K-Means Clustering

Clustering - grouping unlabelled data based on their similarity or proximity.

```
In [2]: #Importing the libraries
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
```

Importing the dataset

```
df = pd.read csv(r"C:\Users\GIRISH\Desktop\INTROTALLENT\PYTHON\ML PROJECT\Customers spen
In [3]:
         df.head()
In [4]:
Out[4]:
            CustomerID
                        Genre Age Annual Income (k$) Spending Score (1-100)
         0
                                19
                                                                       39
                    1
                         Male
                                                  15
         1
                    2
                         Male
                                21
                                                  15
                                                                       81
         2
                    3 Female
                                20
                                                  16
                                                                        6
         3
                    4 Female
                                23
                                                  16
                                                                       77
         4
                    5 Female
                                31
                                                  17
                                                                       40
```

retrieve the values from columns 3 and 4 of the df (all rows) and return them as a NumPy array.

```
x=df.iloc[:,[3, 4]].values
In [5]:
In [6]:
        array([[ 15,
                        391,
Out[6]:
                [ 15,
                        81],
                [ 16,
                        6],
                [ 16,
                        77],
                [ 17,
                       40],
                [ 17,
                        76],
                [ 18,
                        6],
                [ 18,
                        94],
                [ 19,
                        3],
                        72],
                [ 19,
                [ 19,
                        14],
                [ 19,
                        991,
                [ 20,
                        15],
                [ 20,
                        77],
                [ 20,
                        13],
                [ 20,
                        79],
                [ 21,
                        35],
                [ 21,
                       66],
                [ 23,
                        29],
                [ 23,
                       98],
                [ 24,
                        35],
                 [ 24,
                        73],
                 [ 25,
                       51,
                 [ 25, 73],
```

```
[ 28,
        14],
[ 28,
        82],
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        32],
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        61],
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[ 29,
        87],
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        55],
[ 54,
        41],
[ 54,
        44],
[ 54,
        57],
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[54,

[58,

[57,

[57,

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46],

58],

55],

60],

46],

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[ 59,
        55],
[ 59,
        41],
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        49],
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        43],
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        10],
[ 74,
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[ 75,
         5],
[ 75,
        93],
[ 76,
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[ 76,
        87],
  77,
[
        12],
[ 77,
        97],
[ 77,
        36],
[ 77,
        74],
[ 78,
        22],
[ 78,
        90],
[ 78,
        17],
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78,

[78,

[78,

[78,

[78,

88],

20],

76],

16],

89],

```
1],
[ 78,
[ 78, 78],
[ 78, 1],
[ 78, 73],
[ 79, 35],
[ 79, 83],
[ 81, 5],
[ 81, 93],
[ 85, 26],
[ 85, 75],
[ 86, 20],
[ 86, 95],
[ 87, 27],
[ 87, 63],
[ 87, 13],
[ 87,
      75],
[ 87, 10],
[ 87, 92],
[ 88, 13],
[ 88, 86],
[ 88, 15],
[ 88, 69],
[ 93, 14],
[ 93, 90],
[ 97, 32],
[ 97, 86],
[ 98, 15],
[ 98, 88],
[ 99, 39],
[ 99, 97],
[101, 24],
[101, 68],
[103, 17],
[103, 85],
[103, 23],
[103, 69],
[113, 8],
[113, 91],
[120, 16],
[120, 79],
[126, 28],
[126, 74],
[137, 18],
```

[137, 83]], dtype=int64)

Using the elbow method to find the optimal number of clusters

The elbow method is a technique used in clustering analysis to determine the optimal number of clusters to use for partitioning a dataset.

WCSS (within-cluster sum of squares)

- The WCSS measures the sum of the squared distances between each data point and the centroid of its assigned cluster.
- In general, as you increase the number of clusters, the WCSS will decrease since data points will be closer to the centroids of their respective clusters. However, adding too many clusters may lead to

Elbow Point

The "elbow point" on the WCSS plot is the point at which the rate of decrease sharply changes, creating an "elbow" shape. This point represents a balance between minimizing WCSS and avoiding too many clusters. The idea is to choose the number of clusters at this elbow point

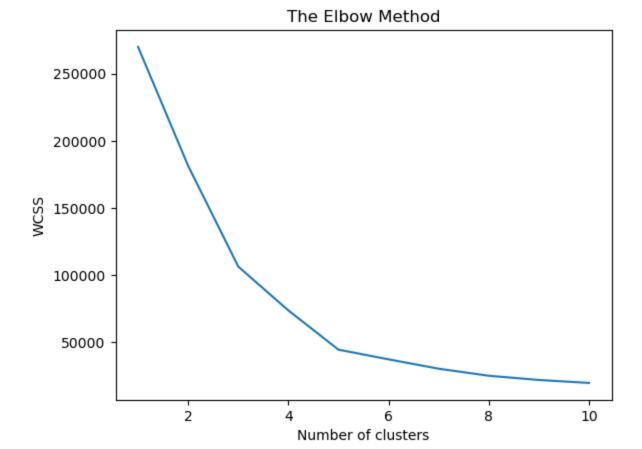
```
In [7]: from sklearn.cluster import KMeans
        wcss = []
        for i in range (1, 11):
            kmeans = KMeans(n clusters = i, init = 'k-means++', random state = 42)
           kmeans.fit(x)
           wcss.append(kmeans.inertia)
        plt.plot(range(1, 11), wcss)
        plt.title('The Elbow Method')
        plt.xlabel('Number of clusters')
        plt.ylabel('WCSS')
        plt.show()
        C:\Users\GIRISH\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.py:870: FutureWarnin
        g: The default value of `n init` will change from 10 to 'auto' in 1.4. Set the value of
        `n init` explicitly to suppress the warning
         warnings.warn(
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        g: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks
        than available threads. You can avoid it by setting the environment variable OMP NUM THR
        EADS=1.
         warnings.warn(
        C:\Users\GIRISH\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.py:870: FutureWarnin
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EADS=1.
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`n init` explicitly to suppress the warning
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g: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks
than available threads. You can avoid it by setting the environment variable OMP NUM THR
```

EADS=1.

warnings.warn(



Training the K-Means model on the dataset

```
In [8]: kmeans = KMeans(n_clusters = 5, init = 'k-means++', random_state = 100)
    y_kmeans = kmeans.fit_predict(x)

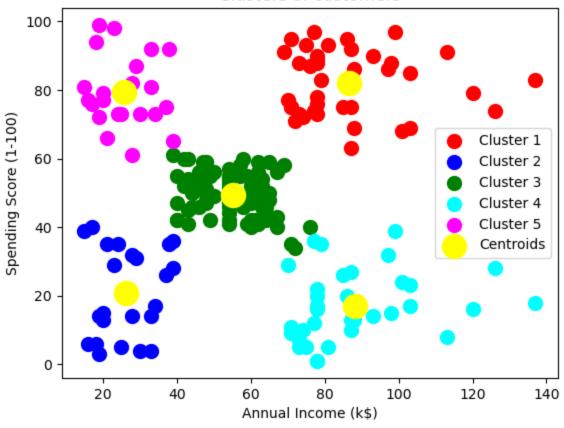
C:\Users\GIRISH\anaconda3\lib\site-packages\sklearn\cluster\_kmeans.py:870: FutureWarnin
    g: The default value of `n_init` will change from 10 to 'auto' in 1.4. Set the value of
    `n_init` explicitly to suppress the warning
        warnings.warn(
    C:\Users\GIRISH\anaconda3\lib\site-packages\sklearn\cluster\_kmeans.py:1382: UserWarnin
    g: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks
    than available threads. You can avoid it by setting the environment variable OMP_NUM_THR
    EADS=1.
    warnings.warn(
```

Visualising the clusters

plt.scatter(x[y_kmeans == 2, 0], x[y_kmeans == 2, 1], s = 100, c = 'green', label = 'Clu plt.scatter(x[y kmeans == 3, 0], x[y kmeans == 3, 1], s = 100, c = 'cyan', label = 'Clus

```
plt.scatter(x[y_kmeans == 4, 0], x[y_kmeans == 4, 1], s = 100, c = 'magenta', label = 'C
plt.scatter(kmeans.cluster_centers_[:, 0], kmeans.cluster_centers_[:, 1], s = 300, c = '
plt.title('Clusters of customers')
plt.xlabel('Annual Income (k$)')
plt.ylabel('Spending Score (1-100)')
plt.legend()
plt.show()
```

Clusters of customers



In []: