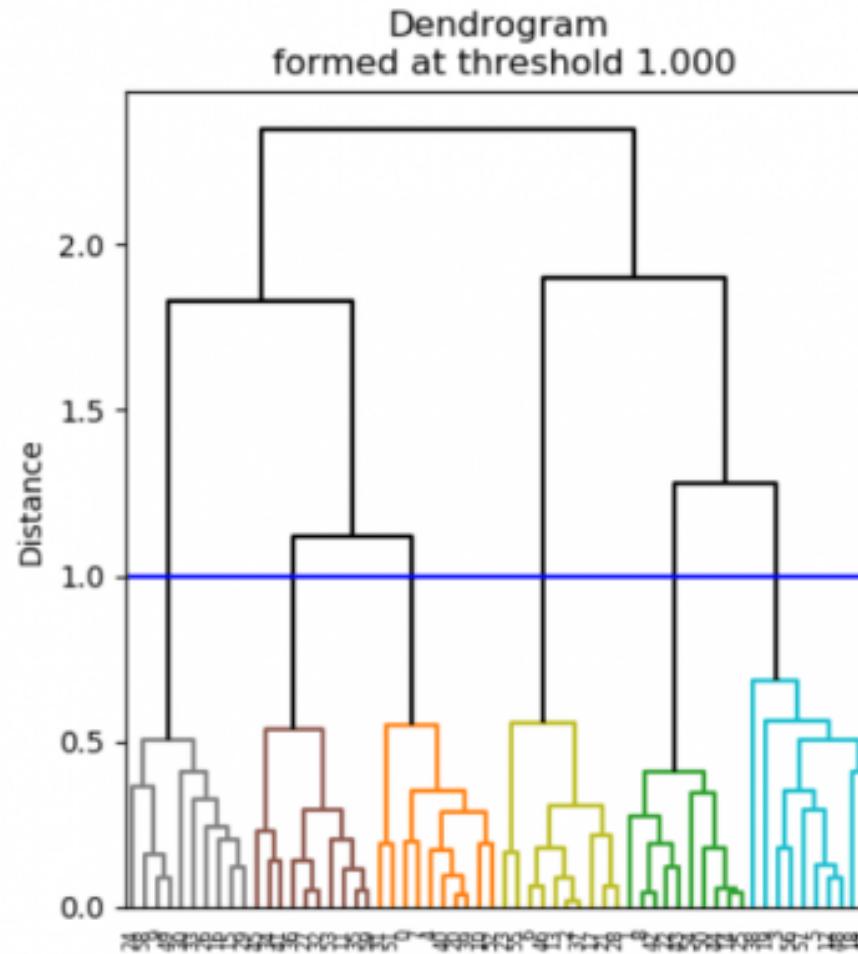
Complete linkage: compact clusters

- It may not be easy to identify the correct number of clusters by the dendrogram

# Hierarchical clustering



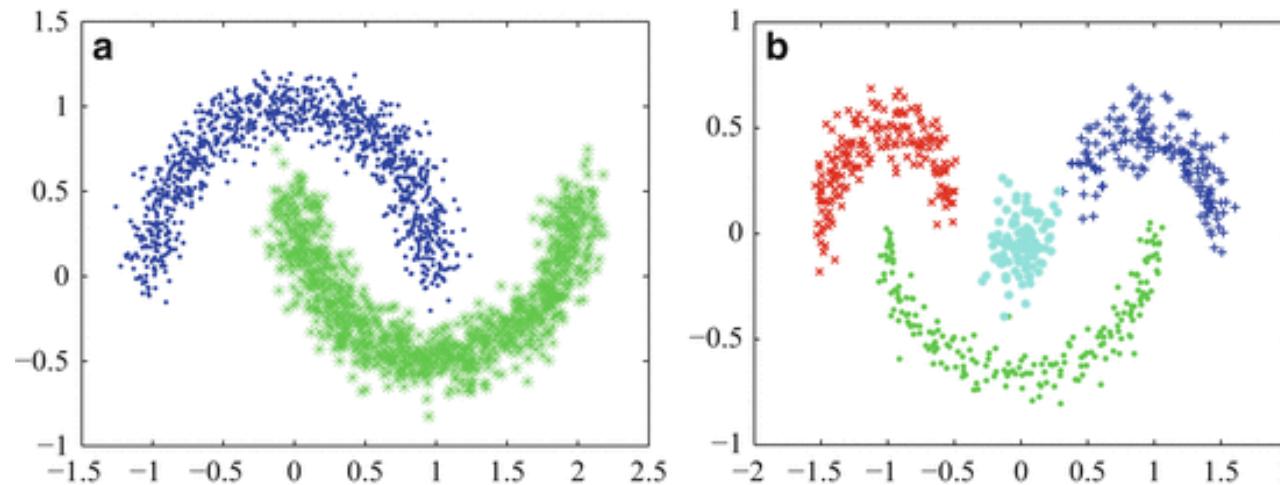
# Hierarchical clustering





# Hierarchical clustering

- Can hierarchical clustering method handle?

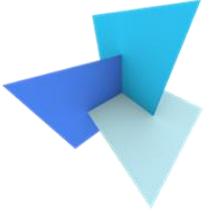




# Agenda

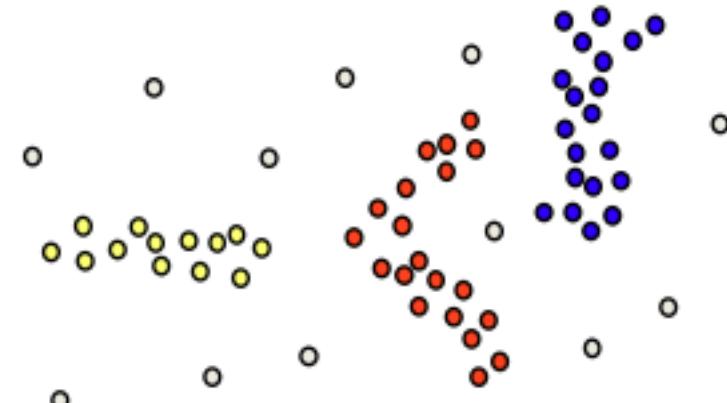
- Overview
  - What is clustering?
  - Distance measure (similarity measure)
  - Types of clustering algorithms
- Clustering algorithms
  - K-means clustering
  - Hierarchical clustering
  - Density-based clustering
- Nearest neighbor classification
- Features





# Density-based clustering

- Basic ideas
  - Clusters are contiguous regions of high density in the data space, separated by regions of lower data density
  - A cluster is defined as a maximal set of density connected points
- DBSCAN
  - Density-Based Spatial Clustering of Applications with Noise





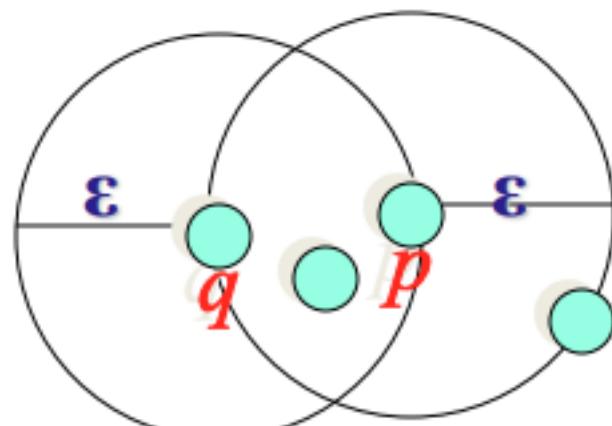
# Density definition: two parameters

- $\epsilon$ -neighborhood: objects within a radius of  $\epsilon$  from a cluster

$$N_\epsilon(p) : \{q \mid d(p, q) \leq \epsilon\}$$

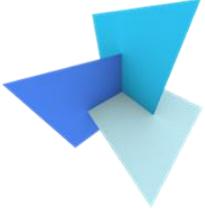
- The minimum number of points required to form a cluster

- High density:  $\epsilon$ -neighborhood of an object contains at least  $minPts$  of objects.

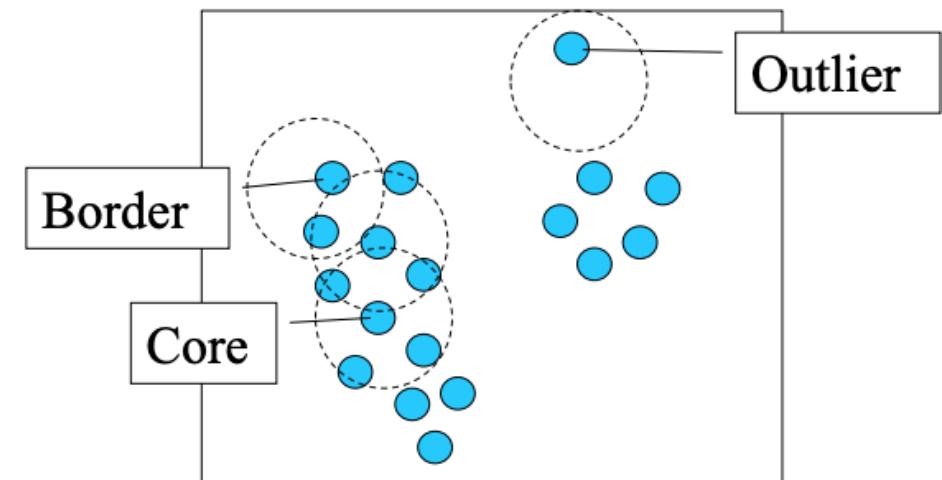


$\epsilon$ -neighborhood of  $p$  and  $q$

# Three types of data points



- Given  $\epsilon$  and  $minPts$ 
  - Core point: has at least  $minPts$  neighbors within its  $\epsilon$ -neighborhood
    - At the interior of a cluster
  - Border point
    - has fewer than  $minPts$  neighbors within its  $\epsilon$ -neighborhood
    - is within the  $\epsilon$ -neighborhood of a core point
  - Outlier/Noise
    - Any point that is neither core nor border



$\epsilon$  = circle radius,  $minPts$  = 5

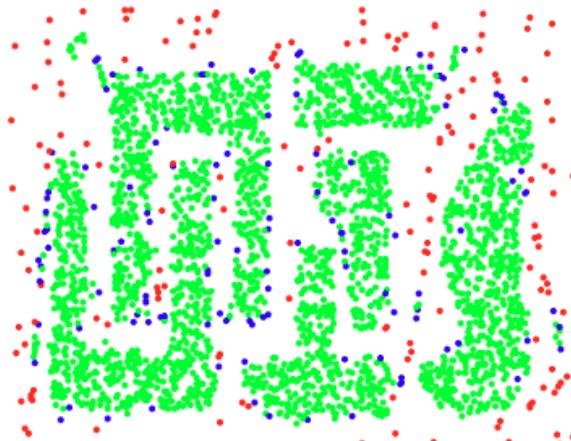


# Three types of data points

- Example



Original Points



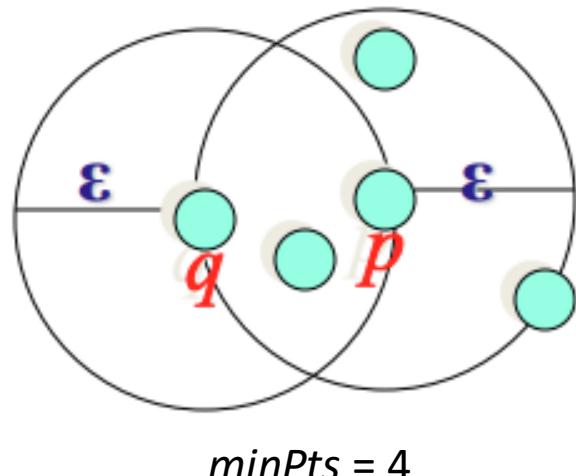
$\epsilon = 10, \ minPts = 4$

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



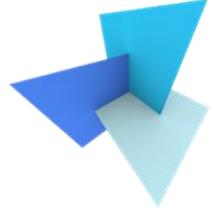
# Density definition: two concepts

- Density reachability
  - A point  $q$  is said to be directly reachable from a point  $p$  if
    - $p$  is a **core point** (i.e., has at least  $minPts$  points within  $\epsilon$ -neighborhood)
    - $q$  is within distance  $\epsilon$  from  $p$



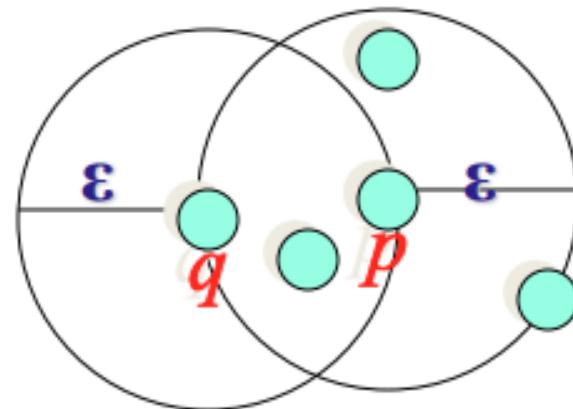
In this example,  $q$  is density reachable from  $p$ .  
Is  $p$  also density reachable from  $q$ ?



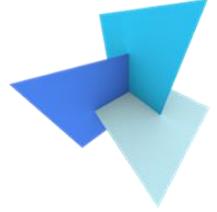


# Density definition: two concepts

- Density reachability
  - A point  $q$  is said to be directly reachable from a point  $p$  if
    - $p$  is a **core point** (i.e., has at least  $minPts$  points within  $\epsilon$ -neighborhood)
    - $q$  is within distance  $\epsilon$  from  $p$
  - Density reachability is asymmetric
    - Only core points can reach other points

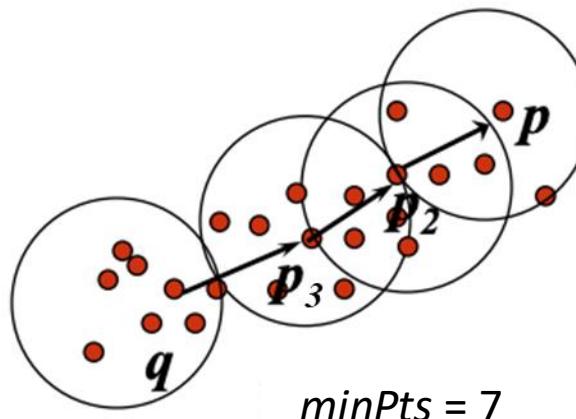


$minPts = 4$



# Density definition: two concepts

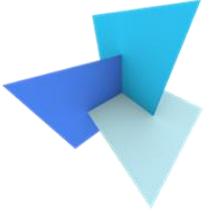
- Density reachability
- Density connectivity
  - A point  $p$  is said to be reachable from  $q$  if
    - There is a path  $p_1, \dots, p_n$  with  $p_1 = p$  and  $p_n = q$ , where each  $p_i$  is directly reachable from  $p_{i+1}$
  - Density connectivity is transitive (i.e., it forms a chain)



Example:

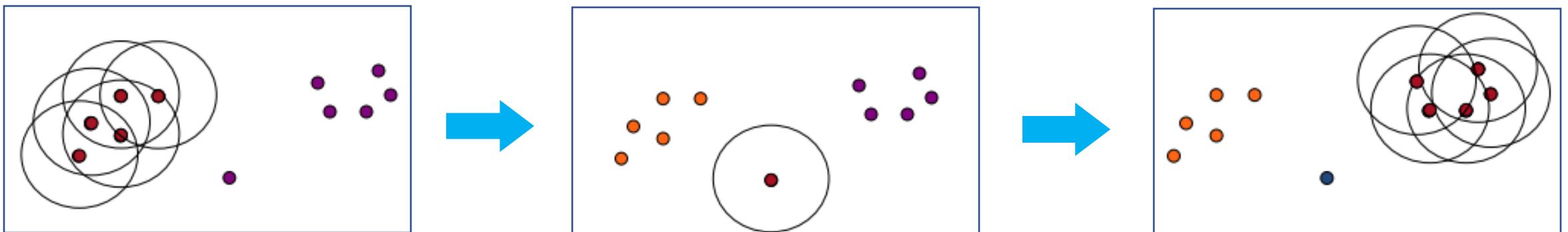
- $p$  is directly reachable from  $p_2$
- $p_2$  is directly reachable from  $p_3$
- $p_3$  is directly reachable from  $q$

So we say:  $p$  is reachable from  $q$  ( $p$  and  $q$  density connected)



# DBSCAN algorithm

```
for each  $o \in D$  do
  if  $o$  is not yet classified then
    if  $o$  is a core-object then
      collect all objects density-connected by  $o$ ,
      and assign them to a new cluster.
    else
      assign  $o$  to NOISE
```

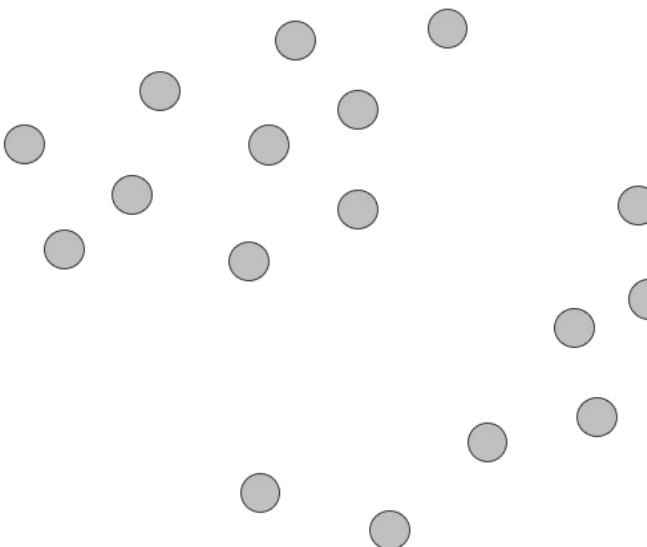


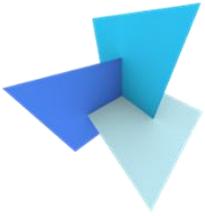
An example of DBSCAN clustering:  $minPts = 3$



# DBSCAN algorithm

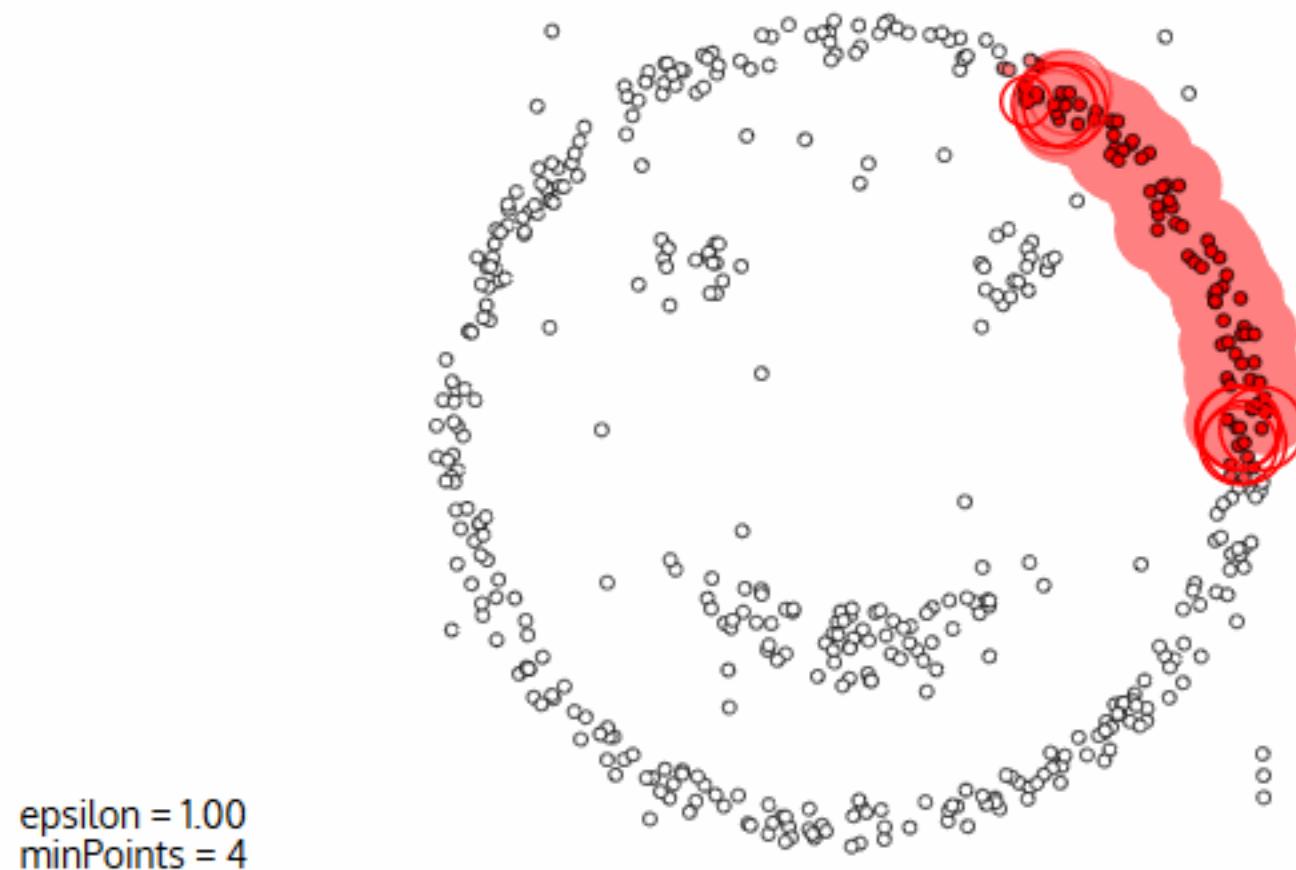
- Illustration

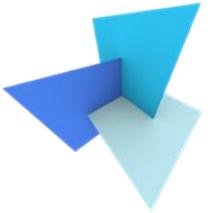




# DBSCAN algorithm

- Illustration



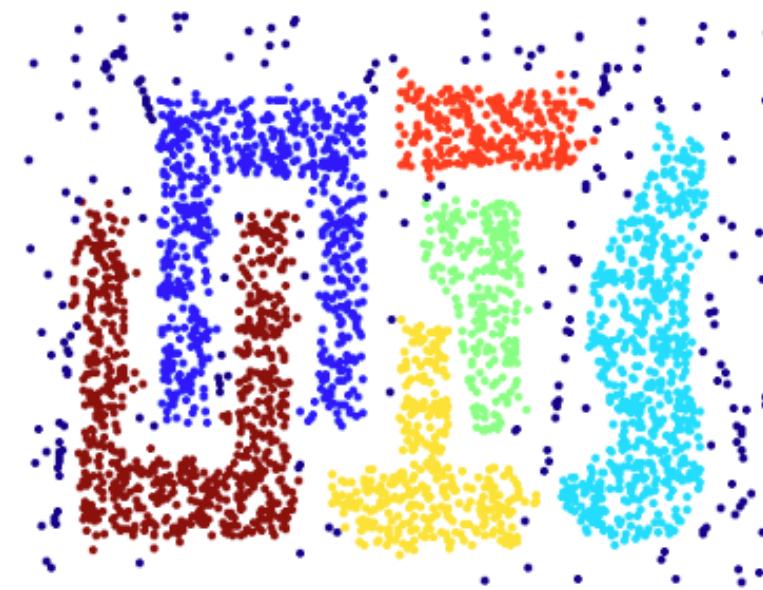


# DBSCAN algorithm

- Example



**Original Points**



**Clusters**



# DBSCAN algorithm

- Determining the two parameters

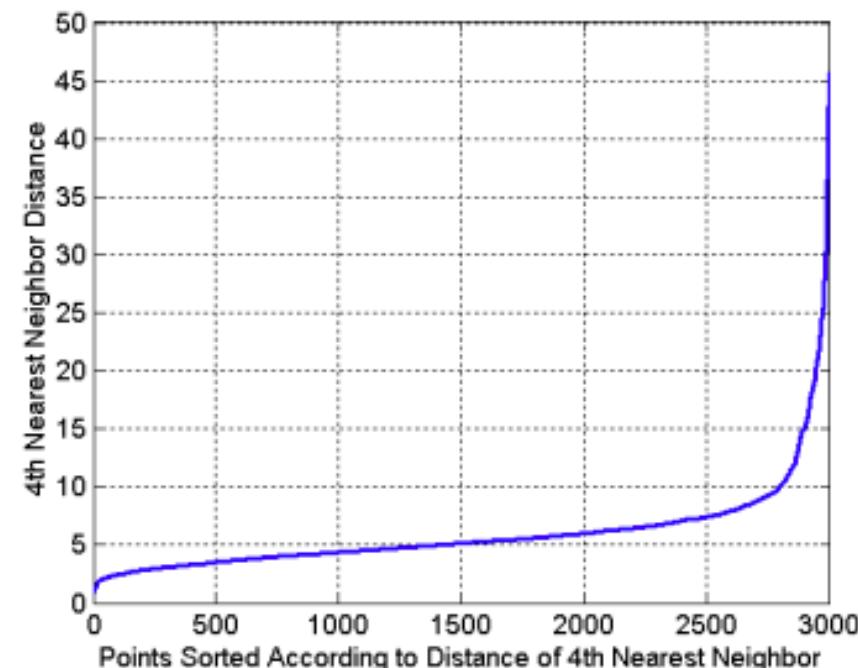
- $minPts$

- $minPts = 1$ ?
    - $minPts = 2$  (same as single linkage hierarchical method, with dendrogram cut at height  $\epsilon$ )
    - $minPts = 2 * \text{dimension}$

- $\epsilon$  (distance threshold)

- k-distance graph ( $k = minPts - 1$ )
    - Look for the “elbow”
      - The point with maximum curvature

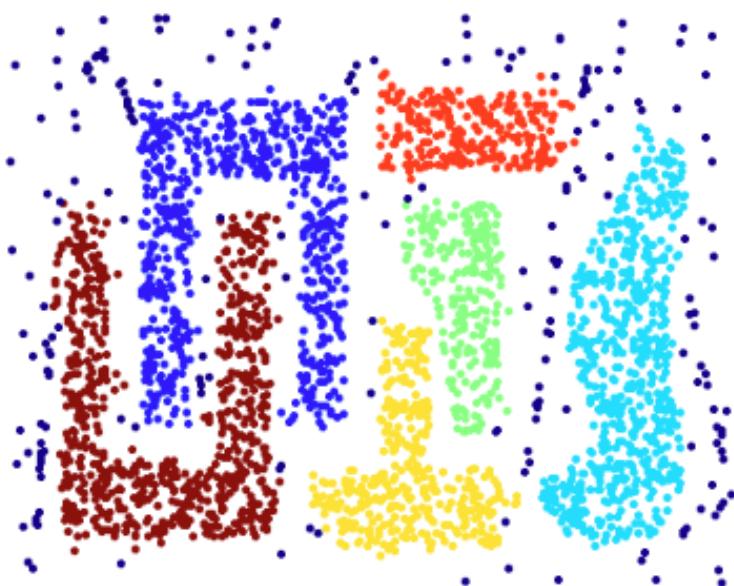
K-distance graph plots the k-th nearest neighbor distance of all points sorted from smallest to largest.

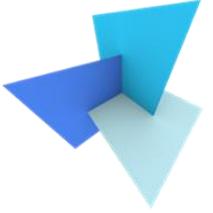




# DBSCAN algorithm

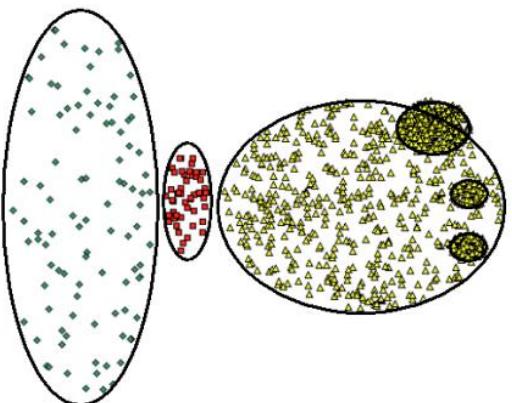
- Advantages
  - Resistant to Noise
  - Robust to clusters of different shapes and sizes



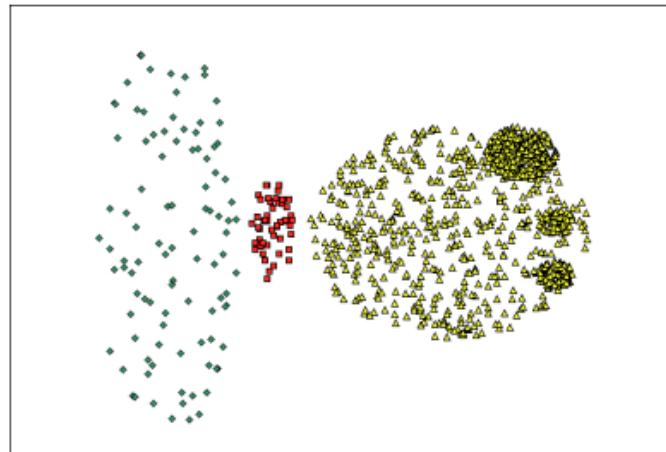


# DBSCAN algorithm

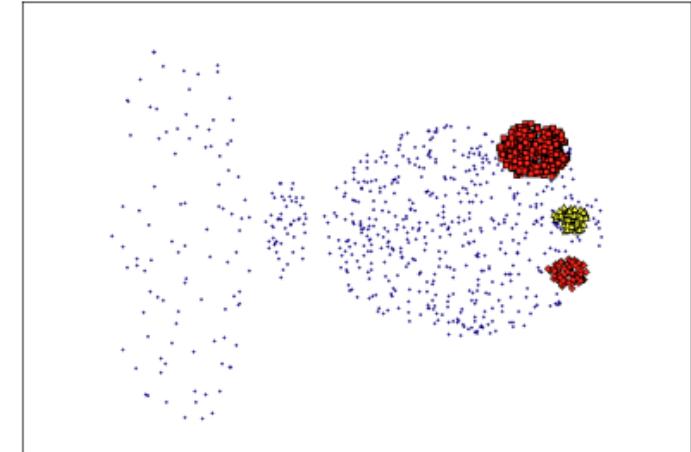
- Limitations
  - Cannot handle varying densities
  - Hard to determine a good set of parameters



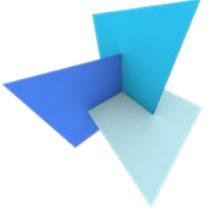
Original points



$minPts = 4, \epsilon = 9.92$



$minPts = 4, \epsilon = 75$

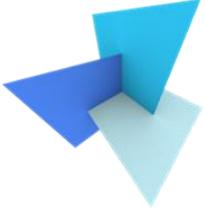


# Question

- Which method (DBSCAN or k-means) was used to produce each result?



# Agenda



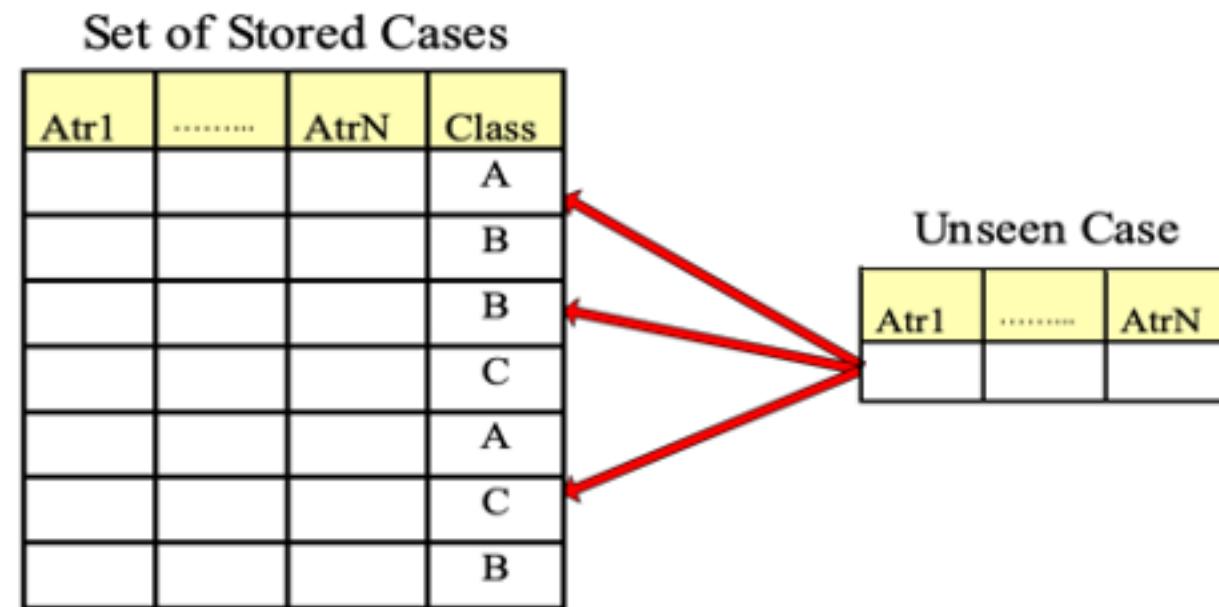
- Overview
  - What is clustering?
  - Distance measure (similarity measure)
  - Types of clustering algorithms
- Clustering algorithms
  - K-means clustering
  - Hierarchical clustering
  - Density-based clustering
- Nearest neighbor classification
- Features

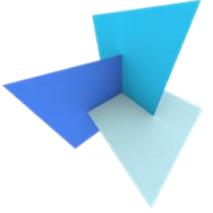




# Nearest neighbor classification

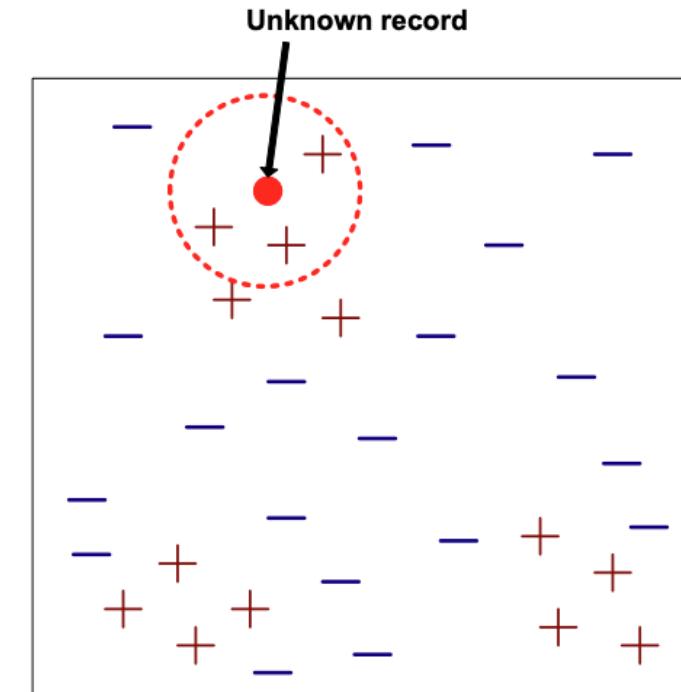
- Basic ideas
  - Store the training records
  - Use training records to predict the class label of unseen cases





# Nearest neighbor classification

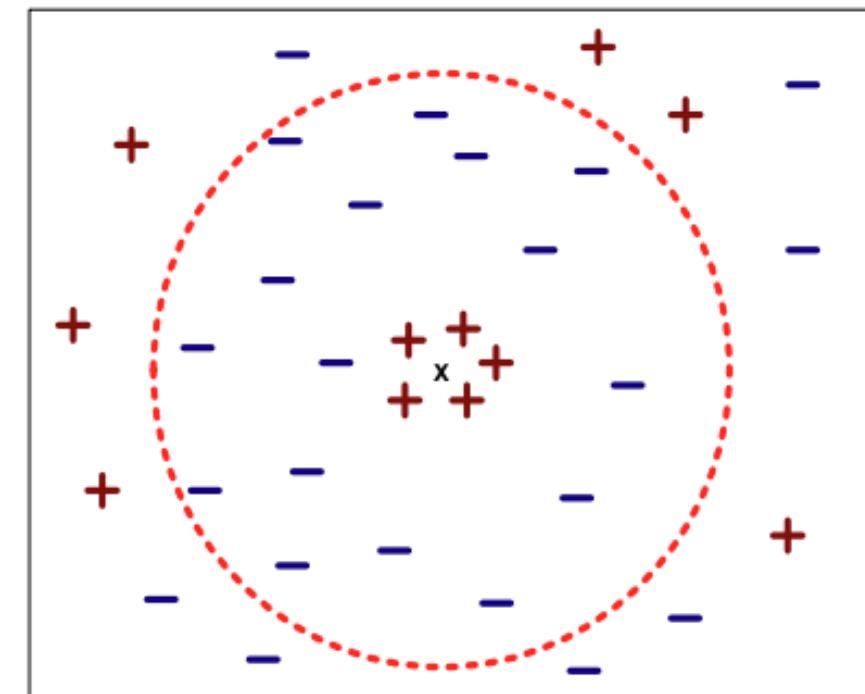
- Requires three things
  - The set of stored records
  - Distance metric to compute distance between records
  - The value of  $k$ , the number of nearest neighbors to retrieve
- To classify an unknown record
  - Compute distance to other training records
  - Identify  $k$  nearest neighbors
  - Use class labels of the  $k$  nearest neighbors to determine the class label of the unknown record (e.g., by taking majority vote)



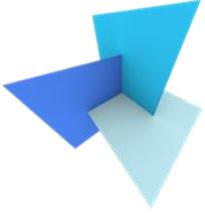


# Nearest neighbor classification

- Choosing the value of  $k$ 
  - If  $k$  is too small, sensitive to noise points
  - If  $k$  is too large, neighborhood may include points from other classes



# Agenda



- Overview
  - What is clustering?
  - Distance measure (similarity measure)
  - Types of clustering algorithms
- Clustering algorithms
  - K-means clustering
  - Hierarchical clustering
  - Density-based clustering
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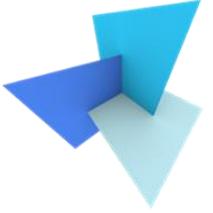




# Features

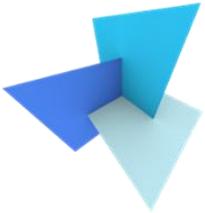
- A set of attributes of an object
- Typically stored as a vector – feature vector
- Scaling issue: distance measure dominated by one of the attributes
  - Example
    - height of a person [1.5m, 1.8m]
    - weight of a person [40kg, 100kg]
    - income of a person [€10K, €1M]
  - Solution
    - Normalization, i.e., 
$$\frac{\text{each attribute value}}{\text{max possible value of this attribute}}$$

$$d(\mathbf{p}, \mathbf{q}) = \sqrt{\sum_{i=1}^n (q_i - p_i)^2}$$



# You should have learned

- Clustering
  - The basic ideas, strengths, and weaknesses of the 3 clustering methods
  - K-means
    - How is K-means interpreted as an optimization problem?
  - Hierarchical clustering
    - Several ways of defining inter-cluster distance
  - Density-based clustering
    - The parameters and the definitions of neighborhood and density in DBSCAN
- Classification
  - The basic idea of k-nearest neighbor classifier



# Next Lecture

- Bayesian classification & logistic regression

