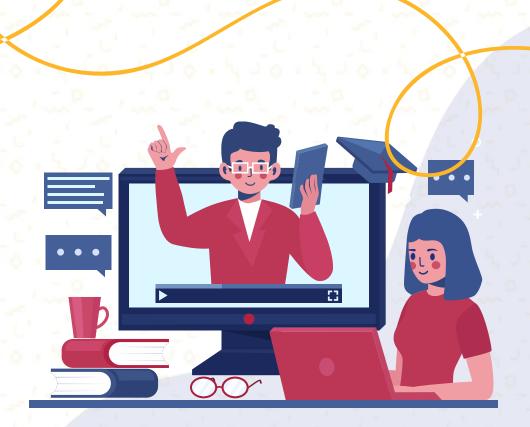






# **Table of Content What will We Learn Today?**

- 1. DBSCAN
- 2. Hierarchical Clustering

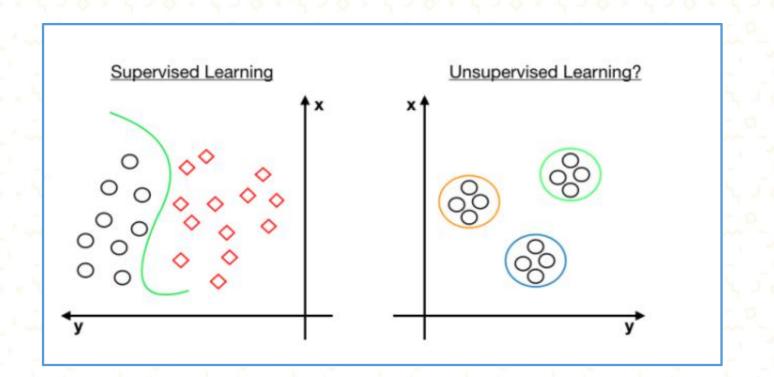






# Supervised vs Unsupervised

- Supervised = Learn to predict the outcome.
  - We know the target label, so we make the model that try to predict the label.
- Unsupervised = Finding pattern/ characteristic from data.
  - · We do not know our target label, so we make model that try to group the data.









#### Types of clustering algorithms

- Connectivity models
  - Based on the notion that the data points closer in data space exhibit more similarity to each other than the data points lying farther away. Example: hierarchical clustering algorithm.
- Centroid models:
  - The notion of similarity is derived by the closeness of a data point to the centroid of the clusters. Example: K-means
- Distribution models
  - The notion of how probable is it that all data points in the cluster belong to the same distribution (For example: Normal, Gaussian).
- Density models
  - These models search the data space for areas of varied density of data points in the data space. Example: DBSCAN





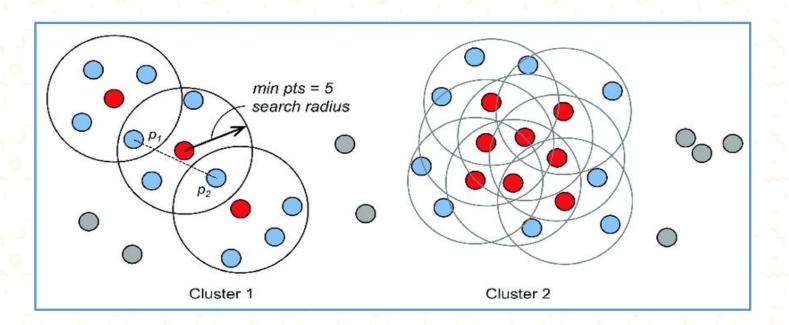
# **DBSCAN**







- Density-Based Spatial Clustering of Applications with Noise (DBSCAN) is
  - algorithm for density-based clustering
  - proposed by Martin Ester, Hans-Peter Kriegel, Jörg Sander and Xiaowei Xu in 1996

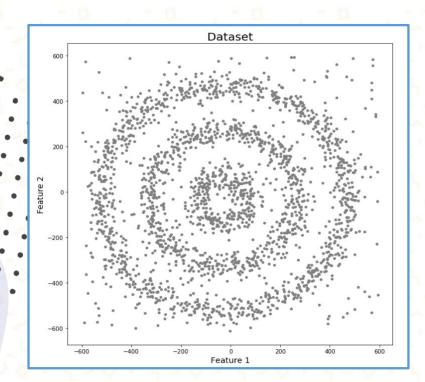


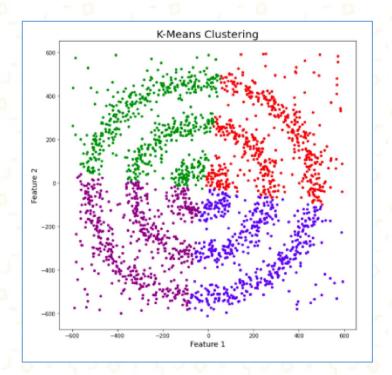


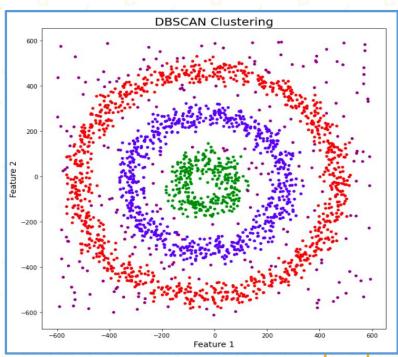




# Why DBSCAN?







DBSCAN can cluster the data points correctly, and also detects noise





# **How DBSCAN works?**

- Group objects in dense region
- Major features
  - Discover clusters of arbitrary shape
  - Handle noise
  - One scan
  - Need density parameters as termination condition
- Density parameters
  - **Radius**  $\varepsilon$ : distance to determine the neighborhood
  - MinPts: Minimum number of points in neighborhood

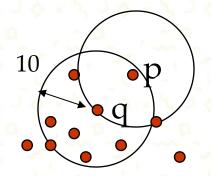






#### **Definitions**

- Core object
  - $\circ$   $\varepsilon$ -neighborhood contains *MinPts* objects
- Directly density-reachable
  - p is directly density-reachable from q if
     q is a core object, and p is ε-neighborhood of q



$$\varepsilon = 10$$
  
MinPts = 5

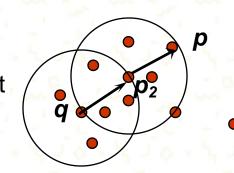


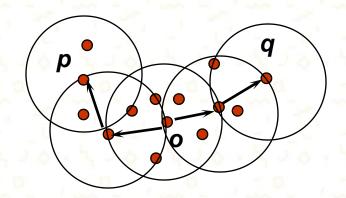




#### **Definitions**

- Density-reachable
  - p is density-reachable from q if
     there are objects p<sub>1</sub>, p<sub>2</sub>, ... p<sub>n</sub>, p<sub>1</sub> = q, p<sub>n</sub> = p such that
     p<sub>i+1</sub> is directly density-reachable from p<sub>i</sub>
- Density-connected
  - p is density-connected to q if there is an object o such that
    p and q are density-reachable from o











#### **DBSCAN**

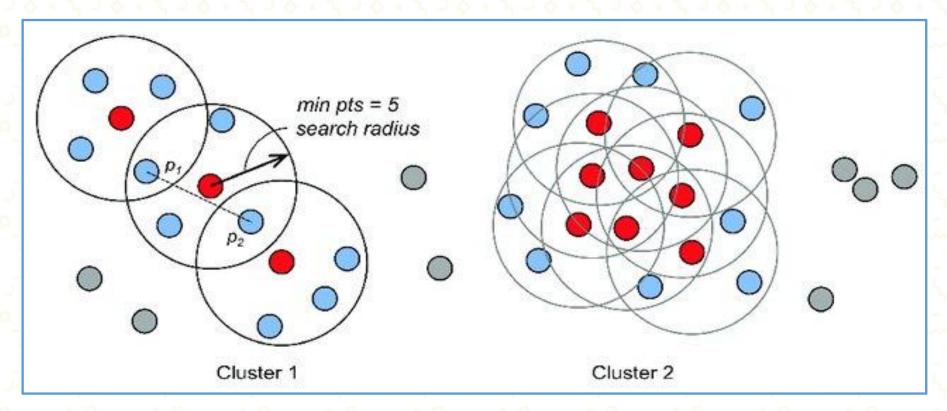
- A cluster a maximal set of density-connected points
- Discovers clusters of arbitrary shape in databases with noise
  - 1. Arbitrary select a point **p**
  - 2. Retrieve all  $\varepsilon$ -neighborhood of  $\boldsymbol{p}$
  - 3. If **p** is a core object, a cluster is formed
  - 4. From each core object **p**, iteratively collects directly density-reachable objects (may merge clusters)
  - 5. Continue the process until no new points can be added
- Problem with DBSCAN
  - Selecting parameters ε and MinPts







#### **DBSCAN**



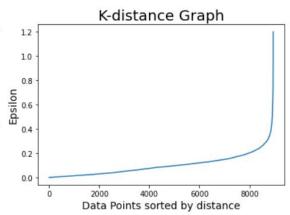
- There are 3 types of point
  - core point (red)
  - border points (blue)
  - noise points (grey)





#### How to determine MinPts and eps

- MinPts
  - Using domain knowledge
  - The larger the data set, the larger the value of MinPts should be
  - If the data set is noisier, choose a larger value of MinPts
  - Generally, MinPts should be greater than or equal to the dimensionality of the data set
  - For 2-dimensional data, use DBSCAN's default value of MinPts = 4 (Ester et al., 1996).
  - If your data has more than 2 dimensions, choose MinPts = 2\*dim, where dim= the dimensions of your data set (Sander et al., 1998).
- Epsilon
  - sorted k-dist graph









# **Hierarchical Clustering**

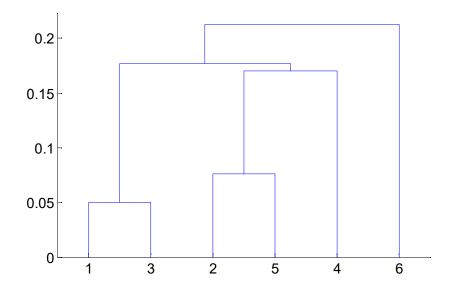






# Hierarchical clustering

- Produces a set of nested clusters organized as a hierarchical tree
- Can be visualized as a dendrogram
  - A tree like diagram that records the sequences of merges or splits









# Strengths of Hierarchical Clustering

- Do not have to assume any particular number of clusters
  - Any desired number of clusters can be obtained by 'cutting' the dendogram at the proper level
- They may correspond to meaningful taxonomies
  - Example in biological sciences (e.g., phylogeny reconstruction, ...)

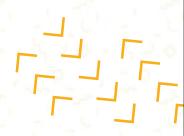






### Hierarchical clustering

- Two main types of hierarchical clustering
  - Agglomerative (bottom-up):
    - 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 (top-down):
    - Start with one, all-inclusive cluster
    - At each step, split a cluster until each cluster contains a point (or there are k clusters)
- Traditional hierarchical algorithms use a similarity or distance matrix
  - Merge or split one cluster at a time







# **Agglomerative Clustering Algorithm**

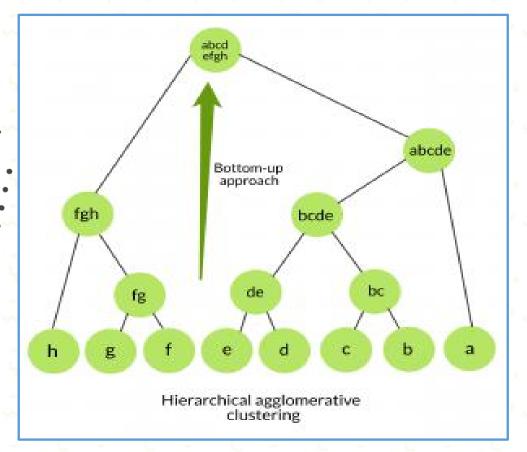
- More popular hierarchical clustering technique
- Basic algorithm is straightforward
  - 1. Compute the proximity matrix
  - 2. Let each data point be a cluster
  - 3. Repeat
  - 4. Merge the two closest clusters
  - 5. Update the proximity matrix
  - **6. Until** only a single cluster remains
- Key operation is the computation of the proximity of two clusters
  - Different approaches to defining the distance between clusters distinguish the different algorithms

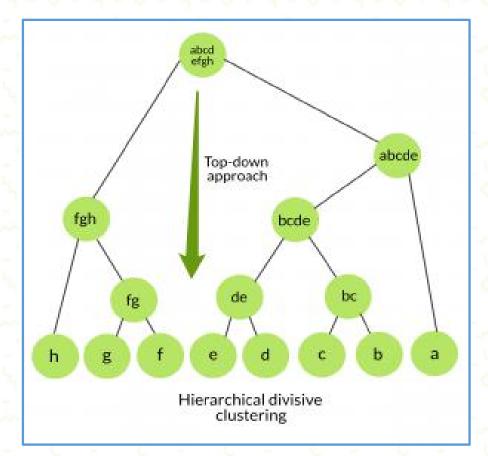






# Hierarchical clustering







https://www.geeksforgeeks.org/ml-hierarchical-clustering-agglomerative-and-divisive-clustering/





# **Lets Coding!**





# Thank YOU

