

## ASSIGNMENT - 02

Date: 08/08/2020

Title: Clustering

Problem

definition: Consider a suitable dataset. For clustering of data instances in different groups, apply different clustering techniques (min. 2).  
Visualize clusters using suitable tools.

SLW & Hlw

requirements: • R tool / Anaconda Python  
• PIV, 2GB RAM, 500 GB HDD.

Learning objectives: Use R functions / scikit-learn functions to create K-means clustering models and hierarchical clustering models.

Learning outcomes: Visualize the effects of K-means and hierarchical clustering using graphic capabilities.

Theory:

1] K-means Clustering.

- It is a type of unsupervised learning, which is used when you have unlabelled data.
- The goal of this algorithm is to find groups in the data, with the number of groups



represented by the variable  $K$ .

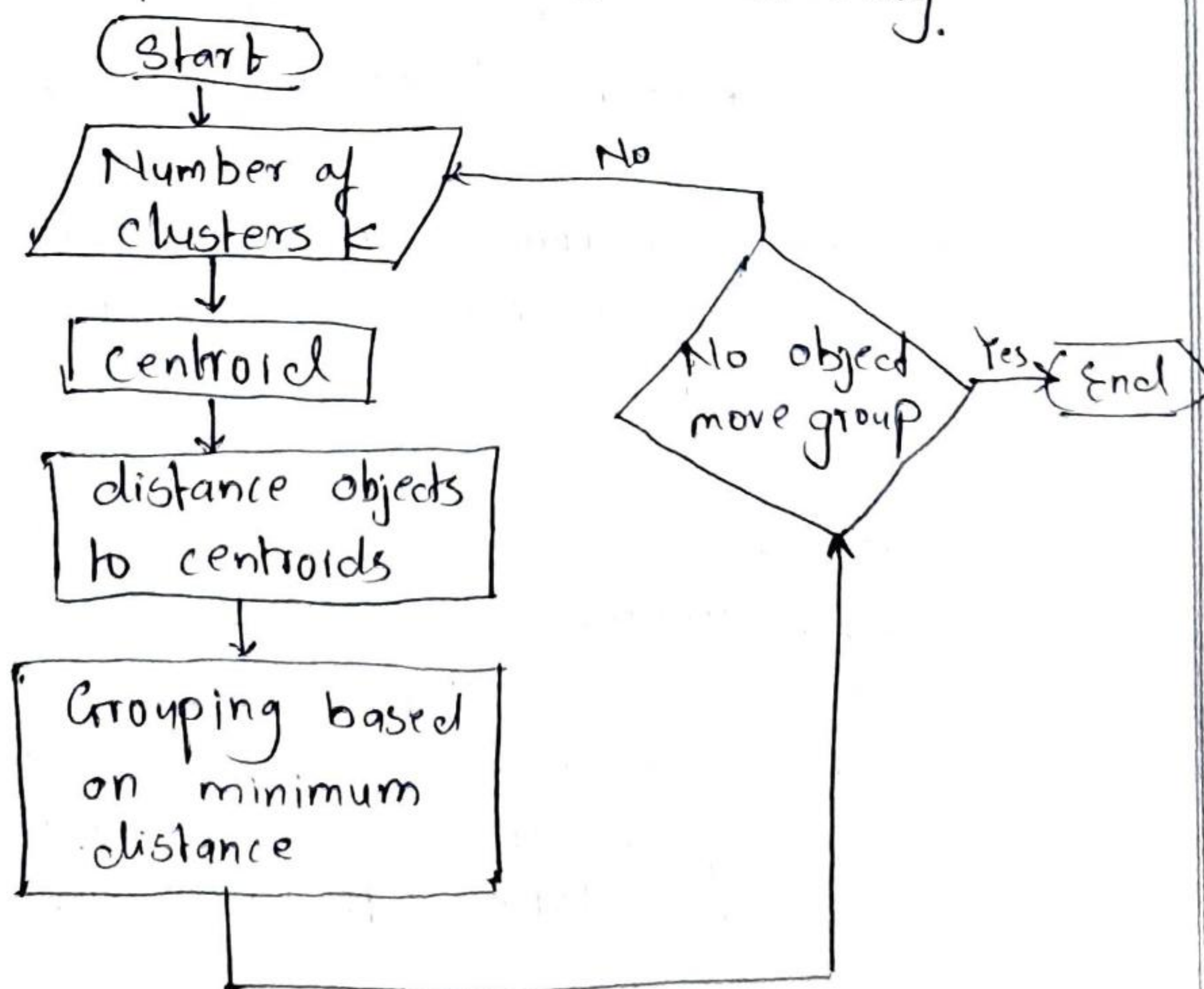
- The algorithm works iteratively to assign each data point to one of  $K$  groups based on features that are provided.
- Data points are clustered based on feature similarity.
- The results of K-means clustering algorithm are:

1. The centroids of the  $K$ -clusters, which can be used to label new data.

2. Labels for training data (each data point is assigned to a single cluster).

- Rather than defining groups before looking at the data, clustering allows you to find and analyze the groups that have been formed organically.

→ steps to perform K-means Clustering.





## B] Hierarchical Clustering

Hierarchical clustering involves creating clusters that have pre-determined ordering from top to bottom.

There are two types of hierarchical clustering.

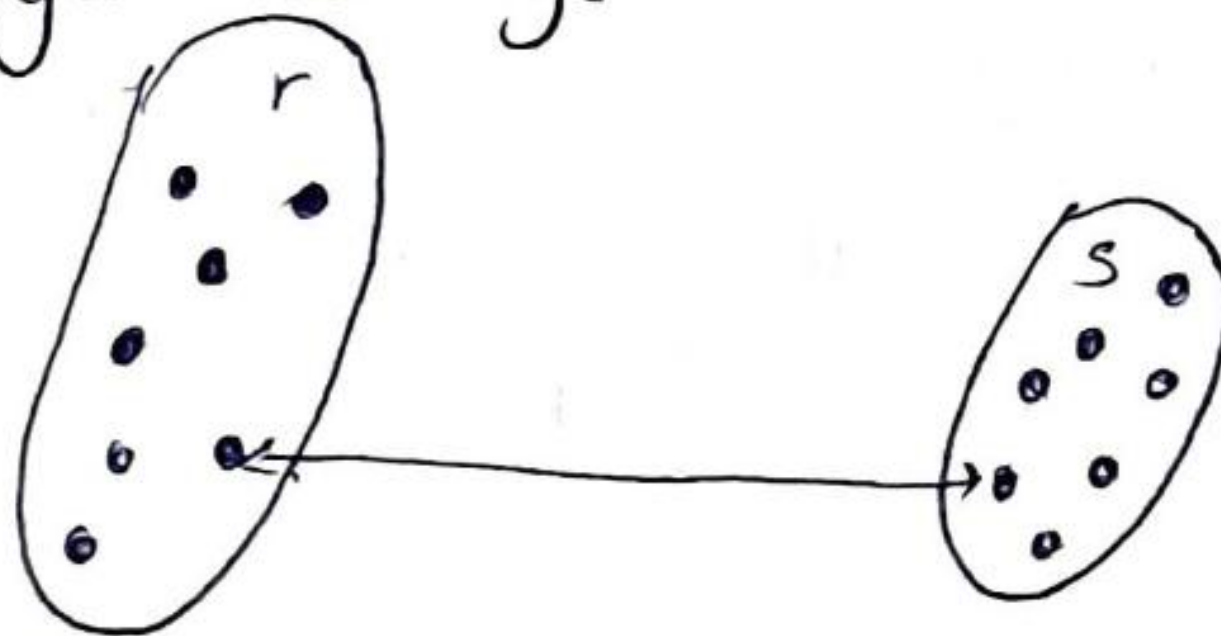
### 1. Divisive method

- It is also known as top-down clustering. We assign all of observations to a single cluster and then partition cluster into two least similar clusters.
- Finally, we proceed recursively on each cluster until there is one cluster for each observation.

### 2. Agglomerative method.

- It is also known as bottom-up clustering.
- We assign each observation to its own cluster.
- Computation algorithm:-
  1. Compute the proximity matrix.
  2. Let each data point be a cluster.
  3. Repeat: Merge the two closest clusters and update the proximity matrix.
  4. Until only a single cluster remains
- Following are the methods to determine proximity matrix.

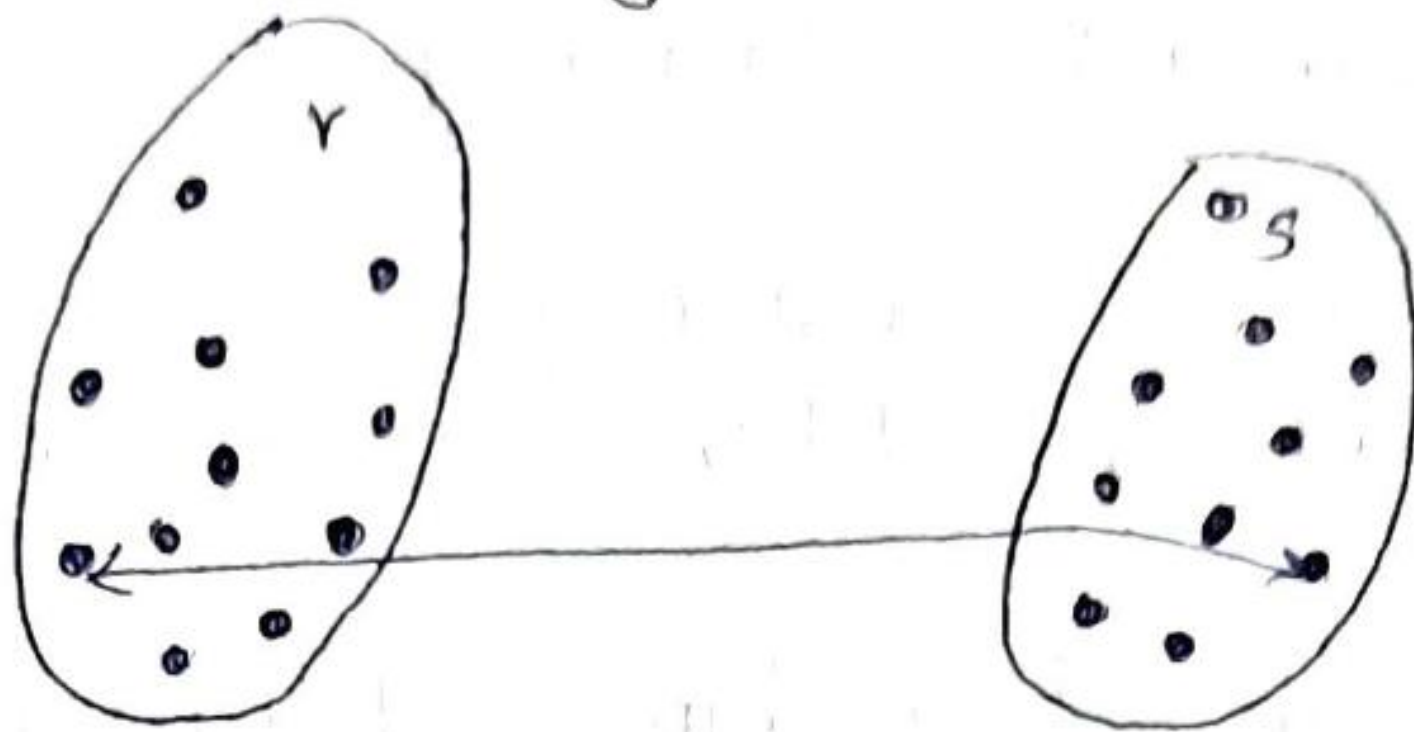
- Single linkage



$$L(r, s) = \min(D(x_{ri}, x_{sj}))$$

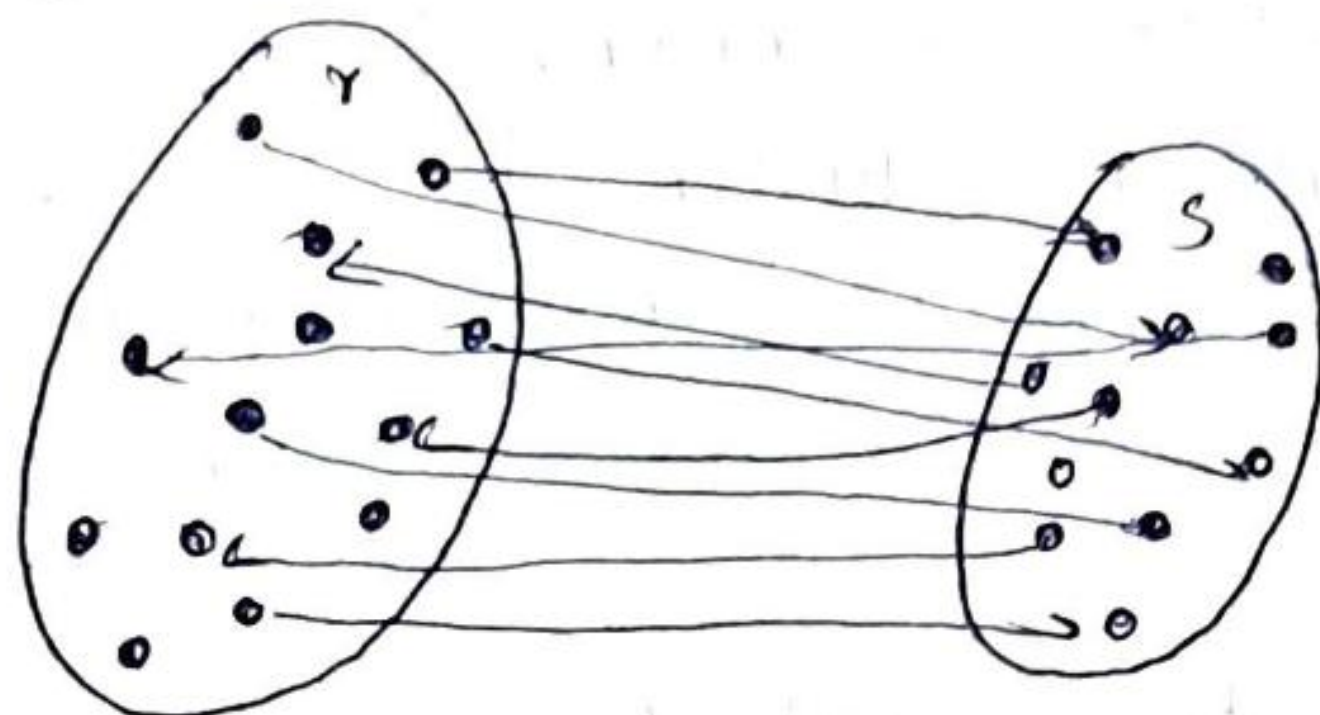


## 2. Complete linkage.



$$L(r, s) = \max (D(x_{ri}, x_{sj}))$$

## 3. Average Linkage.



$$L(r, s) = \frac{1}{n_r n_s} \sum_{i=1}^{n_r} \sum_{j=1}^{n_s} D(x_{ri}, x_{sj})$$

## Test cases:

Sr.no.	Description.	Expected o/p	Actual o/p
1.	In hierarchical clustering construct a dendrogram using "ward laverage" method.	No. of clusters rendered = 5	Success
2.	Visuale cluster using single, complete and average linkages.	A clusters are displayed by means of scatter plot	Success
3.	While fitting k-means to dataset, put random-state = 42.	Success	Success.



Conclusion: Hence, we have successfully implemented hierarchical clustering and K-means clustering algorithm in python using jupyter notebooks.