## PREDICTION USING UNSUPERVISED MACHINE LEARNING MODEL

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**Iris Versicolor** 

Iris Setosa

Iris Virginica

Task- Predict the optimum number of clusters that will be formed

We have the famous 'iris dataset' and our task is to find out the number of clusters that can be formed with the help of K- means clustering.

The algorithm follows a simple and easy way to group a given data set into a certain number of coherent subsets called as clusters. The idea is to find K centres, called as cluster centroids, one for each cluster, hence the name K-means clustering. It has the aims to divide 'n' observations into k clusters in which each observation belongs to the cluster with the nearest mean, serving as a prototype of the cluster.

#### Steps involved in K-means clustering-

1. Select the number 'k' to determine the clusters which needs to be form.

- 2. Select random K points or centroids
- 3.Each dataset point need to be assign with each of the three centroids, which ever will be the closest to them.
- 4. Calculate the euclidean distance between them .
- 5. Calculate the variance and place a new centroid of each cluster.
- 6. Reassign each data point again with the nearest centroids
- 7.If there is no change in any assignment of the datasets to another cluster, we say that our clustering is over.

### Importing libraries

```
In []: import pandas as pd
   import numpy as np
   import matplotlib.pyplot as plt
   import seaborn as sns
   from sklearn.cluster import KMeans
   from sklearn.metrics import silhouette_score
   from sklearn.preprocessing import MinMaxScaler
```

#### Reading dataset

```
In [7]: | iris = pd.read csv("C:/Users/91913/Downloads/Iris.csv")
        x = iris.iloc[:, [0, 1, 2, 3]].values
In [9]: | iris.info()
        iris[0:10]
        <class 'pandas.core.frame.DataFrame'>
        RangeIndex: 150 entries, 0 to 149
        Data columns (total 6 columns):
         150 non-null int64
         1 SepalLengthCm 150 non-null float64
2 SepalWidthCm 150 non-null float64
3 PetalLengthCm 150 non-null float64
         4 PetalWidthCm 150 non-null float64
                           150 non-null object
         5 Species
        dtypes: float64(4), int64(1), object(1)
        memory usage: 7.2+ KB
Out[9]:
```

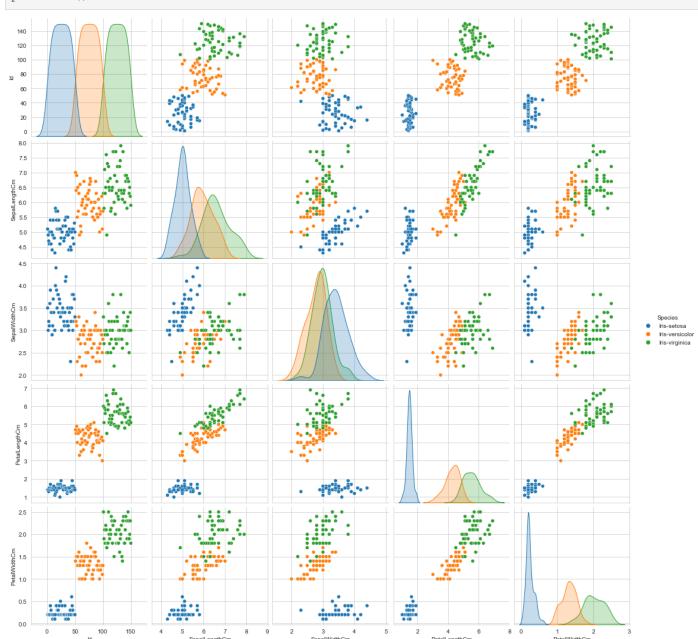
	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
0	1	5.1	3.5	1.4	0.2	Iris-setosa
1	2	4.9	3.0	1.4	0.2	Iris-setosa
2	3	4.7	3.2	1.3	0.2	Iris-setosa
3	4	4.6	3.1	1.5	0.2	Iris-setosa

4	5	5.0	3.6	1.4	0.2 Iris-setosa
5	6	5.4	3.9	1.7	0.4 Iris-setosa
6	7	4.6	3.4	1.4	0.3 Iris-setosa
7	8	5.0	3.4	1.5	0.2 Iris-setosa
8	9	4.4	2.9	1.4	0.2 Iris-setosa
9	10	4.9	3.1	1.5	0.1 Iris-setosa

```
In [13]: iris_setosa=iris.loc[iris["Species"]=="Iris-setosa"]
    iris_virginica=iris.loc[iris["Species"]=="Iris-virginica"]
    iris_versicolor=iris.loc[iris["Species"]=="Iris-versicolor"]
```

### **Scatter plot**

```
In [21]: sns.set_style("whitegrid")
    sns.pairplot(iris, hue="Species", height=3);
    plt.show()
```

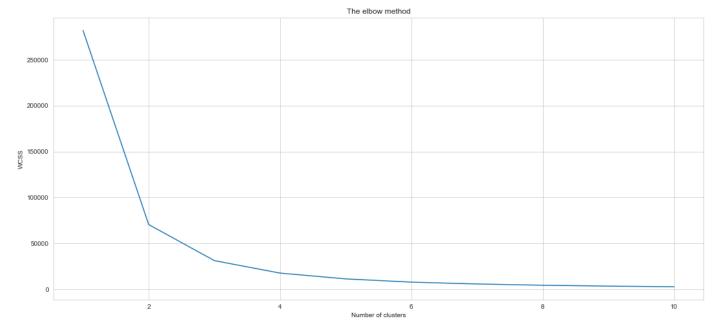


#### How do we know that what value we will choose for K?

The performance of the K-means clustering algorithm depends upon highly efficient clusters that it forms. But choosing the optimal number of clusters is a big task. There are some different ways to find the optimal number of clusters, but here we are discussing the most appropriate method to find the number of clusters or value of K.

# Using the elbow method to determine the optimal number of clusters for k-means clustering

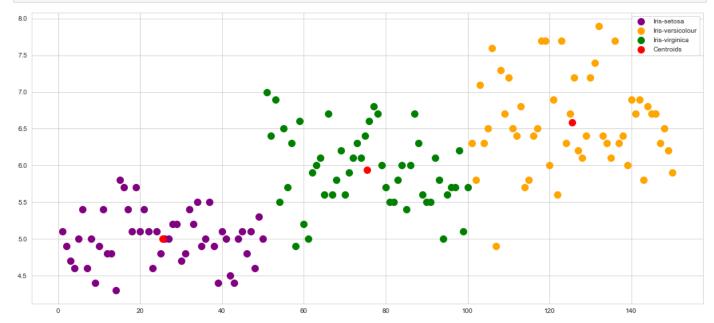
```
In [26]: plt.figure(figsize=(18,8))
   plt.plot(range(1, 11), wcss)
   plt.title('The elbow method')
   plt.xlabel('Number of clusters')
   plt.ylabel('WCSS') #within cluster sum of squares
   plt.show()
```



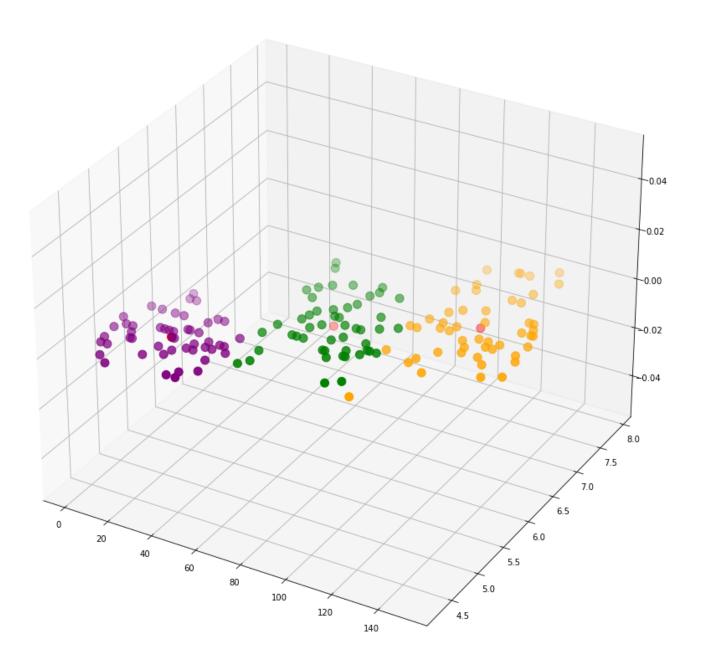
#### Implementing K-Means Clustering

The Value of K that we can get from the graph will be 3.

```
In [30]: #Visualising the clusters
         plt.figure(figsize=(18,8))
         plt.scatter(x[y kmeans == 0, 0],
                     x[y kmeans == 0, 1],
                     s = 100, c = 'purple', label = 'Iris-setosa')
         plt.scatter(x[y kmeans == 1, 0],
                     x[y kmeans == 1, 1],
                     s = 100, c = 'orange', label = 'Iris-versicolour')
         plt.scatter(x[y kmeans == 2, 0],
                     x[y kmeans == 2, 1],
                     s = 100, c = 'green', label = 'Iris-virginica')
         #Plotting the centroids of the clusters
         plt.scatter(kmeans.cluster_centers_[:, 0],
                     kmeans.cluster centers [:,1],
                     s = 100, c = 'red', label = 'Centroids')
         plt.legend()
         plt.show()
```



```
s = 100, c = 'red', label = 'Centroids')
plt.show()
```



In [ ]: