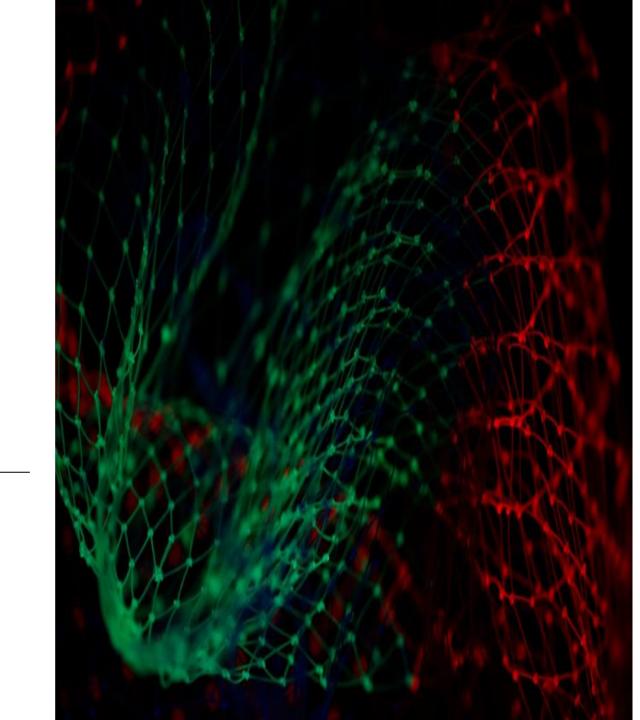
# Machine Learning

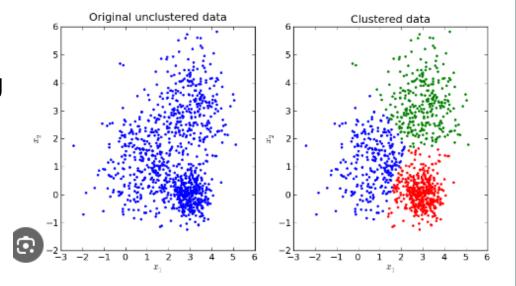
#### **CLUSTERING ALGORITHMS**

KUPPUSAMY S



## 1.K-means Clustering

- K-means clustering is a partitioning method that divides a dataset into K distinct, non-overlapping subsets (clusters).
- Clustering Approach- Partitional (flat)
- The algorithm aims to minimize the variance within each cluster.





- Select K initial cluster centroids randomly.
- Assign each data point to the nearest centroid, forming K clusters.
- Recalculate the centroids of each cluster based on the assigned points.
- Repeat the assignment and centroid update steps until convergence (no change in centroids or minimal change)



#### **Advantages:**

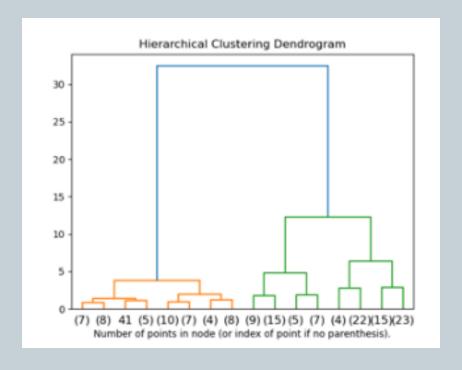
- Simple and easy to implement.
- Efficient for large datasets.
- Works well with spherical-shaped clusters.

- Requires specifying the number of clusters (K) in advance.
- Sensitive to initial centroid placement.
- Not suitable for clusters with varying sizes or densities.



## 2.Agglomerative Clustering

- Agglomerative clustering is a type of hierarchical clustering that builds nested clusters by merging pairs of data points successively.
- The process continues until all points belong to a single cluster or a stopping criterion is met.





- Start with each data point as its own cluster.
- Merge the closest pair of clusters based on a distance metric (e.g., Euclidean distance).
- Update the distance matrix and repeat until a single cluster remains or a predefined number of clusters is reached.



#### **Advantages:**

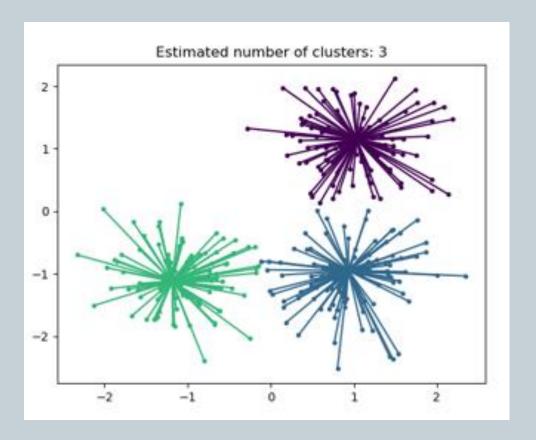
- Does not require specifying the number of clusters in advance.
- •Captures a hierarchy of clusters, useful for understanding data structure.

- Computationally expensive for large datasets.
- •Sensitive to the choice of distance metric and linkage criterion.
- •Once a merge is made, it cannot be undone.



# 3. Affinity Propagation Clustering

- Affinity Propagation is a clustering algorithm that identifies exemplars (representative points) by exchanging messages between data points.
- It does not require specifying the number of clusters beforehand.





- Each data point sends messages to all other points indicating how suitable they are as exemplars.
- Messages are updated iteratively based on "responsibility" and "availability" measures.
- Clusters are formed around points with the highest responsibility-availability scores.



#### **Advantages:**

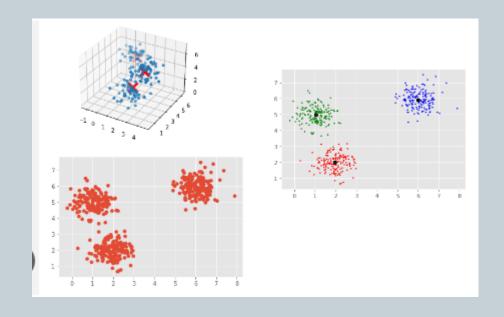
- Automatically determines the number of clusters.
- Can find clusters of varying sizes and densities.

- Computationally expensive for large datasets.
- •Sensitive to the choice of preference parameter.



# 4.MeanShift Clustering

- •MeanShift is a non-parametric, densitybased clustering algorithm that does not require specifying the number of clusters.
- •It works by shifting points towards areas of higher data density (modes of the density function).





- •Place a window over each data point and compute the mean of points within the window.
- Shift the window to the mean and repeat this process until convergence.
- •Points that converge to the same mode are assigned to the same cluster.



#### **Advantages:**

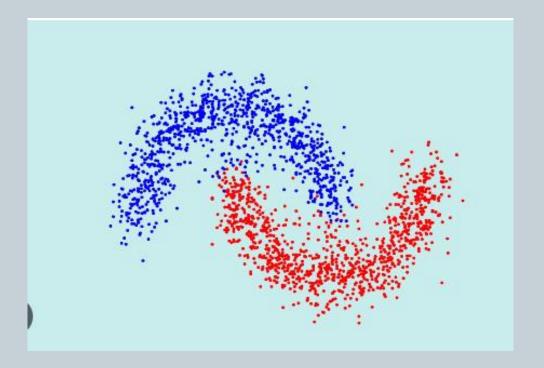
- •Does not require pre-specifying the number of clusters.
- •Can detect clusters of arbitrary shapes and sizes.

- Computationally expensive, especially for large datasets.
- •Requires selecting a suitable bandwidth parameter for the window, which can be challenging.



# **5.**Spectral Clustering

- Spectral Clustering is a graph-based algorithm that uses the eigenvalues of a similarity matrix to reduce dimensionality before clustering.
- It is effective for detecting nonconvex clusters.





- Construct a similarity graph based on the data points.
- Compute the Laplacian matrix of the graph.
- •Perform eigenvalue decomposition and select the top eigenvectors.
- Apply K-means clustering on the selected eigenvectors to form clusters.



#### Advantages:

- Suitable for non-convex clusters.
- Can handle complex data distributions.

- Computationally expensive for large datasets.
- •Requires a similarity matrix, which can be difficult to define.

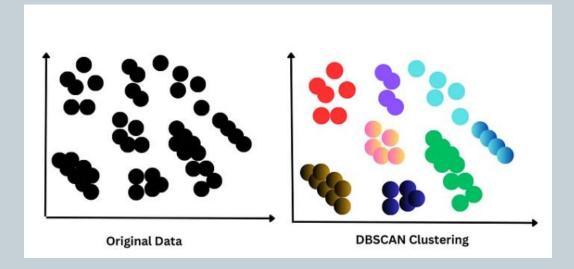


## **6.DBSCAN Clustering**

#### Definition:

•DBSCAN (Density-Based Spatial Clustering of Applications with Noise) is a density-based algorithm that clusters points based on the density of their neighborhood.

•It can identify arbitrarily shaped clusters and outliers.





- •Define core points with at least min\_samples neighbors within eps distance.
- •Expand clusters from core points by adding density-reachable points.
- •Mark points not reachable by any core point as noise.



#### Advantages:

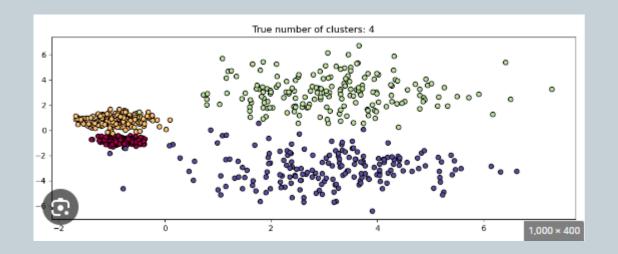
- Does not require specifying the number of clusters.
- Can find arbitrarily shaped clusters and detect outliers.

- Sensitive to parameter settings (eps and min\_samples).
- Not suitable for datasets with varying densities.



### 7.HDBSCAN Clustering

- •HDBSCAN (Hierarchical DBSCAN) extends DBSCAN by converting it into a hierarchical clustering algorithm.
- •It allows for varying cluster densities and provides a hierarchy of clusters.





- •Construct a minimum spanning tree of the distance-weighted data points.
- •Generate a hierarchy of clusters by varying the density threshold.
- •Extract clusters using a stability measure to find the most persistent clusters.



#### Advantages:

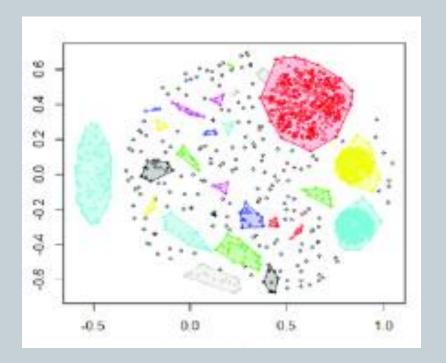
- No need to specify the number of clusters.
- •Handles varying cluster densities and noise.

- Computationally more expensive than DBSCAN.
- Parameter tuning can be complex.



# **8.OPTICS Clustering**

- •OPTICS (Ordering Points To Identify the Clustering Structure) is a density-based clustering algorithm that extends DBSCAN to handle varying densities.
- •It produces an ordering of the points that reflects their density-based clustering structure.





- •The algorithm orders data points based on their density reachability distance.
- A reachability plot is generated, and clusters are identified by valleys in the plot.
- •Points with similar densities form clusters, while points with no neighbors are considered noise.



#### **Advantages:**

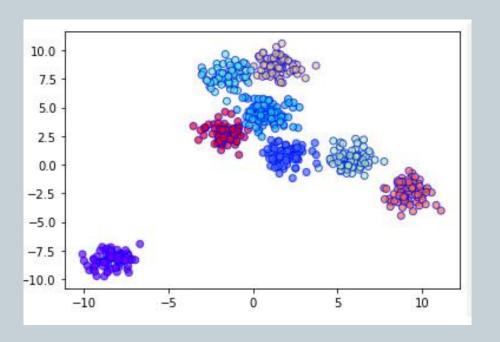
- •Handles clusters of varying densities well.
- •Does not require specifying the number of clusters.
- •Can identify nested clusters and noise points.

- Computationally expensive compared to DBSCAN.
- •Interpreting reachability plots can be complex.



# 9.BIRCH Clustering

- •BIRCH (Balanced Iterative Reducing and Clustering using Hierarchies) is a hierarchical clustering algorithm designed for large datasets.
- •It builds a Clustering Feature tree structure (CF tree) to summarize the data, and clusters are formed by traversing this structure.





- •Build a CF tree, where each node represents a cluster of data points summarized by their centroid.
- Points are inserted into the tree, and sub-clusters are formed dynamically.
- •Clusters are refined during the global clustering phase, using methods like K-means or agglomerative clustering.



#### **Advantages:**

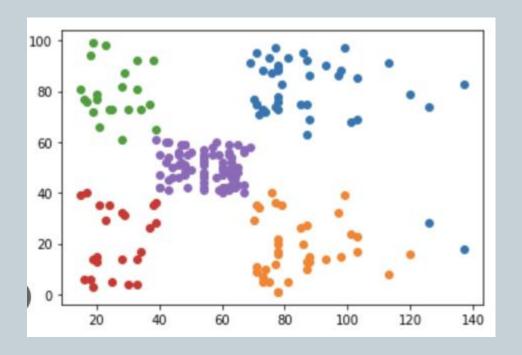
- Scalable to large datasets.
- Can incrementally process new data points.
- •Handles noise and outliers effectively.

- Performance can degrade with high-dimensional data.
- •The structure of the CF tree depends heavily on the input order of data.



## 10.BisectingKMeans Clustering

- •Bisecting K-means is a hierarchical (top-down) clustering algorithm that repeatedly splits clusters using K-means.
- •It combines elements of both divisive hierarchical clustering and K-means partitioning.





- •Start with all data points in a single cluster.
- •At each iteration, the current cluster is bisected using K-means to form two sub-clusters.
- •The bisection that minimizes intra-cluster variance is chosen, and this process continues until the desired number of clusters is reached.



#### Advantages:

- Often more efficient and scalable than standard K-means.
- •Allows control over the number of splits and clusters.

- Sensitive to initial cluster centroids.
- •May result in suboptimal splits if poor bisections are made early.

