

Class: BE 3

Batch: P3

Roll no. 41310

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Assignment 2

Problem Statement:

Consider a suitable dataset. For clustering of data instances into different groups apply different clustering techniques (minimum 2). Visualize the clusters using suitable tool.

Learning Objective:

- To understand different clustering algorithms. Ex. Hierarchical Clustering, K-means Clustering.
- To understand how to visualize data using python libraries.

Software Requirements:

Anaconda, Spyder etc.

Hardware Requirement:

2GB RAM, 500 GB HDD.

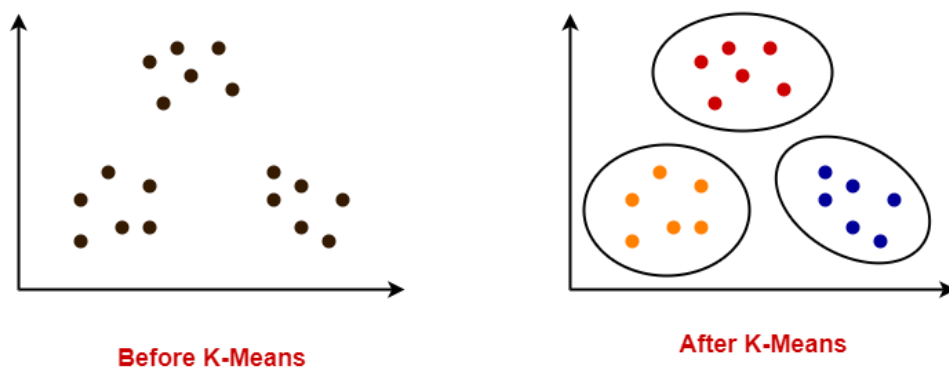
Theory:

K-means Clustering:

K-Means is probably the most well-known clustering algorithm. It's taught in a lot of introductory data science and machine learning classes. It's easy to understand and implement in code! Check out the graphic below for an illustration.

1. To begin, we first select a few classes/groups to use and randomly initialize their respective center points. To figure out the number of classes to use, it's good to take a quick look at the data and try to identify any distinct groupings. The center points are vectors of the same length as each data point vector and are the "X's" in the graphic above.
2. Each data point is classified by computing the distance between that point and each group center, and then classifying the point to be in the group whose center is closest to it.
3. Based on these classified points, we recompute the group center by taking the mean of all the vectors in the group.
4. Repeat these steps for a set number of iterations or until the group centers don't change much between iterations. You can also opt to randomly initialize the group centers a few times, and then select the run that looks like it provided the best results.

K-Means has the advantage that it's pretty fast, as all we're really doing is computing the distances between points and group centers; very few computations! It thus has a linear complexity $O(n)$.



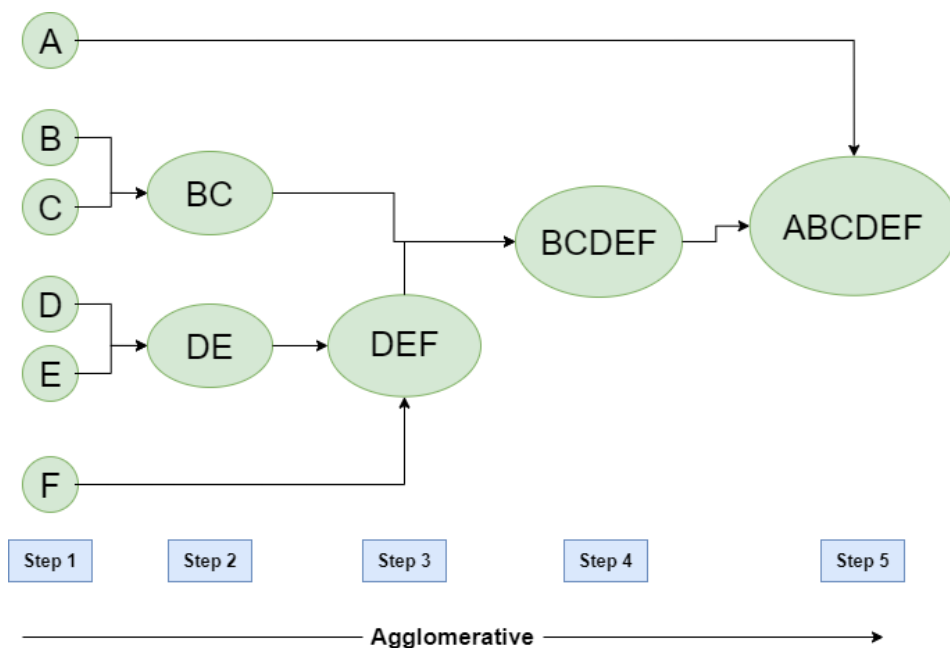
Agglomerative Hierarchical Clustering:

Hierarchical clustering algorithms fall into 2 categories: top-down or bottom-up. Bottom-up algorithms treat each data point as a single cluster at the outset and then successively merge (or agglomerate) pairs of clusters until all clusters have been merged into a single cluster that contains all data points. Bottom-up hierarchical clustering is therefore called hierarchical agglomerative clustering or HAC. This hierarchy of clusters is represented as a tree (or dendrogram). The root of the tree is the unique cluster that gathers all the samples, the leaves being the clusters with only one sample.

1. We begin by treating each data point as a single cluster i.e if there are X data points in our dataset then we have X clusters. We then select a distance metric that measures the distance between two clusters. As an example, we will use average linkage which defines the distance between two clusters to be the average distance between data points in the first cluster and data points in the second cluster.
2. On each iteration, we combine two clusters into one. The two clusters to be combined are selected as those with the smallest

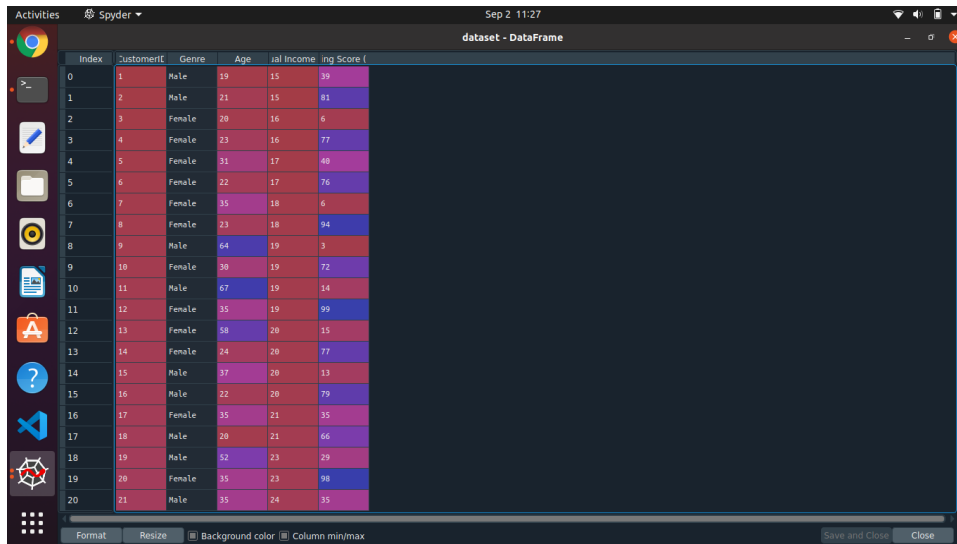
average linkage. I.e. according to our selected distance metric, these two clusters have the smallest distance between each other and therefore are the most similar and should be combined.

3. Step 2 is repeated until we reach the root of the tree i.e., we only have one cluster which contains all data points. In this way we can select how many clusters we want in the end, simply by choosing when to stop combining the clusters i.e. when we stop building the tree!



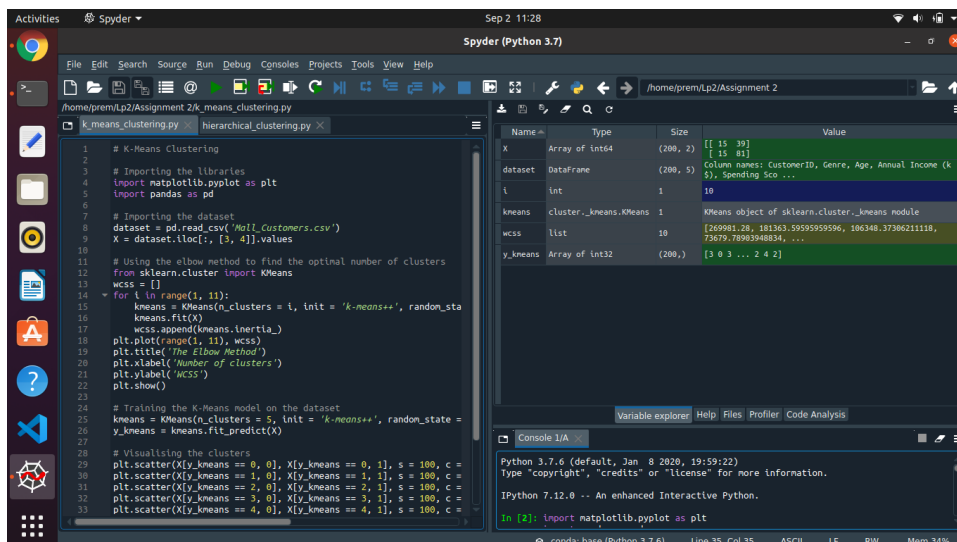
Python Implementation:

Dataset: Mall_Customer.csv



The screenshot shows the Spyder Python IDE interface. The main window displays a DataFrame titled 'dataset - DataFrame'. The DataFrame contains 21 rows (indexed 0 to 20) and 5 columns: 'Index', 'CustomerID', 'Genre', 'Age', and 'Annual Income (k\$)'. The data is as follows:

Index	CustomerID	Genre	Age	Annual Income (k\$)
0	1	Male	19	15
1	2	Male	21	15
2	3	Female	20	16
3	4	Female	23	16
4	5	Female	31	17
5	6	Female	22	17
6	7	Female	35	18
7	8	Female	23	18
8	9	Male	64	19
9	10	Female	30	19
10	11	Male	67	19
11	12	Female	35	19
12	13	Female	58	20
13	14	Female	24	20
14	15	Male	37	20
15	16	Male	22	20
16	17	Female	35	21
17	18	Male	20	21
18	19	Male	52	23
19	20	Female	35	23
20	21	Male	35	24



The screenshot shows the Spyder Python IDE interface with a Python script for K-Means clustering. The code is as follows:

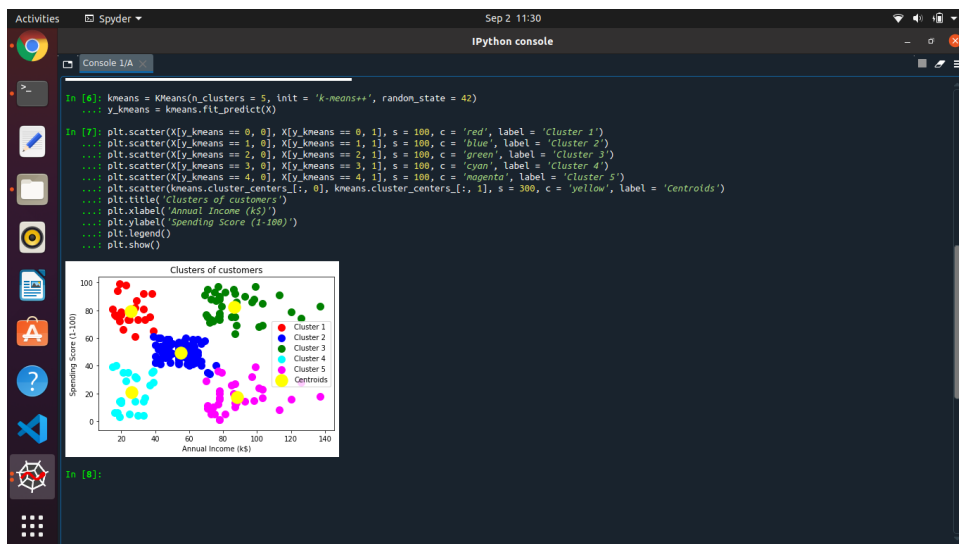
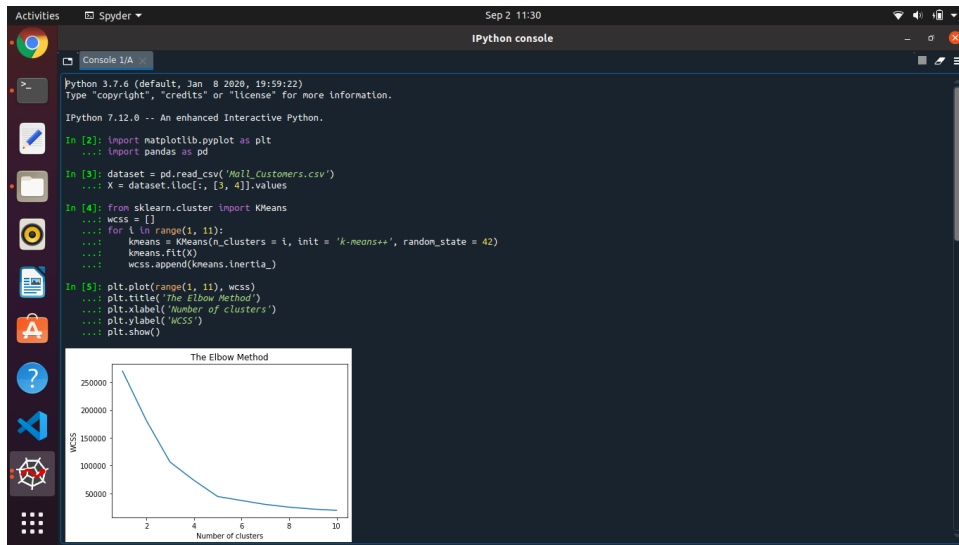
```
1 # K-Means Clustering
2
3 # Importing the libraries
4 import matplotlib.pyplot as plt
5 import pandas as pd
6
7 # Importing the dataset
8 dataset = pd.read_csv('Mall_Customers.csv')
9 X = dataset.iloc[:, [3, 4]].values
10
11 # Using the elbow method to find the optimal number of clusters
12 from sklearn.cluster import KMeans
13 wcss = []
14 for i in range(1, 11):
15     kmeans = KMeans(n_clusters = i, init = 'k-means++', random_state = 0)
16     kmeans.fit(X)
17     wcss.append(kmeans.inertia_)
18 plt.plot(range(1, 11), wcss)
19 plt.title('The Elbow Method')
20 plt.xlabel('Number of clusters')
21 plt.ylabel('WCSS')
22 plt.show()
23
24 # Training the K-Means model on the dataset
25 kmeans = KMeans(n_clusters = 5, init = 'k-means++', random_state = 0)
26 y_kmeans = kmeans.fit_predict(X)
27
28 # Visualising the clusters
29 plt.scatter(X[y_kmeans == 0, 0], X[y_kmeans == 0, 1], s = 100, c = 'red', label = 'Cluster 1')
30 plt.scatter(X[y_kmeans == 1, 0], X[y_kmeans == 1, 1], s = 100, c = 'blue', label = 'Cluster 2')
31 plt.scatter(X[y_kmeans == 2, 0], X[y_kmeans == 2, 1], s = 100, c = 'green', label = 'Cluster 3')
32 plt.scatter(X[y_kmeans == 3, 0], X[y_kmeans == 3, 1], s = 100, c = 'orange', label = 'Cluster 4')
33 plt.scatter(X[y_kmeans == 4, 0], X[y_kmeans == 4, 1], s = 100, c = 'purple', label = 'Cluster 5')
```

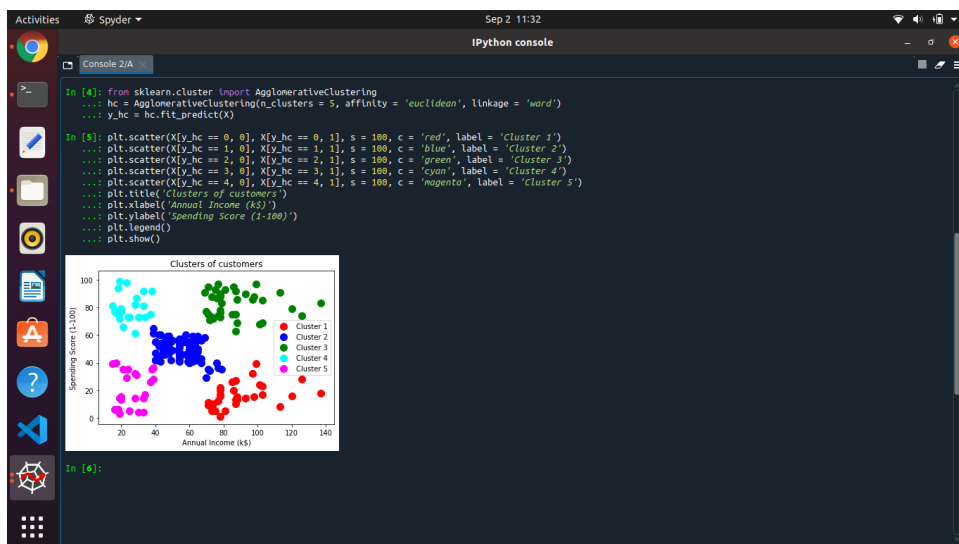
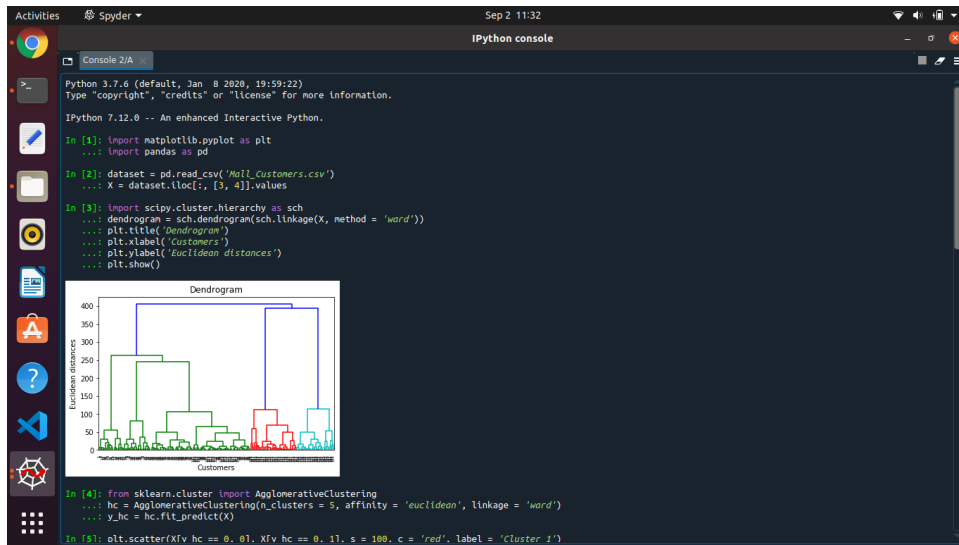
The Variable explorer on the right shows the following variables:

Name	Type	Size	Value
X	Array of int64	(200, 2)	[[15 39] [15 81] ...
dataset	DataFrame	(200, 5)	Column names: CustomerID, Genre, Age, Annual Income (k\$), Spending Score ...
i	int	1	10
kmeans	cluster_kmeans.KMeans	1	KMeans object of sklearn.cluster_kmeans module
wcss	list	10	[269981.28, 181363.59595959596, 106348.37306211118, 78579.7893348834, ...]
y_kmeans	Array of int32	(200,)	[3 0 3 ... 2 4 2]

The Console window at the bottom shows the following output:

```
Python 3.7.6 (default, Jan 8 2020, 19:59:22)
Type "copyright", "credits" or "license" for more information.
IPython 7.12.0 -- An enhanced Interactive Python.
In [2]: import matplotlib.pyplot as plt
```





Conclusion:

Thus, Clustering techniques like K means and Hierarchical Clustering were used for performing clustering on Mall Customer.csv dataset. Moreover, the clusters were visualize using matplotlib. The assignment was implemented successfully using Spyder.