Data Set: titanic

Full Tree

graph

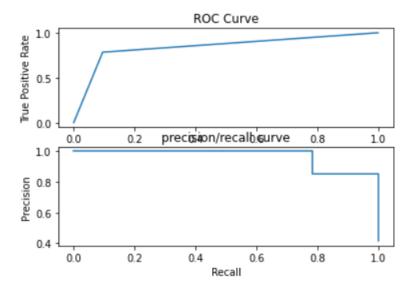
```
3 import pandas as pd
import matplotlib.pyplot as plt
# Set a random seed
      import random
from sklearn.model_selection import train_test_split
from sklearn import tree
from sklearn.tree import DecisionTreeClassifier
      from sklearn.tree import export_graphviz from sklearn.metrics import precision_recall_curve,precision_score,recall_score,accuracy_score,confusion_matrix,roc_curve,classification_report from graphviz import Source
      import graphviz
from IPython.display import display,SVG
      # Load the dataset
in_file = 'titanic.csv'
full_data = pd.read_csv(in_file)
      outcomes = full_data['Survived']
features_raw = full_data.drop('Survived', axis = 1)
features = pd.get_dummles(features_raw)
features = features.fillna(0.0)
      X_train, X_test, y_train, y_test = train_test_split(features, outcomes, test_size=0.2, random_state=42)
# Import the classifier from sklearn
     # Import the classifier from skiearn
model - PecisionTreeClassifier(max_depth=6, min_samples_leaf=6, min_samples_split=10)
model.fit(X_train,y_train)
# Making predictions
y_train_pred = model.predict(X_train)
y_test_pred = model.predict(X_test)
      accuracy = accuracy_score(y_test, y_test_pred)
      recall = recall_score(y_test, y_test_pred)
recall = recall_score(y_test, y_test_pred)
recall = recall_score(y_test, y_test_pred)
classification = classification_report(y_test, y_test_pred)
confusion = confusion_matrix(y_test, y_test_pred)
print('The accuracy is', accuracy)
print('The precision is', precision)
rint('The recall is' recall)
      print('The recall is',recall)
print('The classification report is\n',classification)
      print('The confusion matrix is',confusion)
      fpr, tpr, thersholds = roc_curve(y_test, y_test_pred)
pre, rec, thresholds2 = precision_recall_curve(y_test, y_test_pred)
       plt.figure()
       plt.subplot(2,1,1)
      plt.plot(fpr, tpr)
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
      plt.title('ROC Curve')
       plt.subplot(2,1,2)
      plt.step(rec, pre)
       plt.xlabel('Recall')
      plt.ylabel('Precision')
plt.title('precision/recall curve')
```

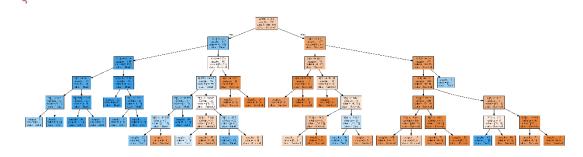
graph = Source(tree.export_graphviz(model, out_file=None, class_names=['Survived','Dead'], impurity=False,filled=True))

The accuracy is 0.8547486033519553 The precision is 0.8529411764705882 The recall is 0.7837837837837838 The classification report is

	precision	recall	f1-score	support
0	0.86	0.90	0.88	105
1	0.85	0.78	0.82	74
accuracy			0.85	179
macro avg	0.85	0.84	0.85	179
weighted avg	0.85	0.85	0.85	179

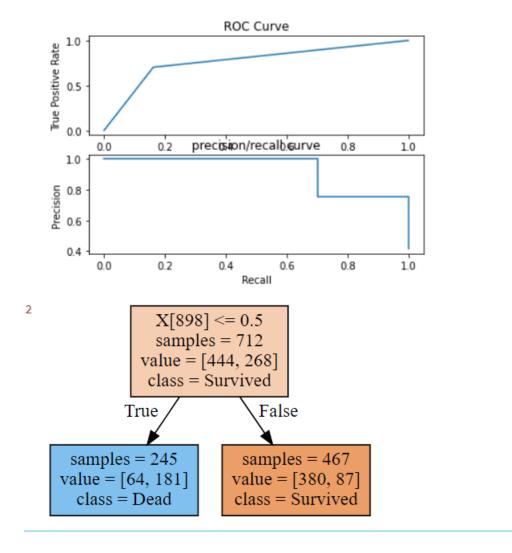
The confusion matrix is [[95 10] [16 58]]





```
12 model1 = DecisionTreeClassifier(max_depth=1, min_samples_leaf=6, min_samples_split=10)
   model1.fit(X_train,y_train)
   # Making predictions
   y_test_pred1 = model1.predict(X_test)
   1. The accuracy of your model on the test data
   2. The precision and recall values
   3. A classification report (scikit-learn has a function that can create this for you)
   4. The confusion matrix for this experiment
   5. An ROC curve
   6. A Precision/Recall curve
   accuracy1 = accuracy_score(y_test, y_test_pred1)
   precision1 = precision_score(y_test, y_test_pred1)
   recall1 = recall_score(y_test, y_test_pred1)
   classification1 = classification_report(y_test, y_test_pred1)
   confusion1 = confusion_matrix(y_test, y_test_pred1)
   print('The accuracy is', accuracy1)
print('The precision is', precision1)
   print('The recall is',recall1)
   print('The classification report is\n',classification1)
   print('The confusion matrix is',confusion1)
   fpr1, tpr1, thersholds1 = roc_curve(y_test, y_test_pred1)
   pre1, rec1, thresholds21 = precision_recall_curve(y_test, y_test_pred1)
   plt.figure()
   plt.subplot(2,1,1)
   plt.plot(fpr1, tpr1)
   plt.xlabel('False Positive Rate')
```

```
plt.ylabel('True Positive Rate')
plt.title('ROC Curve')
plt.subplot(2,1,2)
plt.step(rec1, pre1)
plt.xlabel('Recall')
plt.ylabel('Precision')
plt.title('precision/recall curve')
plt.show()
graph = Source(tree.export_graphviz(model1, out_file=None, class_names=['Survived','Dead'], impurity=F
#graph.render(view=True, format="pdf", filename="decisiontree_depth1")
The accuracy is 0.7821229050279329
The precision is 0.7536231884057971
The recall is 0.7027027027027027
The classification report is
              precision
                          recall f1-score support
          0
                  0.80
                        0.84
                                    0.82
                                               105
                  0.75
                        0.70
                                   0.73
                                                74
                                     0.78
                                               179
   accuracy
                 0.78
                          0.77
   macro avg
                                    0.77
                                                179
                0.78
                           0.78
                                     0.78
                                               179
weighted avg
The confusion matrix is [[88 17]
 [22 52]]
```



Compare

When using one level depth decision tree, the accurate value of the result is lower. Judging from the conflict matrix, the correct result is not as good as the full depth decision tree, and the area of the ROC curve is not as good as the full depth decision tree, so the full depth decision tree will have better prediction results