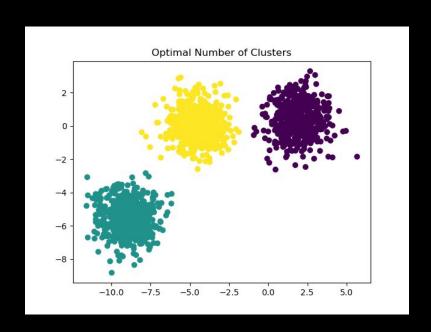
# K-Means Clustering: Unveiling Hidden Patterns



# Introduction to Unsupervised Learning

What is Unsupervised Learning?

Learning from unlabeled data.

Discovering hidden patterns and structures.

No target variable to predict.

# What is K-Means Clustering?

### **Definition:**

- An iterative algorithm that partitions data into K distinct clusters.
- Each data point belongs to the cluster with the nearest mean (centroid).

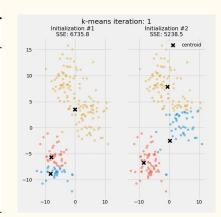
### Goal:

 Minimize the within-cluster variance (WCSS) or Sum of Squared Errors (SSE) that is defined as the sum of the squared Euclidean distances.

# The K-Means Algorithm: Step-by-Step

## **Algorithm 1** k-means algorithm

- 1: Specify the number k of clusters to assign.
- 2: Randomly initialize k centroids.
- 3: repeat
- 4: **expectation:** Assign each point to its closest centroid.
- 5: maximization: Compute the new centroid (mean) of each cluster.
- 6: until The centroid positions do not change.



## **Mathematical Formulation**

### **Distance Metric:**

- Typically Euclidean distance:  $d(x, \mu_k) = \sqrt{\sum_{i=1}^n (x_i \mu_{ki})^2}$
- Where x is a data point,  $\mu_k$  is the k-th centroid, and n is the number of dimensions.

## **Objective Function (WCSS or SSE):**

$$J = \sum_{k=1}^{K} \sum_{x_i \in C_k} d(x_i, \mu_k)^2$$

• Where  $C_k$  is the k-th cluster.

## **Centroid Update:**

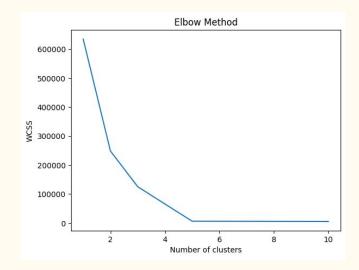
$$\mu_k = \frac{1}{|C_k|} \sum_{x_i \in C_k} x_i$$

• Where  $|C_k|$  is the number of points in cluster  $C_k$ .

# Determining the Optimal Number of Clusters (K)

### **Elbow Method:**

- Plot the WCSS (inertia) for different values of K.
- Identify the 'elbow' point where the rate of decrease in WCSS slows down.
- This point represents the optimal K.



## Performance Metrics

## Within-Cluster Sum of Squares (WCSS or SSE) / Inertia:

- Measures the compactness of clusters.
- Lower WCSS indicates tighter clusters.

#### **Silhouette Score:**

- Measures how similar an object is to its own cluster compared to other clusters.
- Ranges from -1 to 1: 1 indicates well-separated clusters, -1 indicates misclassification.

## **Adjusted Rand Index (ARI):**

- Unlike the silhouette coefficient, the ARI uses true cluster assignments to measure the similarity between true and predicted labels.
- The ARI output values range between -1 and 1. A score close to 0 indicates random assignments, and a score close to 1 indicates perfectly labeled clusters.

# Hyperparameters

### Number of Clusters (K):

- The most critical parameter.
- Determines the number of clusters to form.
- Must be chosen carefully, using the elbow method, or silhouette score

#### **Initialization Method (k-means++, random):**

- k-means++ helps to select better initial centroids, leading to faster convergence and better results.

### **Maximum Iterations (max\_iter):**

Limits the number of iterations to avoid infinite loops.

### Random State (random\_state):

 Controls the randomness of the initial centroid choice. By using a constant random state, the results can be reproduced."

# Real-World Applications

## **Customer Segmentation:**

- Grouping customers based on purchasing behavior, demographics, etc.

## **Anomaly Detection:**

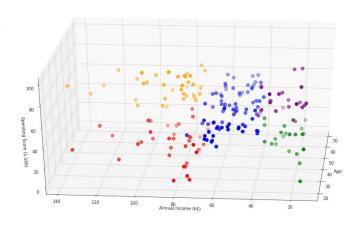
- Identifying unusual data points that deviate from normal patterns.
- Fraud detection, network security.

### **Document Clustering:**

- Grouping similar documents based on their content.
- Topic modeling, information retrieval.

### Genetics:

- Grouping genes with similar expression patterns.
- Disease classification.



# Advantages and Disadvantages

### Advantages:

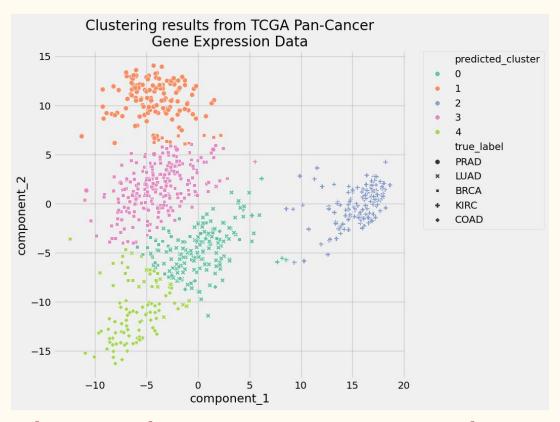
- Simple and easy to implement.
- Efficient for large datasets.
- Relatively scalable

### Disadvantages:

- Sensitive to initial centroid selection.
- Assumes spherical clusters.
- Requires pre-specifying K.
- Sensitive to outliers



# Coding



k means clustering programacion ii.ipynb