

Amey Ghadge_TSF Data Science intern

Task-2 Prediction using unsupervised machine learning

```
In [1]: #Importing the Libraraies
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [2]: df=pd.read_csv("Iris.csv")
```

```
In [3]: df
```

```
Out[3]:
```

| | Id | 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 |
| ... | ... | ... | ... | ... | ... | ... |
| 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 |

150 rows × 6 columns

```
In [4]: df.isnull().any()
```

```
Out[4]: Id                False
SepalLengthCm          False
SepalWidthCm           False
PetalLengthCm          False
PetalWidthCm           False
Species                False
dtype: bool
```

```
In [5]: df.shape
```

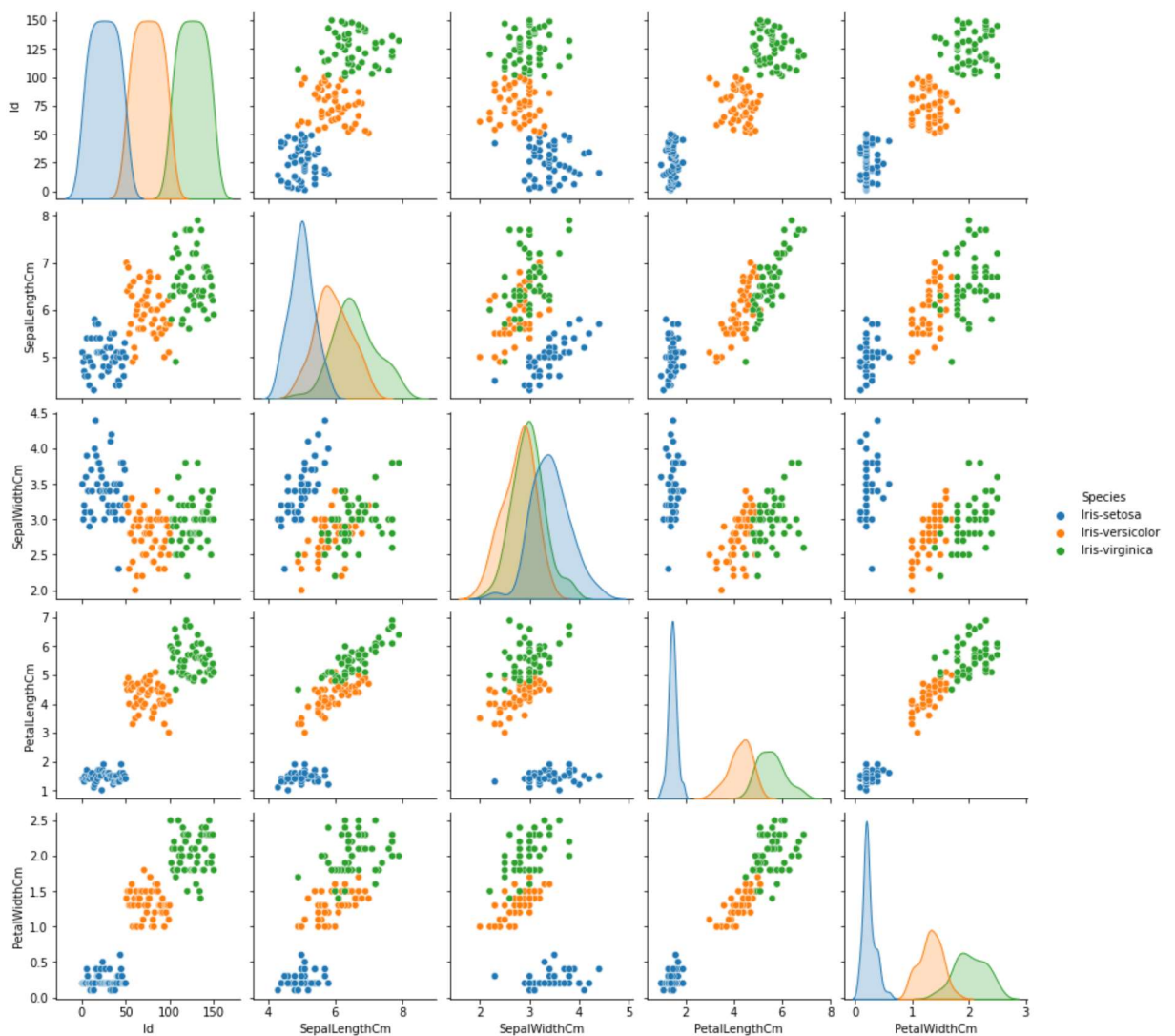
```
Out[5]: (150, 6)
```

```
In [6]: #getting information of dataset
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
dtypes: float64(4), int64(1), object(1)
memory usage: 7.2+ KB
```

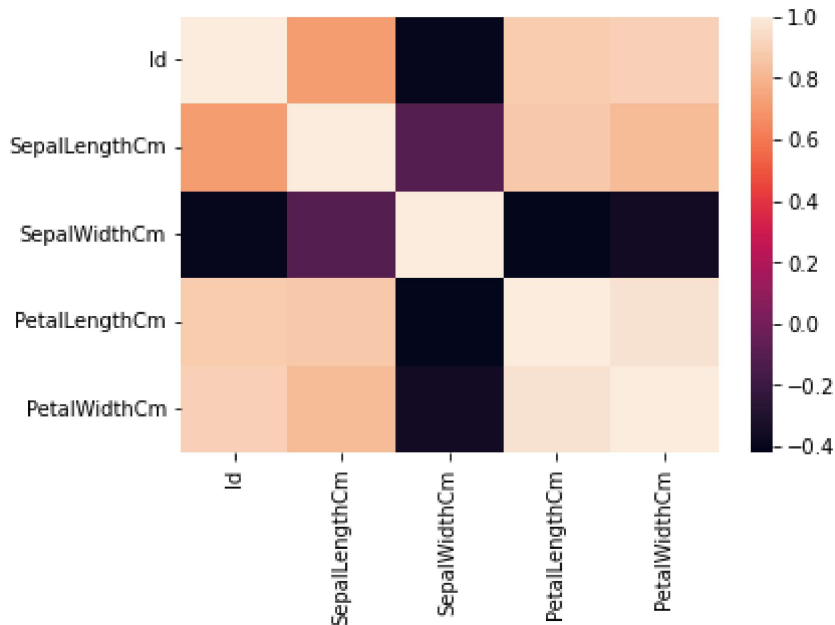
```
In [7]: # Let's plot pair plot to visualise the attributes all at once
sns.pairplot(data=df, hue = 'Species')
```

```
Out[7]: <seaborn.axisgrid.PairGrid at 0x2b51ad53df0>
```



```
In [8]: # correlation matrix
sns.heatmap(df.corr())
```

```
Out[8]: <AxesSubplot:>
```



```
In [9]: x = df.iloc[:, [0,1,2,3]].values
```

```
In [10]: #Finding the optimum number of clusters for k-means classification

from sklearn.cluster import KMeans

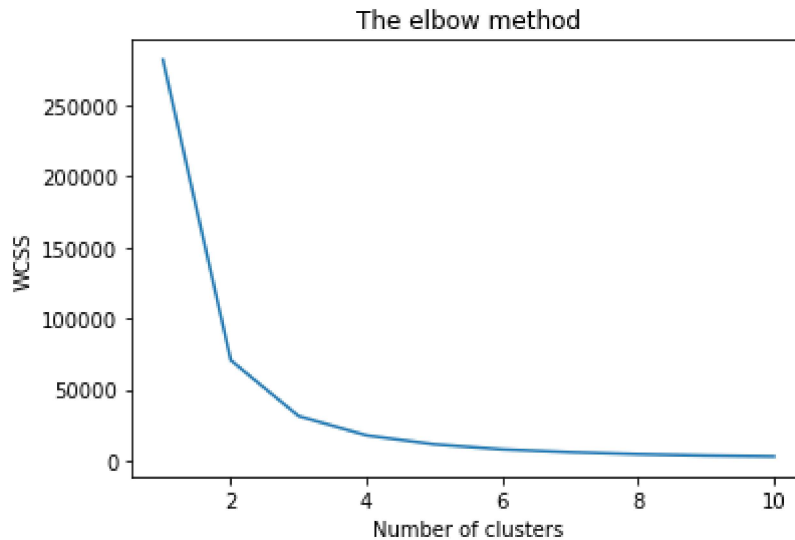
wcss = []

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

#Plotting the results onto a Line graph, allowing us to observe 'The elbow'
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()
```

C:\Users\Amay\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:881: UserWarning: 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_THREADS=1.

```
warnings.warn(
```

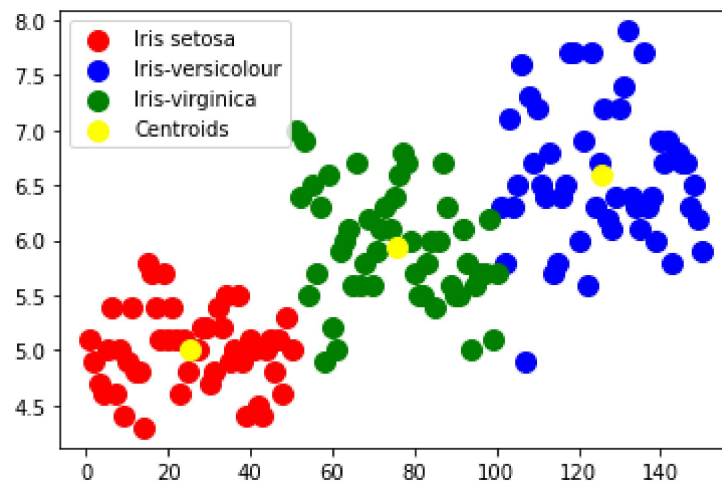


```
In [11]: #Applying kmeans to the dataset/creating the means classifier
kmeans = KMeans (n_clusters = 3, init = "k-means++", max_iter = 300, n_init = 10, ra
y_kmeans = kmeans.fit_predict(x)
```

```
In [12]: #visualising the clusters
plt.scatter(x[y_kmeans == 0, 0], x[y_kmeans == 0, 1], s = 100, c='red', label = "Iri
plt.scatter(x[y_kmeans == 1, 0], x[y_kmeans == 1, 1], s = 100, c= 'blue', label = "I
plt.scatter(x[y_kmeans == 2, 0], x[y_kmeans == 2, 1], s = 100, c='green', label="Iri

#plotting the centroids of the clusters
plt.scatter (kmeans.cluster_centers_[0, 0], kmeans.cluster_centers_[0,1], s= 100, c=
plt.legend()
```

Out[12]: <matplotlib.legend.Legend at 0x2b51e86fbe0>



In []: