# The Sparks Foundation - Data Science & Business Analytics Internship

# TASK 2 - Prediction using Unsupervised Machine Learning

In this task it is required to predict the optimum number of cluster for the iris data set .iris data set consists of 3 types of flower namely Iris-setosa Iris-versicolour and Iris-virginica

## Steps:

\*Step 1 - Importing the dataset

\*Step 2 - Visualisng the data

\*Step 3 - Finding the optimum number of clusters

\*Step 4 - Applying k means clustering on the data

\*Step 5 - Visualising the clusters

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# STEP-1 Importing the data

In this step we will import the required libraries and data set with the help of pandas library

#### In [1]:

```
# Importing the required Libraries
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from sklearn import datasets
from sklearn.cluster import KMeans

# To ignore the warnings
import warnings as wg
wg.filterwarnings("ignore")
```

#### In [2]:

```
# Reading data iris dataset
df = pd.read_csv("taskdata.csv")
```

```
In [3]:
```

```
df.head()
```

## Out[3]:

	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

# Step 2 - Visualisng the data

In this setp we will try to visualize our dataset

## In [4]:

```
df.tail()
```

## Out[4]:

	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm	Species
145	146	6.7	3.0	5.2	2.3	Iris-virginica
146	147	6.3	2.5	5.0	1.9	Iris-virginica
147	148	6.5	3.0	5.2	2.0	Iris-virginica
148	149	6.2	3.4	5.4	2.3	Iris-virginica
149	150	5.9	3.0	5.1	1.8	Iris-virginica

## In [5]:

```
df.shape
```

#### Out[5]:

(150, 6)

## In [6]:

```
df.columns
```

## Out[6]:

## In [7]:

```
df['Species'].unique()
```

## Out[7]:

array(['Iris-setosa', 'Iris-versicolor', 'Iris-virginica'], dtype=object)

## In [8]:

```
df.info()
```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 6 columns):

#	Column	Non-Null Count	Dtype
0	Id	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
5	Species	150 non-null	object
dtyp	es: float64(4),	int64(1), objec	t(1)

utypes. 110ato4(4), 111to4(1), 01

memory usage: 7.2+ KB

## In [9]:

```
df.describe()
```

## Out[9]:

	ld	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
count	150.000000	150.000000	150.000000	150.000000	150.000000
mean	75.500000	5.843333	3.054000	3.758667	1.198667
std	43.445368	0.828066	0.433594	1.764420	0.763161
min	1.000000	4.300000	2.000000	1.000000	0.100000
25%	38.250000	5.100000	2.800000	1.600000	0.300000
50%	75.500000	5.800000	3.000000	4.350000	1.300000
75%	112.750000	6.400000	3.300000	5.100000	1.800000
max	150.000000	7.900000	4.400000	6.900000	2.500000

## In [10]:

```
# now we will drop the label column because it is an unsupervised learning problem
iris = pd.DataFrame(df)
iris_df = iris.drop(columns= ['Species' ,'Id'] )
iris_df.head()
```

#### Out[10]:

	SepalLengthCm	SepalWidthCm	PetalLengthCm	PetalWidthCm
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

## Step 3 - Finding the optimum number of clusters

Before clustering the data using kmeans, we need to specify the number of clusters. In order to find the optimum number of clusters, there are various methods available like Silhouette Coefficient and the Elbow method. Here, the elbow method is used.

#### Brief about the Elbow method

In this method, the number of clusters are varies within a certain range. For each number, within-cluster sum of square (wss) value is calculated and stored in a list. These value are then plotted against the range of number of clusters used before. The location of bend in the 2d plot indicates the appropriate number of clusters.

#### In [11]:

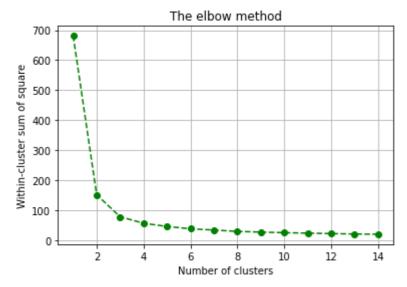
```
# Calculating the within-cluster sum of square
within_cluster_sum_of_square = []

clusters_range = range(1,15)
for k in clusters_range:
    km = KMeans(n_clusters=k)
    km = km.fit(iris_df)
    within_cluster_sum_of_square.append(km.inertia_)
```

#### In [12]:

```
# Plotting the "within-cluster sum of square" against clusters range

plt.plot(clusters_range, within_cluster_sum_of_square, 'go--', color='green')
plt.title('The elbow method')
plt.xlabel('Number of clusters')
plt.ylabel('Within-cluster sum of square')
plt.grid()
plt.show()
```



we can clearly see why it is called 'The elbow method' from the above graph, the optimum clusters is where the elbow occurs. This is when the within cluster sum of squares (WCSS) doesn't decrease significantly with every iteration.

From this we choose the number of clusters as

"3"

# Step 4 - Applying k means clustering on the data

#### In [13]:

```
from sklearn.cluster import KMeans
model = KMeans(n_clusters = 3, init = 'k-means++', max_iter = 300, n_init = 10, random_stat
predictions = model.fit_predict(iris_df)
```

## Step 5 - Visualising the clusters

#### In [14]:

```
x = iris_df.iloc[:, [0, 1, 2, 3]].values
plt.scatter(x[predictions == 0, 0], x[predictions == 0, 1], s = 25, c = 'red', label = 'Iri
plt.scatter(x[predictions == 1, 0], x[predictions == 1, 1], s = 25, c = 'blue', label = 'Ir
plt.scatter(x[predictions == 2, 0], x[predictions == 2, 1], s = 25, c = 'green', label = 'I

# Plotting the cluster centers

plt.scatter(model.cluster_centers_[:, 0], model.cluster_centers_[:,1], s = 100, c = 'yellow
plt.legend()
plt.grid()
plt.show()
```

