

Assignment 6

```
In [1]: import pandas as pd
import numpy as np
```

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

Out[2]:

	sepalength	sepalwidth	petallength	petalwidth	class
0	5.1	3.5	1.4	0.2	Iris-setosa
1	4.9	3.0	1.4	0.2	Iris-setosa
2	4.7	3.2	1.3	0.2	Iris-setosa
3	4.6	3.1	1.5	0.2	Iris-setosa
4	5.0	3.6	1.4	0.2	Iris-setosa
...
145	6.7	3.0	5.2	2.3	Iris-virginica
146	6.3	2.5	5.0	1.9	Iris-virginica
147	6.5	3.0	5.2	2.0	Iris-virginica
148	6.2	3.4	5.4	2.3	Iris-virginica
149	5.9	3.0	5.1	1.8	Iris-virginica

150 rows × 5 columns

```
In [3]: df.dtypes
```

```
Out[3]: sepalength    float64
sepalwidth    float64
petallength    float64
petalwidth    float64
class          object
dtype: object
```

```
In [4]: df['class']=df['class'].astype('category')
df.dtypes
```

```
Out[4]: sepalength    float64
sepalwidth    float64
petallength    float64
petalwidth    float64
class          category
dtype: object
```

```
In [5]: df['class']=df['class'].cat.codes
df
```

Out[5]:

	sepalength	sepalwidth	petallength	petalwidth	class
0	5.1	3.5	1.4	0.2	0
1	4.9	3.0	1.4	0.2	0
2	4.7	3.2	1.3	0.2	0
3	4.6	3.1	1.5	0.2	0
4	5.0	3.6	1.4	0.2	0
...
145	6.7	3.0	5.2	2.3	2
146	6.3	2.5	5.0	1.9	2
147	6.5	3.0	5.2	2.0	2
148	6.2	3.4	5.4	2.3	2
149	5.9	3.0	5.1	1.8	2

150 rows × 5 columns

```
In [6]: df.isnull().sum()
```

```
Out[6]: sepalength    0
sepalwidth    0
petallength    0
petalwidth    0
class         0
dtype: int64
```

```
In [7]: (df <= 0).sum()
```

```
Out[7]: sepalength    0
sepalwidth    0
petallength    0
petalwidth    0
class        50
dtype: int64
```

```
In [8]: print(df.shape)
```

(150, 5)

```
In [9]: # co-relation matrix
def DetectOutlier(df,var):
    Q1 = df[var].quantile(0.25)
    Q3 = df[var].quantile(0.75)
    IQR = Q3 - Q1
    high, low = Q3+1.5*IQR, Q1-1.5*IQR

    print("Highest allowed in variable:", var, high)
    print("lowest allowed in variable:", var, low)

    count = df[(df[var] > high) | (df[var] < low)][var].count()

    print('Total outliers in:',var,':',count)

DetectOutlier(df,'sepallength')

DetectOutlier(df,'sepalwidth')

DetectOutlier(df,'petallength')

DetectOutlier(df,'petalwidth')
```

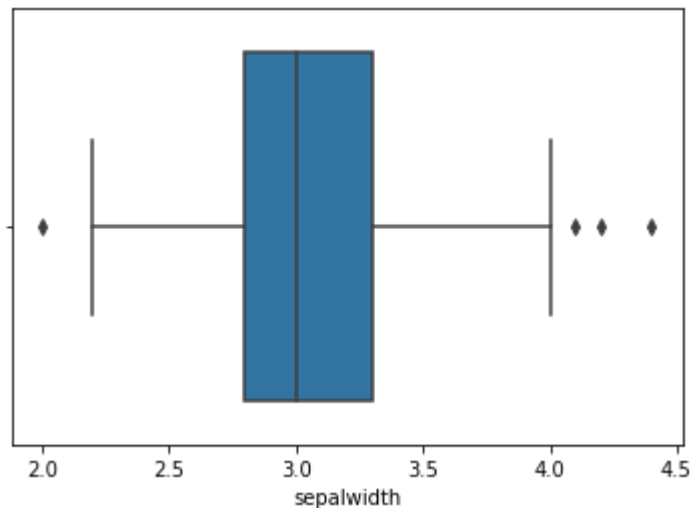
```
Highest allowed in variable: sepallength 8.350000000000001
lowest allowed in variable: sepallength 3.1499999999999986
Total outliers in: sepallength : 0
Highest allowed in variable: sepalwidth 4.05
lowest allowed in variable: sepalwidth 2.05
Total outliers in: sepalwidth : 4
Highest allowed in variable: petallength 10.349999999999998
lowest allowed in variable: petallength -3.6499999999999999
Total outliers in: petallength : 0
Highest allowed in variable: petalwidth 4.05
lowest allowed in variable: petalwidth -1.95
Total outliers in: petalwidth : 0
```

```
In [10]: import seaborn as sns
sns.boxplot(df['sepalwidth'])
```

C:\Users\DELL\AppData\Local\Programs\Python\Python39\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword argument: x. From version 0.12, the only valid positional argument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation.

```
warnings.warn(
```

```
Out[10]: <AxesSubplot:xlabel='sepalwidth'>
```



```
In [11]: def OutlierRemoval(df,var):
    Q1 = df[var].quantile(0.25)
    Q3 = df[var].quantile(0.75)
    IQR = Q3 - Q1
    high, low = Q3+1.5*IQR, Q1-1.5*IQR

    print("Highest allowed in variable:", var, high)
    print("lowest allowed in variable:", var, low)

    count = df[(df[var] > high) | (df[var] < low)][var].count()

    print('Total outliers in:',var,':',count)

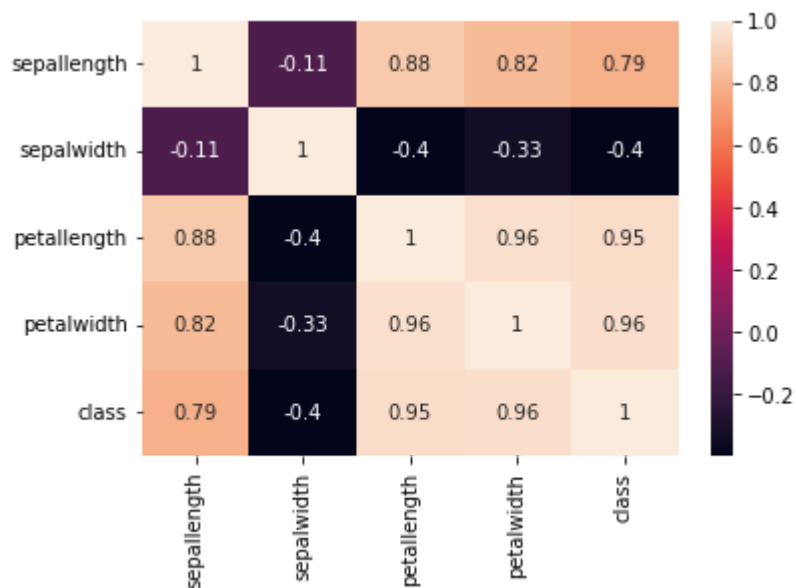
    df = df[((df[var] >= low) & (df[var] <= high))]
    return df
```

```
In [12]: print(df.shape)
df = OutlierRemoval(df, 'sepalwidth')
print(df.shape)
```

```
(150, 5)
Highest allowed in variable: sepalwidth 4.05
lowest allowed in variable: sepalwidth 2.05
Total outliers in: sepalwidth : 4
(146, 5)
```

```
In [13]: import seaborn as sns
sns.heatmap(df.corr(),annot=True)
```

Out[13]: <AxesSubplot:>



```
In [14]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns
```

```
In [15]: # split the data into inputs and outputs
X = df.iloc[:, [0,2,3]].values
y = df.iloc[:, 4].values
```

```
In [16]: # training and testing data
from sklearn.model_selection import train_test_split

# assign test data size 25%
X_train, X_test, y_train, y_test = train_test_split(X,y,test_size= 0.25, random_st
```

```
In [17]: # importing standard scaler  
from sklearn.preprocessing import StandardScaler  
  
# scaling the input data  
sc_X = StandardScaler()  
X_train = sc_X.fit_transform(X_train)  
X_test = sc_X.fit_transform(X_test)
```

```
In [18]: # importing standard scaler  
from sklearn.preprocessing import StandardScaler  
  
# scaling the input data  
sc_X = StandardScaler()  
X_train = sc_X.fit_transform(X_train)  
X_test = sc_X.fit_transform(X_test)
```

```
In [19]: # import Gaussian Naive Bayes classifier  
from sklearn.naive_bayes import GaussianNB  
  
# create a Gaussian Classifier  
classifier1 = GaussianNB()  
  
# training the model  
classifier1.fit(X_train, y_train)  
  
# testing the model  
y_pred1 = classifier1.predict(X_test)
```

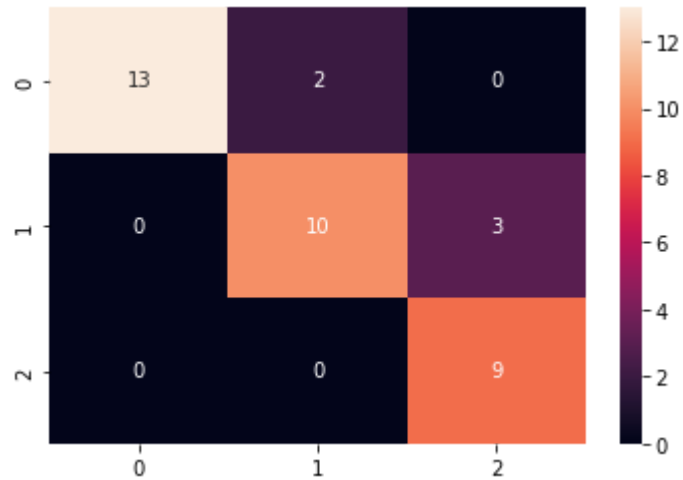
```
In [20]: # importing accuracy score  
from sklearn.metrics import accuracy_score  
  
# printing the accuracy of the model  
print(accuracy_score(y_test,y_pred1))
```

0.8648648648648649

```
In [21]: # importing the required modules
import seaborn as sns
from sklearn.metrics import confusion_matrix

# passing actual and predicted values
cm = confusion_matrix(y_test, y_pred1)

# true write data values in each cell of the matrix
sns.heatmap(cm, annot=True)
plt.savefig('confusion.png')
```



```
In [22]: # importing classification report
from sklearn.metrics import classification_report

# printing the report
print(classification_report(y_test, y_pred1))
```

	precision	recall	f1-score	support
0	1.00	0.87	0.93	15
1	0.83	0.77	0.80	13
2	0.75	1.00	0.86	9
accuracy			0.86	37
macro avg	0.86	0.88	0.86	37
weighted avg	0.88	0.86	0.87	37