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Course: Bil470

Importing the Dependincies

```
import numpy as np
In [39]:
         import pandas as pd
         from sklearn import svm
         from sklearn import metrics
         from sklearn.metrics import accuracy score, precision score
          from sklearn.metrics import confusion matrix
          from sklearn.metrics import precision recall fscore support
          from sklearn.metrics import classification_report
          import matplotlib.pyplot as plt
          import seaborn as sns
          from sklearn.metrics import roc curve, auc
          from sklearn.preprocessing import label_binarize
          from sklearn.preprocessing import LabelEncoder
          from sklearn.model selection import train_test_split
          from dt import DecisionTreeClassifier
          from tabulate import tabulate
          from sklearn.model selection import learning curve
```

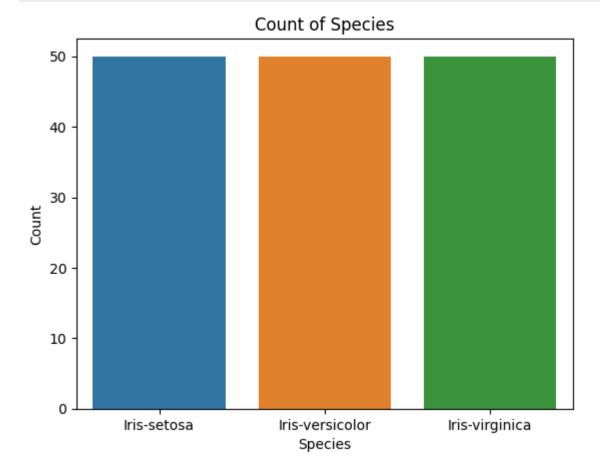
Exploratory Data Analysis (EDA) for IRIS

```
In [2]: iris_data = pd.read_csv('Iris.csv')
          iris data.head()
            Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm
Out[2]:
                                                                                 Species
             1
                            5.1
                                           3.5
                                                           1.4
                                                                          0.2 Iris-setosa
             2
                            4.9
                                           3.0
                                                                          0.2 Iris-setosa
                                                           1.4
             3
                            4.7
                                           3.2
          2
                                                           1.3
                                                                          0.2 Iris-setosa
                                                                          0.2 Iris-setosa
          3
                            4.6
                                           3.1
                                                           1.5
            5
                            5.0
                                           3.6
                                                           1.4
                                                                          0.2 Iris-setosa
         iris data = iris data.drop('Id',axis=1)
In [4]: iris_data.describe()
```

Out[4]:

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

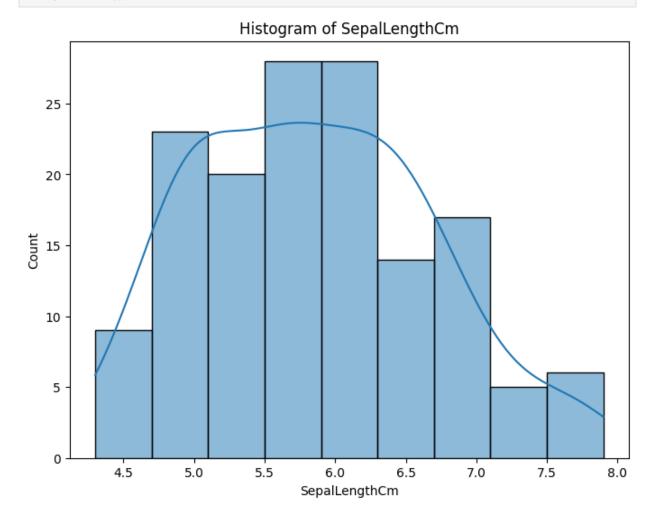
```
In [5]: sns.countplot(data=iris_data, x='Species')
    plt.xlabel('Species')
    plt.ylabel('Count')
    plt.title('Count of Species')
    plt.show()
```



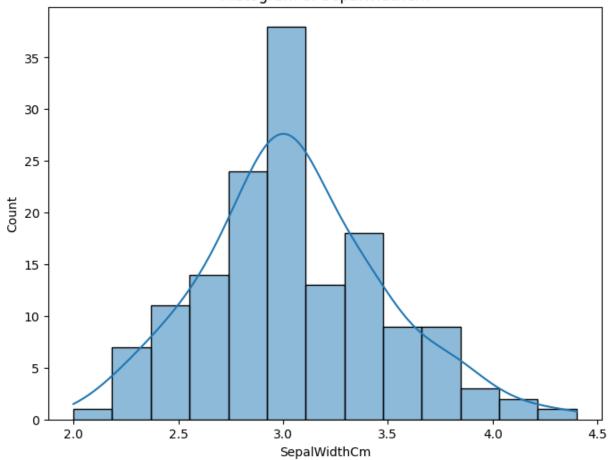
```
In [6]: features = iris_data.columns[:-1]

for feature in features:
    plt.figure(figsize=(8, 6))
    sns.histplot(data= iris_data, x=feature, kde=True) #kde is density
    plt.xlabel(feature)
    plt.ylabel('Count')
    plt.title(f'Histogram of {feature}')
```

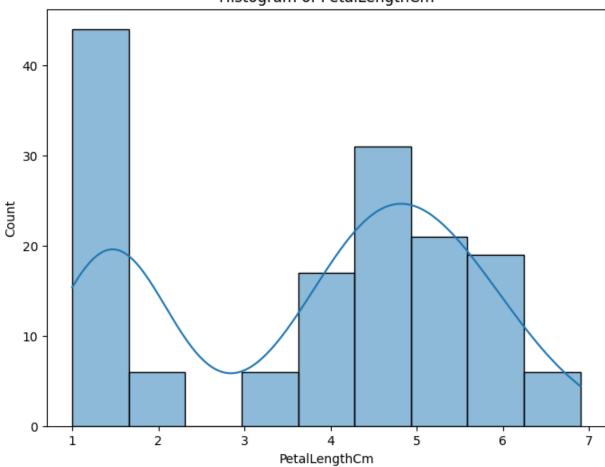
plt.show()



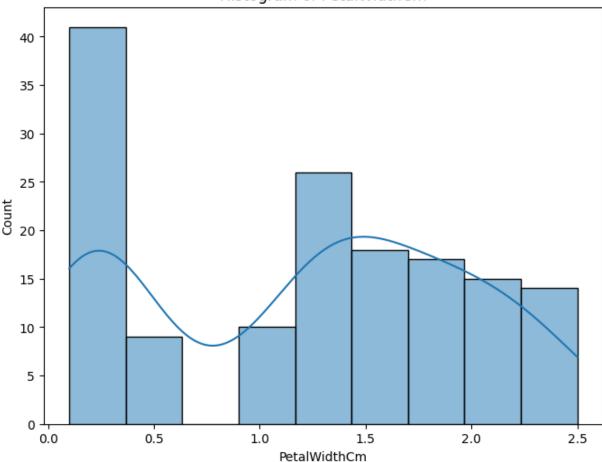
Histogram of SepalWidthCm



Histogram of PetalLengthCm

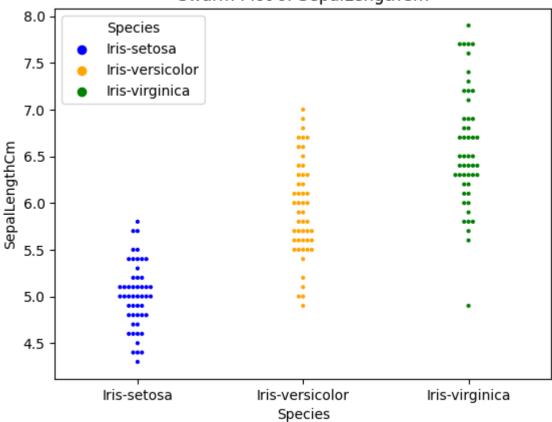


Histogram of PetalWidthCm

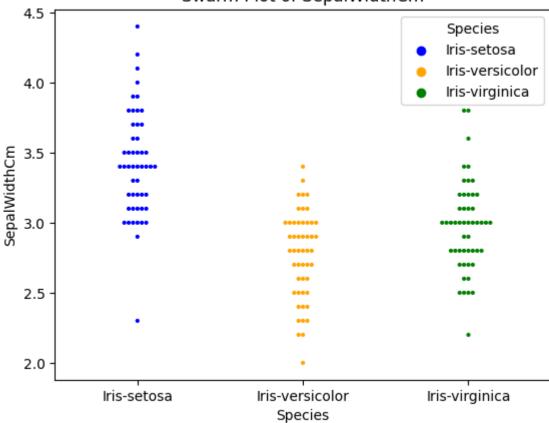


```
In [7]: species_colors = {'setosa': 'blue', 'versicolor': 'orange', 'virginica': 'green'} 
for feature in features:
    plt.figure()
    sns.swarmplot(data=iris_data, x='Species', y=feature, hue='Species', palette=speciplt.xlabel('Species')
    plt.ylabel(feature)
    plt.title(f'Swarm Plot of {feature}')
    plt.show()
```

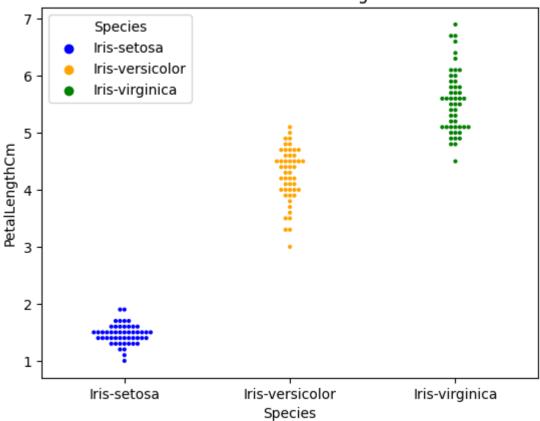




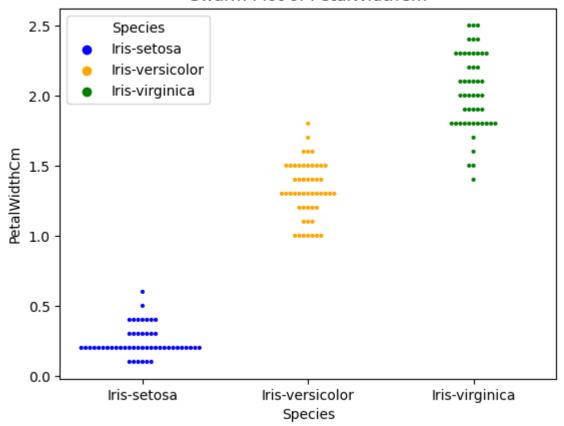




Swarm Plot of PetalLengthCm

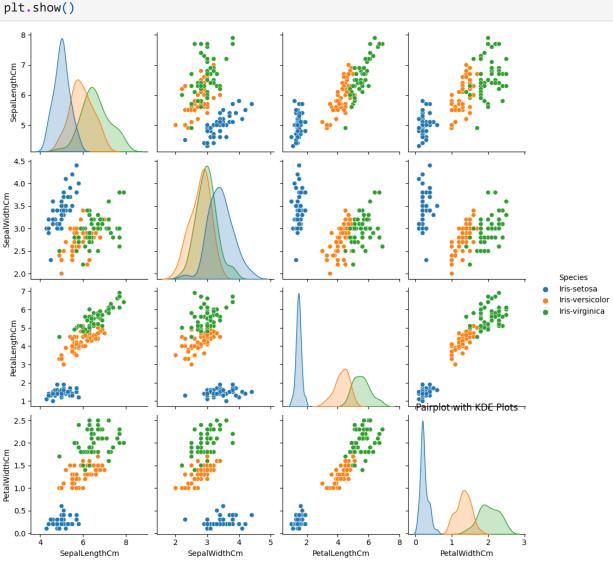


Swarm Plot of PetalWidthCm



In [8]: sns.pairplot(data=iris_data, hue='Species', diag_kind='kde')
plt.title('Pairplot with KDE Plots')





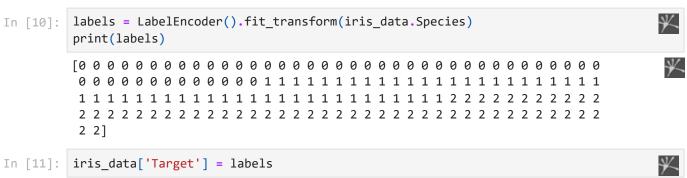
```
In [9]: correlation_matrix = iris_data.corr()

plt.figure(figsize=(15, 10))
    sns.heatmap(correlation_matrix, annot=True, linewidths=0.5, fmt='.2f')
    plt.title("Correlation Matrix Heatmap")
    plt.show()
```

C:\Users\Berkay\AppData\Local\Temp\ipykernel_16236\2869641057.py:1: FutureWarning: e default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_o nly to silence this warning.

correlation matrix = iris data.corr()





Train the classifier

```
In [12]: clf = DecisionTreeClassifier(max_depth=5)
In [13]: X = iris_data.drop(['Species','Target'], axis=1)
    Y = iris_data['Target']
In [14]: X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size = 0.2, shuffle
In [15]: X_train_list = X_train.values.tolist()
    Y_train_list = Y_train.values.tolist()
In [16]: clf.fit(X_train_list,Y_train_list)
```

```
<dt.Node at 0x12e8b76e860>
Out[16]:
In [17]: clf.print_tree()
       X 2 <= 1.9 ? 0.32464769647696473
        left:0
        right:X_3 <= 1.7 ? 0.3870826209910089
        left:X 2 <= 4.9 ? 0.06530864197530856
          left:X 3 <= 1.6 ? 0.04875000000000007
             left:1
             right:2
          left:2
             left:1
                   right:2
        right:X_2 <= 4.8 ? 0.016557097638178662
          left:1
             right:2
          right:2
```

Predict Class of Train values

```
print(Y_train.values.tolist())
In [18]:
         predictOfTrain = clf.predict(X train.values.tolist())
         print(predictOfTrain)
        [1, 0, 0, 2, 1, 2, 2, 1, 2, 1, 1, 2, 0, 0, 2, 1, 2, 0, 2, 1, 2, 1, 0, 2, 1, 0, 0, 0]
        2, 1, 1, 2, 1, 0, 0, 0, 2, 1, 1, 0, 0, 0, 2, 2, 1, 2, 2, 1, 1, 1, 2, 1, 0, 0, 2, 0,
        1, 2, 1, 0, 2, 0, 1, 2, 1, 0, 1, 2, 0, 1, 2, 1, 2, 1, 0, 2, 2, 0, 2, 0, 1, 2, 1, 0,
        2, 1, 2, 1, 0, 1, 2, 2, 2, 0, 0, 1, 1, 1, 1, 0, 2, 2, 0, 1, 0, 0, 1, 1, 0, 0, 0, 2,
        2, 1, 2, 2, 0, 1, 0, 1]
        2, 1, 1, 2, 1, 0, 0, 0, 2, 1, 1, 0, 0, 0, 2, 2, 1, 2, 2, 1, 1, 1, 2, 1, 0, 0, 2, 0,
        1, 2, 1, 0, 2, 0, 1, 2, 1, 0, 1, 2, 0, 1, 2, 1, 2, 1, 0, 2, 2, 0, 2, 0, 1, 2, 1, 0,
        2, 1, 2, 1, 0, 1, 2, 2, 2, 0, 0, 1, 1, 1, 1, 0, 2, 2, 0, 1, 0, 0, 1, 1, 0, 0, 0, 2,
        2, 1, 2, 2, 0, 1, 0, 1]
        differences = [y true != y pred for y true, y pred in zip(Y train, predictOfTrain)]
In [19]:
         difference rate = sum(differences) / len(Y train)
         print(difference rate)
                                                                                     ¥
        0.0
```

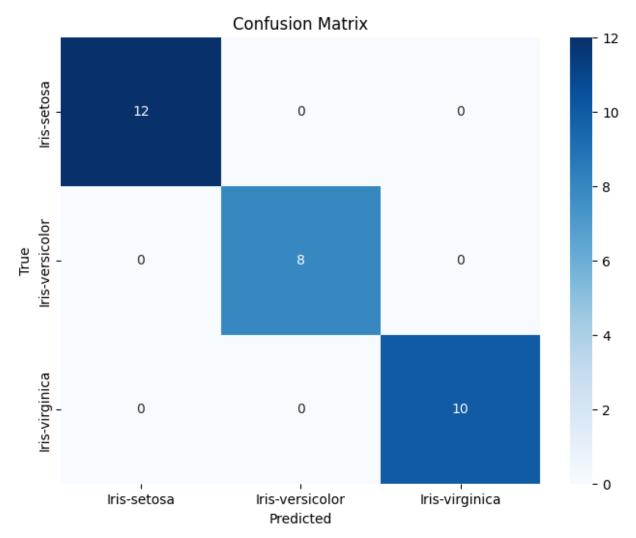
Predict Class of Test values

```
print(difference_rate)
0.0
```

Results

Confusion Matrix of Test

```
In [22]: cm = confusion_matrix(Y_test.values.tolist(), predictOfTest)
         # Print the confusion matrix
         print("Confusion Matrix:")
         print(cm)
         Confusion Matrix:
         [[12 0 0]
          [0 8 0]
          [ 0 0 10]]
         class_labels = list(LabelEncoder().fit(iris_data.Species).classes_)
In [23]:
         plt.figure(figsize=(8, 6))
         sns.heatmap(cm, annot=True, cmap='Blues', fmt='d', xticklabels=class_labels, yticklabe
         plt.xlabel('Predicted')
         plt.ylabel('True')
         plt.title('Confusion Matrix')
         plt.show()
```



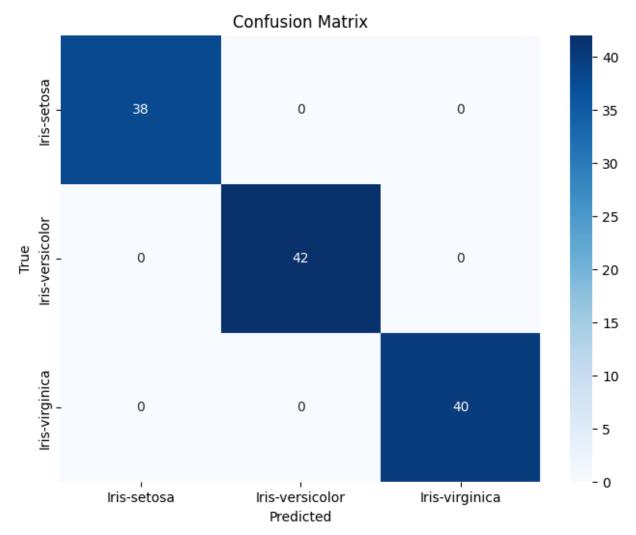
Confusion Matrix of Train

```
In [24]: cm = confusion_matrix(Y_train.values.tolist(), predictOfTrain)

# Print the confusion matrix
print("Confusion Matrix:")
print(cm)

Confusion Matrix:
[[38 0 0]
       [ 0 42 0]
       [ 0 0 40]]

In [25]: plt.figure(figsize=(8, 6))
sns.heatmap(cm, annot=True, cmap='Blues', fmt='d', xticklabels=class_labels, yticklabe() plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()
```



F1-Score

```
In [26]:
          report = classification_report(Y_train.values.tolist(), predictOfTrain, target_nam
          print("Classification Report:")
          print("-" * 60)
          print(report)
         Classification Report:
                                        recall f1-score
                           precision
                                                            support
              Iris-setosa
                               1.000
                                         1.000
                                                    1.000
                                                                 38
         Iris-versicolor
                               1.000
                                         1.000
                                                    1.000
                                                                 42
           Iris-virginica
                               1.000
                                          1.000
                                                    1.000
                                                                 40
                                                    1.000
                                                                120
                 accuracy
                macro avg
                               1.000
                                         1.000
                                                    1.000
                                                                120
             weighted avg
                               1.000
                                          1.000
                                                    1.000
                                                                120
```

F1-Score, Precision, Recall and Accuracy of Train

```
precision, recall, f1, support = precision_recall_fscore_support(Y_train, predict0
In [27]:
         accuracy = accuracy score(Y train, predictOfTrain)
         for i, label in enumerate(class labels):
             print(f"Class: {label}")
             print(f"Precision: {precision[i]:.2f}")
             print(f"Recall: {recall[i]:.2f}")
             print(f"F1-Score: {f1[i]:.2f}")
             print("")
         print(f"Overall Accuracy: {accuracy:.2f}")
         Class: Iris-setosa
         Precision: 1.00
         Recall: 1.00
         F1-Score: 1.00
         Class: Iris-versicolor
         Precision: 1.00
         Recall: 1.00
         F1-Score: 1.00
         Class: Iris-virginica
         Precision: 1.00
         Recall: 1.00
         F1-Score: 1.00
         Overall Accuracy: 1.00
```

F1-Score, Precision, Recall and Accuracy of Train

Class: Iris-setosa
Precision: 1.00
Recall: 1.00
F1-Score: 1.00

Class: Iris-versicolor
Precision: 1.00
Recall: 1.00
F1-Score: 1.00

Class: Iris-virginica
Precision: 1.00
Recall: 1.00
F1-Score: 1.00

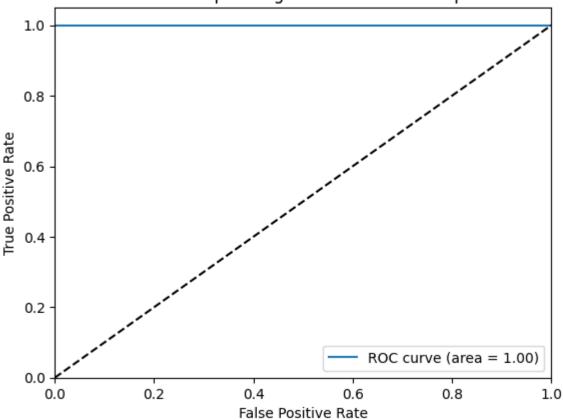
Overall Accuracy: 1.00



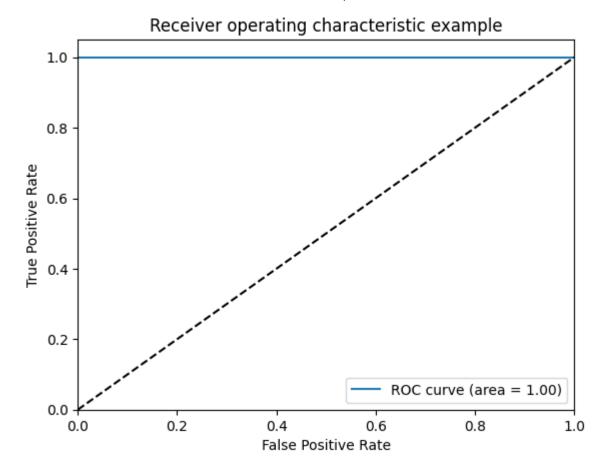
Plot of ROC Curve (Test/Train) and Value of AUC (Test/Train)

```
In [29]: y = label_binarize(pd.Series(Y_test.values.tolist()), classes=[0, 1, 2])
                                                                                             ¥
         y2 = label_binarize(pd.Series(predictOfTest), classes=[0, 1, 2])
         fpr = dict()
         tpr = dict()
          roc_auc = dict()
         for i in range(3):
          fpr[i], tpr[i], _ = roc_curve(y[:, i], y2[:, i])
          roc_auc[i] = auc(fpr[i], tpr[i])
         fpr["micro"], tpr["micro"], _ = roc_curve(y.ravel(), y2.ravel())
          roc_auc["micro"] = auc(fpr["micro"], tpr["micro"])
          plt.figure()
          plt.plot(fpr[2], tpr[2], label='ROC curve (area = %0.2f)' % roc auc[2])
         plt.plot([0, 1], [0, 1], 'k--')
          plt.xlim([0.0, 1.0])
          plt.ylim([0.0, 1.05])
         plt.xlabel('False Positive Rate')
         plt.ylabel('True Positive Rate')
          plt.title('Receiver operating characteristic example')
          plt.legend(loc="lower right")
         plt.show()
```





```
In [30]: y = label_binarize(pd.Series(Y_train.values.tolist()), classes=[0, 1, 2])
          y2 = label binarize(pd.Series(predictOfTrain), classes=[0, 1, 2])
          fpr = dict()
          tpr = dict()
          roc_auc = dict()
          for i in range(3):
           fpr[i], tpr[i], _ = roc_curve(y[:, i], y2[:, i])
           roc_auc[i] = auc(fpr[i], tpr[i])
          fpr["micro"], tpr["micro"], _ = roc_curve(y.ravel(), y2.ravel())
roc_auc["micro"] = auc(fpr["micro"], tpr["micro"])
          plt.figure()
          plt.plot(fpr[2], tpr[2], label='ROC curve (area = %0.2f)' % roc_auc[2])
          plt.plot([0, 1], [0, 1], 'k--')
          plt.xlim([0.0, 1.0])
          plt.ylim([0.0, 1.05])
          plt.xlabel('False Positive Rate')
          plt.ylabel('True Positive Rate')
          plt.title('Receiver operating characteristic example')
          plt.legend(loc="lower right")
          plt.show()
```



Evaluation

Kendi yazdığım karar ağacı modelinin üzerinden eğittiğim iris data seti test ve train verileri üzerinde hatasız çalışmıştır. Bu hatasız çalışmanın sebepleri arasında verinin kolayca ayırt edilebilen özelliklere sahip olması ve yüksek performans elde etmek mümkün olması, train veri setinin overfitlenmiş olabilmesi ve test veri setinin çok büyük olmaması sonucu ayırt edilememiş olması veya modelin bu eğitimde tam doğru noktalardan karar ağacını ayırması olabilir.

Ideal Depth for DT

Kendi modelimdeki derinlik 5, bu değeri öylesine seçmiştim fakat aşağıdaki grafikte de görüldüğü üzere ideal derinlik 2 çıktı. Grafiğe bakıldığında görebileceğimiz gibi derinlik 2'den sonra daima aynı. Bu gibi durumlarda (aynı performansı gösteren) en basit modeli seçmek en mantıklıdır.

In [34]: clf.print_tree()

```
X 2 <= 1.9 ? 0.32464769647696473
 left:0
 right:X_3 <= 1.7 ? 0.3870826209910089
 left:X_2 <= 4.9 ? 0.06530864197530856
    left:X 3 <= 1.6 ? 0.04875000000000007
       left:1
       right:2
    left:2
        left:1
               right:2
  right:X_2 <= 4.8 ? 0.016557097638178662
    left:1
       right:2
    right:2
Level 0: Root node (X_2 \le 1.9)
Level 1: Left branch (0.32464769647696473)
Level 1: Right branch (X_3 \le 1.7)
Level 2: Left branch (X_2 \le 4.9)
Level 3: Left branch (X_3 \le 1.6)
Level 4: Left branch (1)
Level 4: Right branch (2)
Level 3: Right branch (X_3 \le 1.5)
Level 4: Left branch (2)
Level 4: Right branch (X_0 \le 6.7)
Level 5: Left branch (1)
Level 5: Right branch (2)
Level 2: Right branch (X_2 \le 4.8)
Level 3: Left branch (X_0 \le 5.9)
Level 4: Left branch (1)
Level 4: Right branch (2)
Level 3: Right branch (2)
```

Level 0: X_2 <= 1.9



```
Level 1: 0.3246 Level 1: X_3 <= 1.7
                                 Level 2: X_2 <= 4.8
           Level 2: X 2 <= 4.9
             Level 3: X_3 \leftarrow 1.6 Level 3: X_3 \leftarrow 1.5 Level 3: X_0 \leftarrow 5.9
   Level 3: 2
     Level 4: 1
                           Level 4: 2
                                             Level 4: 1
   Level 4: 2
depth_values = [1, 2, 3, 4, 5, 6, 7, 8, 9, 10]
accuracy scores = []
precision_scores = []
for depth in depth_values:
   model = DecisionTreeClassifier(max_depth=depth)
   model.fit(X train.values.tolist(), Y train.values.tolist())
   y pred = model.predict(X test.values.tolist())
   accuracy = accuracy_score(Y_test, y_pred)
   precision = precision_score(Y_test, y_pred, average='macro')
   accuracy scores.append(accuracy)
   precision scores.append(precision)
plt.figure(figsize=(10, 6))
plt.plot(depth_values, accuracy_scores, label='Accuracy')
plt.plot(depth_values, precision_scores, label='Precision')
plt.xlabel('Max Depth')
```

plt.ylabel('Score')

plt.legend()
plt.grid(True)
plt.show()

plt.title('Model Performance for Different Max Depths')

