Decision Tree Classification on Hepatitis Dataset

```
In [1]: import pandas as pd
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import confusion_matrix, precision_score, recall_score, f
from sklearn.metrics import classification_report
```

Data Loading

```
In [2]: url = 'https://github.com/rashakil-ds/Public-Datasets/raw/main/hepatitis.csv'
    df = pd.read_csv(url)
    df.head()
```

Out[2]:

	Class	AGE	SEX	STEROID	ANTIVIRALS	FATIGUE	MALAISE	ANOREXIA	LIVER BIG	LIVER FIRM	P/
0	0	30	2	1.0	2	2	2	2	1.0	2.0	
1	0	50	1	1.0	2	1	2	2	1.0	2.0	
2	0	78	1	2.0	2	1	2	2	2.0	2.0	
3	0	31	1	NaN	1	2	2	2	2.0	2.0	
4	0	34	1	2.0	2	2	2	2	2.0	2.0	
4											

```
In [3]: |df.isnull().sum()
Out[3]: Class
                              0
         AGE
                              0
         SEX
                              0
         STEROID
                              1
                              0
         ANTIVIRALS
                              0
         FATIGUE
        MALAISE
                              0
         ANOREXIA
                              0
                              9
         LIVER BIG
         LIVER FIRM
                             10
                              4
         SPLEEN PALPABLE
         SPIDERS
                              4
                              4
         ASCITES
        VARICES
                              4
                              5
         BILIRUBIN
                             28
         ALK PHOSPHATE
         SGOT
                              3
                             15
         ALBUMIN
         PROTIME
                             66
        HISTOLOGY
                              0
         dtype: int64
```

Check the missing values

```
In [4]: missing_values = df.isnull().sum()
print(missing_values)
```

```
0
Class
AGE
                     0
SEX
                     0
STEROID
                     1
ANTIVIRALS
                     0
                     0
FATIGUE
MALAISE
                     0
ANOREXIA
                     0
LIVER BIG
                     9
LIVER FIRM
                    10
SPLEEN PALPABLE
                     4
                     4
SPIDERS
ASCITES
                     4
VARICES
                     4
BILIRUBIN
                     5
ALK PHOSPHATE
                    28
                     3
SGOT
                    15
ALBUMIN
PROTIME
                    66
HISTOLOGY
                     0
dtype: int64
```

```
In [5]: df.dropna(inplace = True)
```

Split the dataset into features (X) and Target Variable (y)

```
In [6]: X = df.drop('Class', axis = 1)
y = df['Class']
```

Convert the target variable to binary(1 for positive, 0 for negative)

```
In [7]: y = y.apply(lambda x: 1 if x == 1 else 0)
```

Decision Tree Model

```
In [8]: # Split the dataset into training and testing sets
   X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, rando
   # Build a Decision Tree classifier
   clf = DecisionTreeClassifier()

# Train the model on the training set
   clf.fit(X_train, y_train)
```

Out[8]: v DecisionTreeClassifier DecisionTreeClassifier()

Make predictions on the testing set

```
In [9]: clf.score(X_test, y_test)
```

Out[9]: 0.83333333333333334

Evaluate the model

```
In [10]: y_pred = clf.predict(X_test)
```

Evaluate the model using various matrics

```
In [11]: conf_matrix = confusion_matrix(y_test, y_pred)
    prec_score = precision_score(y_test, y_pred)
    rec_score = recall_score(y_test, y_pred)
    f1 = f1_score(y_test, y_pred)
    roc_auc = roc_auc_score(y_test, y_pred)
```

Print the evaluate results

```
In [12]: print('Confusion Matrix:\n', conf_matrix)
    print('Precision Score: ', prec_score)
    print('Recall Score: ', rec_score)
    print('F1 Score: ', f1)
    print('AUC-ROC Score: ', roc_auc)
```

Confusion Matrix:

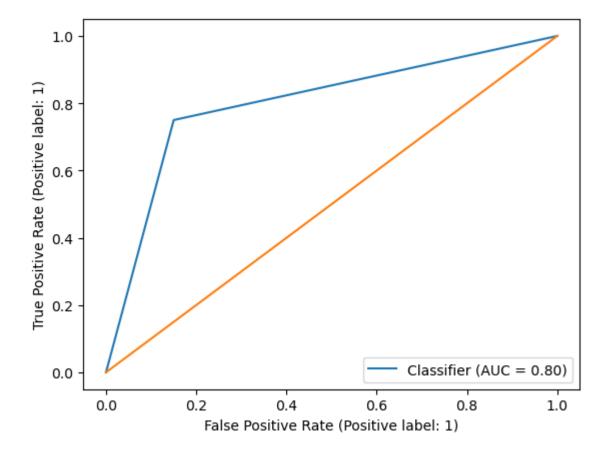
[[17 3] [1 3]]

Precision Score: 0.5
Recall Score: 0.75
F1 Score: 0.6
AUC-ROC Score: 0.8

Classifier for AUC

In [13]: from sklearn.metrics import RocCurveDisplay
import matplotlib.pyplot as plt
RocCurveDisplay.from_predictions(y_test, y_pred)
plt.plot([0,1],[0,1])

Out[13]: [<matplotlib.lines.Line2D at 0x1a04bb3e710>]



Results and Analysis

Summary of Evaluation Metrics

The model exhibits a decent performance with a Precision of 0.5, indicating a balanced prediction of positive outcomes. A Recall of 0.75 suggests that the model captures a substantial portion of the actual positive cases. The F1 Score of 0.6 signifies a reasonable balance between Precision and Recall. An AUC-ROC Score of 0.8 indicates a good ability to discriminate between positive and negative instances.

Strengths and Weaknesses of the Decision Tree Model:

Strengths: Decision trees are interpretable and suitable for simple decision-making scenarios. They handle both numerical and categorical data well. Decision trees provide insights into feature importance.

Weaknesses: The model may be sensitive to noisy data and prone to overfitting. Decision trees might not capture complex relationships in the data as effectively as more sophisticated models. They may struggle with class imbalances.

Recommendations: Consider tuning hyperparameters to potentially improve model performance. Explore ensemble methods or alternative algorithms to address weaknesses.