

GRIP : THE SPARKS FOUNDATION

DATA SCIENCE AND BUSINESS ANALYTICS INTERN

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TASK 2 : Prediction using Unsupervised ML

Aim : To predict the optimum number of clusters and represent it visually.

Importing the Data

```
In [30]: #importing all the libraries
import numpy as np
import pandas as pd
import seaborn as sns
from matplotlib import pyplot as plt
from sklearn import datasets
from sklearn.cluster import KMeans
```

```
In [38]: #Reading the data
df = datasets.load_iris()
df = pd.DataFrame(iris.data, columns = iris.feature_names)
df.head()
```

```
Out[38]:
```

| | sepal length (cm) | sepal width (cm) | petal length (cm) | petal width (cm) |
|---|-------------------|------------------|-------------------|------------------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 |
| 3 | 4.6 | 3.1 | 1.5 | 0.2 |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 |

```
In [43]: df.info
```

```
Out[43]: <bound method DataFrame.info of
etal length (cm)  petal width (cm)
0                5.1          3.5          1.4          0
.2
1                4.9          3.0          1.4          0
.2
2                4.7          3.2          1.3          0
.2
3                4.6          3.1          1.5          0
.2
4                5.0          3.6          1.4          0
.2
..              ...          ...          ...          .
..
145              6.7          3.0          5.2          2
.3
146              6.3          2.5          5.0          1
.9
147              6.5          3.0          5.2          2
.0
148              6.2          3.4          5.4          2
.3
149              5.9          3.0          5.1          1
.8

[150 rows x 4 columns]>
```

```
In [57]: df.shape
```

```
Out[57]: (149, 4)
```

```
In [58]: df.columns
```

```
Out[58]: Index(['sepal length (cm)', 'sepal width (cm)', 'petal length (cm)',
               'petal width (cm)'],
              dtype='object')
```

```
In [62]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 149 entries, 0 to 149
Data columns (total 4 columns):
#   Column                Non-Null Count  Dtype
---  -
0   sepal length (cm)      149 non-null   float64
1   sepal width (cm)       149 non-null   float64
2   petal length (cm)      149 non-null   float64
3   petal width (cm)       149 non-null   float64
dtypes: float64(4)
memory usage: 5.8 KB
```

```
In [63]: df.describe()
```

Out[63]:

| | sepal length (cm) | sepal width (cm) | petal length (cm) | petal width (cm) |
|--------------|-------------------|------------------|-------------------|------------------|
| count | 149.000000 | 149.000000 | 149.000000 | 149.000000 |
| mean | 5.843624 | 3.059732 | 3.748993 | 1.194631 |
| std | 0.830851 | 0.436342 | 1.767791 | 0.762622 |
| min | 4.300000 | 2.000000 | 1.000000 | 0.100000 |
| 25% | 5.100000 | 2.800000 | 1.600000 | 0.300000 |
| 50% | 5.800000 | 3.000000 | 4.300000 | 1.300000 |
| 75% | 6.400000 | 3.300000 | 5.100000 | 1.800000 |
| max | 7.900000 | 4.400000 | 6.900000 | 2.500000 |

In [64]: `df.isnull().sum()`

Out[64]:

```
sepal length (cm)    0
sepal width (cm)     0
petal length (cm)    0
petal width (cm)     0
dtype: int64
```

In [65]: `df.drop_duplicates(inplace=True)`

Data Visualisation

In [52]: `x = df.iloc[:, [0,1,2, 3]].values`

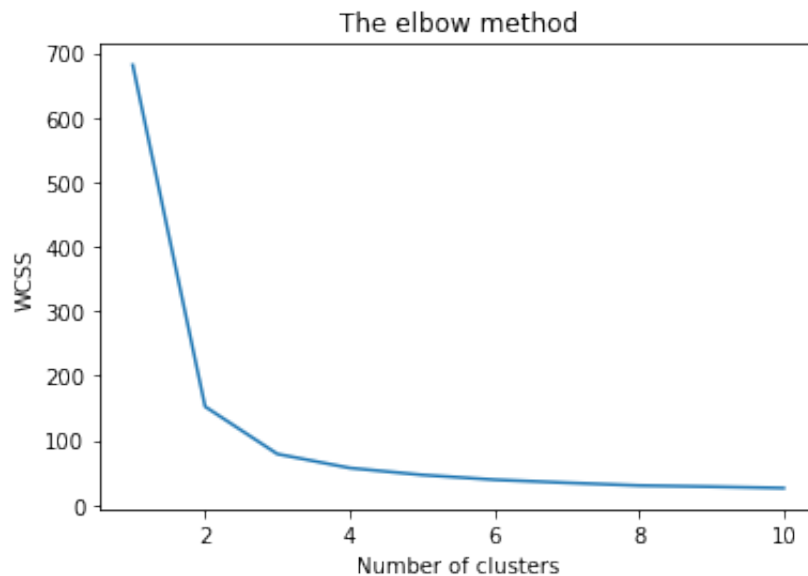
In [82]:

```
from sklearn.cluster import KMeans
wcss = []

for i in range(1, 11):
    kmeans = KMeans(n_clusters = i, init = 'k-means++',
                    max_iter = 300, n_init = 10, random_state = 0)
    kmeans.fit(x)
    wcss.append(kmeans.inertia_)
```

In [80]:

```
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()
```



To determine the optimal number of clusters, the value of k at the elbow is selected i.e. the point after which the distortion start decreasing in a linear fashion. Thus from the above graph, it can be said that the optimum clusters the data is 3.

```
In [84]: # K-Means to the dataset
kmeans = KMeans(n_clusters = 3, init = 'k-means++', random_state = 0)
```

The experimental results show the robustness of the Y-means algorithm as well as its good performance against a set of other well known unsupervised clustering techniques.

```
In [85]: y_kmeans = kmeans.fit_predict(X)
```

```
In [88]: #visualising the data
import seaborn as sns
#plt.figure(figsize=(9,5))
sns.scatterplot(X[y_kmeans == 0, 0], X[y_kmeans == 0, 1], color = 'red', label = 0)
sns.scatterplot(X[y_kmeans == 1, 0], X[y_kmeans == 1, 1], color = 'blue', label = 1)
sns.scatterplot(X[y_kmeans == 2, 0], X[y_kmeans == 2, 1], color = 'green', label = 2)

## Plotting the centroids of the clusters
sns.scatterplot(kmeans.cluster_centers[:, 0], kmeans.cluster_centers[:, 1], label = 'Centroids', s=100)
plt.grid(False)
plt.title('Clusters of Iris')
plt.legend()
plt.show()
```

