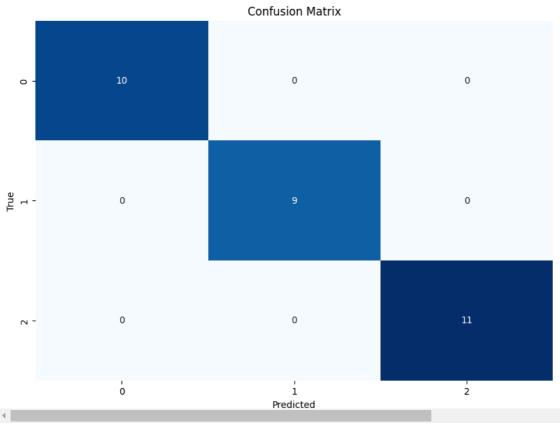
```
# 1.
        Load the basic libraries and packages
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import pydot
from sklearn.tree import export graphviz
from sklearn import metrics
from sklearn import datasets
from sklearn.metrics import confusion_matrix
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix
# 2.
        Load the dataset
iris = datasets.load_iris()
class_name = iris.target_names
data = pd.DataFrame({"Sepal Length" : iris.data[:,0],
                      "Sepal Width" : iris.data[:,1],
                      "Petal Length" : iris.data[:,2],
                      "Petal Width" : iris.data[:,3] ,
                      "Species" : iris.target ,
                      "Species Name" : class_name[iris.target]});
data.head()
\overline{z}
         Sepal Length Sepal Width Petal Length Petal Width Species Species Name
                                                                                           \blacksquare
                   5.1
                                3.5
                                               1.4
                                                             0.2
                                                                        0
                                                                                  setosa
                                                                                           ıl.
      1
                  4.9
                                3.0
                                               1.4
                                                             0.2
                                                                        0
                                                                                  setosa
                   4.7
                                3.2
                                               1.3
                                                             0.2
                                                                        0
                                                                                  setosa
      3
                   4.6
                                3.1
                                               1.5
                                                             0.2
                                                                        0
                                                                                  setosa
                   5.0
                                                             0.2
                                 3.6
                                               1.4
                                                                                  setosa
 Next steps:
              Generate code with data
                                          View recommended plots
                                                                         New interactive sheet
# 3.
        Separate the feature and prediction value columns
X = data.drop(["Species", "Species Name"], axis=1)
y = data["Species"]
# 4. Splitting the Training and Testing Data
 X\_Train \ , \ X\_Test \ , \ y\_train \ , \ Y\_Test = train\_test\_split(X, \ y \ , \ test\_size=0.2 \ , \ random\_state=42) 
# 5. Creating a Model
forest = RandomForestClassifier(n_estimators=200 , random_state = 42)
# 6. Training a Model
forest.fit(X_Train,y_train)
                       RandomForestClassifier
     RandomForestClassifier(n_estimators=200, random_state=42)
# 7. Predicitng the Output Using model
Y_Predicted = forest.predict(X_Test)
# 8. Measuring the Accuracy
metrics.accuracy_score(Y_Test,Y_Predicted)
→<u>*</u> 1.0
```

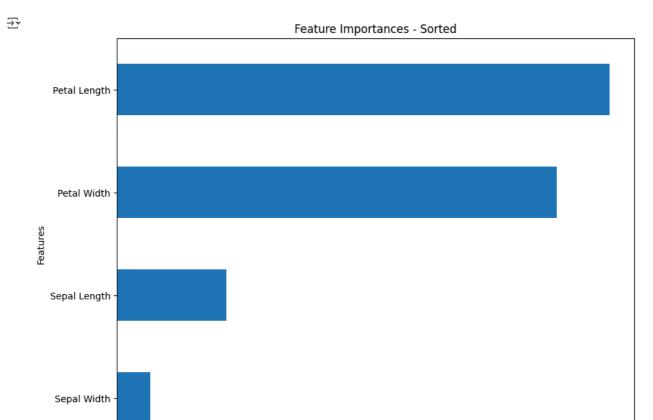
```
# 9. Creating a Confusion Matrix
```

## → Text(0.5, 1.0, 'Confusion Matrix')

plt.title('Confusion Matrix')



```
# Sort the feature importances in descending order
sorted_importances = feature_importances.sort_values(ascending=True)
# Plot the sorted feature importances
sorted_importances.plot(kind='barh', figsize=(10, 8))
# Add labels and title
plt.xlabel('Importance')
plt.ylabel('Features')
plt.title('Feature Importances - Sorted')
plt.show()
```



0.2

Importance

0.3

0.4

0.1

0.0