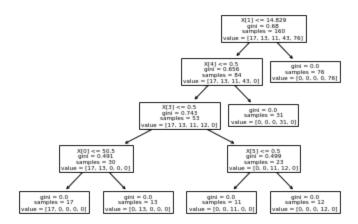
```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score, confusion_matrix
import matplotlib.pyplot as plt
from sklearn import tree
```

Desicion Tree Classification

```
In [2]:
         #Load the dataset
         url = 'https://cf-courses-data.s3.us.cloud-object-storage.appdomain.cloud/IBMDeveloperS
         df = pd.read csv(url)
In [3]:
         df.head()
                          BP
                              Cholesterol Na_to_K
Out[3]:
           Age Sex
                                                  Drug
         0
             23
                  F
                        HIGH
                                   HIGH
                                           25.355 drugY
         1
             47
                  М
                        LOW
                                   HIGH
                                          13.093 drugC
         2
             47
                         LOW
                                          10.114 drugC
                  М
                                   HIGH
         3
             28
                  F NORMAL
                                   HIGH
                                           7.798 drugX
                  F
             61
                         LOW
                                   HIGH
                                          18.043 drugY
In [4]:
         df.info()
         <class 'pandas.core.frame.DataFrame'>
         RangeIndex: 200 entries, 0 to 199
         Data columns (total 6 columns):
                           Non-Null Count Dtype
          #
              Column
          0
              Age
                           200 non-null
                                            int64
          1
                           200 non-null
                                            object
              Sex
          2
                           200 non-null
                                            object
          3
              Cholesterol 200 non-null
                                            object
                           200 non-null
                                            float64
          4
              Na_to_K
                                            object
                           200 non-null
              Drug
         dtypes: float64(1), int64(1), object(4)
         memory usage: 9.5+ KB
In [5]:
         df = pd.get_dummies(df, columns = ['Sex', 'BP', 'Cholesterol'],
                      drop_first=True)
In [6]:
         df.head()
```

```
Out[6]:
            Age Na_to_K Drug Sex_M BP_LOW BP_NORMAL Cholesterol_NORMAL
                   25.355 drugY
                                     0
                                             0
                                                         0
                                                                             0
         0
              23
          1
              47
                   13.093 drugC
                                     1
                                             1
                                                         0
                                                                             0
         2
              47
                   10.114 drugC
                                     1
                                             1
                                                         0
                                                                             0
          3
              28
                   7.798 drugX
                                             0
                                                                             0
                                                         1
              61
                   18.043 drugY
                                             1
                                                         0
                                                                             0
 In [7]:
          #Select relevant features
          X = df.drop('Drug', axis=1)
          y = df['Drug']
 In [8]:
          #Split data into training and test sets
          X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=4
 In [9]:
          print(X_train.shape)
          print(y_train.shape)
          print(X_test.shape)
          print(y_test.shape)
          (160, 6)
          (160,)
          (40, 6)
          (40,)
In [10]:
          #Fit the Decision Tree Classifier
          model = DecisionTreeClassifier(random_state=42)
          model.fit(X_train, y_train)
Out[10]: DecisionTreeClassifier(random_state=42)
In [11]:
          #Make predictions
          y_pred = model.predict(X_test)
In [12]:
          #Evaluate the model
          accuracy = accuracy_score(y_test, y_pred)
In [13]:
          print(f"\nAccuracy on the whole dataset: {accuracy}")
         Accuracy on the whole dataset: 1.0
In [14]:
          tree.plot_tree(model)
          plt.show()
```



Random Forest Classification

```
In [15]:
          from sklearn.ensemble import RandomForestClassifier
          from sklearn.datasets import make_moons
In [16]:
          #Fit the Random Forest Classifier
          forest = RandomForestClassifier(n_estimators=100, random_state=0)
          forest.fit(X_train, y_train)
         RandomForestClassifier(random_state=0)
Out[16]:
In [17]:
          #Make predictions
          y_pred = model.predict(X_test)
In [18]:
          #Evaluate the model
          accuracy = accuracy_score(y_test, y_pred)
In [19]:
          print(f"\nAccuracy on the whole dataset: {accuracy}")
```

Accuracy on the whole dataset: 1.0