## DMML Lab Assignment

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## 1 Aim:

To write a program in python to create a decision tree. The program takes a labeled dataset as input and produces a decision tree. Data set is taken in a ratio of 70:30 for training and data.

## 2 Code and Output:

```
[1]: import pandas as pd
     from sklearn import *
     from sklearn.tree import export_graphviz
     from sklearn.metrics import *
[2]: df = pd.read_csv('hcvdat0.csv')
[3]:
     df.shape
[3]: (615, 14)
[4]:
     df.columns
[4]: Index(['Unnamed: 0', 'Category', 'Age', 'Sex', 'ALB', 'ALP', 'ALT', 'AST',
            'BIL', 'CHE', 'CHOL', 'CREA', 'GGT', 'PROT'],
           dtype='object')
     df[df.isnull().any(axis=1)].head()
[5]:
          Unnamed: 0
                           Category
                                      Age Sex
                                                ALB
                                                       ALP
                                                              ALT
                                                                    AST
                                                                          BIL
                                                                                 CHE
     121
                 122
                      0=Blood Donor
                                       43
                                            m
                                               48.6
                                                      45.0
                                                            10.5
                                                                   40.5
                                                                          5.3
                                                                                7.09
                                                            19.1
     319
                 320 0=Blood Donor
                                       32
                                            f
                                               47.4
                                                      52.5
                                                                  17.1
                                                                          4.6
                                                                               10.19
     329
                 330
                      0=Blood Donor
                                       33
                                               42.4
                                                     137.2 14.2 13.1
                                                                          3.4
                                                                                8.23
                                            f
     413
                 414 0=Blood Donor
                                       46
                                            f
                                               42.9
                                                      55.1 15.2
                                                                   29.8
                                                                          3.6
                                                                                8.37
     424
                 425 0=Blood Donor
                                               45.6
                                                    107.2 24.4 39.0
                                                                        13.8
                                                                                9.77
                                       48
```

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```
CHOL
                 CREA
                         GGT
                              PROT
                 63.0
                       25.1
                              70.0
      121
            NaN
      319
                 63.0
                       23.0
                             72.2
            NaN
      329
            NaN
                 48.0
                       25.7 74.4
      413
                 61.0
                       29.0 71.9
            NaN
      424
            NaN
                 88.0 38.0 75.1
 [6]:
      df = df.dropna()
 [7]:
      df.shape
      (589, 14)
 [7]:
      df.drop(df.columns[0],axis=1,inplace=True)
 [9]: df.head()
                                                      AST
 [9]:
              Category
                         Age Sex
                                   ALB
                                         ALP
                                                ALT
                                                            BIL
                                                                    CHE
                                                                         CHOL
                                                                                CREA \
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         0=Blood Donor
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      1 0=Blood Donor
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                               m
      2 0=Blood Donor
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                                                                   8.84
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                          32
                                  46.9
                                        74.7
                               m
      3 0=Blood Donor
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                                                     22.6
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                                                                         4.74
                                                                                80.0
                               m
      4 0=Blood Donor
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                                                     24.8
                                                            9.6
                                                                   9.15
                                                                         4.32
                                                                                76.0
                               m
          GGT
               PROT
         12.1
               69.0
         15.6 76.5
         33.2 79.3
      3 33.8 75.7
      4 29.9 68.7
[10]: from sklearn.preprocessing import LabelEncoder
[11]: category=LabelEncoder()
      sex=LabelEncoder()
      df['category_df'] = category.fit_transform(df['Category'])
      df['sex_df'] = category.fit_transform(df['Sex'])
      df.head()
                                                      AST
[11]:
              Category
                         Age Sex
                                   ALB
                                         ALP
                                                ALT
                                                            BIL
                                                                    CHE
                                                                         CHOL
                                                                                CREA
                                                                               106.0
         0=Blood Donor
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                                  38.5
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         0=Blood Donor
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                                               18.0
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                                                                 11.17
                                                                         4.80
                                                                                74.0
                                  38.5
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      2 0=Blood Donor
                          32
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                                               36.2
                                                     52.6
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                                                                   8.84
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                                                                                86.0
      3 0=Blood Donor
                          32
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                                        52.0
                                               30.6
                                                     22.6
                                                           18.9
                                                                   7.33
                                                                         4.74
                                                                                80.0
                               m
      4 0=Blood Donor
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                                  39.2
                                        74.1
                                               32.6 24.8
                                                            9.6
                                                                   9.15
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                                                                                76.0
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```

```
GGT PROT category_df
                                 sex_df
      0 12.1 69.0
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      2 33.2 79.3
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                                      1
      3 33.8 75.7
                              0
                                      1
      4 29.9 68.7
                              0
                                      1
[12]: data_class=df['Category'].copy()
      data_class=list(dict.fromkeys(data_class))
      print(data_class)
      df.head()
     ['O=Blood Donor', 'Os=suspect Blood Donor', '1=Hepatitis', '2=Fibrosis',
     '3=Cirrhosis']
                                                                CHE CHOL
[12]:
              Category
                       Age Sex
                                 ALB
                                       ALP
                                             ALT
                                                   AST
                                                         BIL
                                                                            CREA \
                                             7.7 22.1
                                                         7.5
                                                               6.93 3.23
                                                                           106.0
      O O=Blood Donor
                        32
                                38.5
                                      52.5
                             m
      1 0=Blood Donor
                        32
                                38.5
                                      70.3 18.0 24.7
                                                         3.9 11.17 4.80
                                                                            74.0
                             m
      2 0=Blood Donor
                                      74.7
                                            36.2 52.6
                                                         6.1
                                                               8.84 5.20
                                                                            86.0
                        32
                             m
                                46.9
      3 0=Blood Donor
                        32
                                43.2
                                      52.0
                                            30.6 22.6 18.9
                                                               7.33 4.74
                                                                            80.0
                             m
      4 0=Blood Donor
                        32
                                39.2 74.1
                                            32.6 24.8
                                                         9.6
                                                               9.15 4.32
                                                                            76.0
                             m
         GGT PROT category_df
                                 sex_df
      0 12.1 69.0
                              0
                                      1
      1 15.6 76.5
                              0
                                      1
      2 33.2 79.3
                              0
                                      1
                              0
      3 33.8 75.7
                                      1
      4 29.9 68.7
                                      1
[13]: from sklearn.model_selection import train_test_split
[14]: df = df.drop(['Category', 'Sex'], axis=1)
      y=df['category_df'].copy()
      X = df.drop(['category_df'], axis=1)
      X_train, X_test, y_train, y_test =train_test_split(X,y,test_size=0.
       \rightarrow3, random_state=500)
[15]: from sklearn.tree import DecisionTreeClassifier
[16]: model = DecisionTreeClassifier(max_leaf_nodes = 15, random_state = 0)
      model.fit(X_train,y_train)
[16]: DecisionTreeClassifier(max_leaf_nodes=15, random_state=0)
[17]: from sklearn.metrics import accuracy_score,
       →confusion_matrix,classification_report
```

```
[18]: y_predicted = model.predict(X_test)
    print("Accuracy : ", accuracy_score(y_test,y_predicted)*100)
    print("Report : ", classification_report(y_test,y_predicted))
    print(confusion_matrix(y_test,y_predicted))
```

Accuracy: 94.35028248587571

Accuracy: 94.35028248587571									
Report :			precision			recall	f1-score	support	
			0		0.99	0.98	0.98	164	
			1		0.00	0.00	0.00	1	
			2		0.40	0.50	0.44	4	
			3		0.50	0.50	0.50	2	
			4		0.57	0.67	0.62	6	
accuracy							0.94	177	
macro avg					0.49	0.53	0.51	177	
weighted avg				0.95	0.94	0.95	177		
[[160	0	1	0	0	3]				
[ :	1	0	0	0	0]				
[ (	0	1	2	1	0]				
[ (	0	0	1	1	0]				
[ (	0	0	2	0	4]]				

## [20]: !dot -Tpng tree.dot -o tree.png

[]:

