

## D.SAMPLE CODING

```
#import the dataset from specified location
data = pd.read_csv('E:/Datascience/Datasets/indian_liver_patient.csv')
```

```
# showing the data from top 5
data.head()
```

	Age	Gender	Total_Bilirubin	Direct_Bilirubin	Alkaline_Phosphotase	Alamine_Aminotransferase	Aspartate_Aminotransferas
0	65	Female	0.7	0.1	187	16	18
1	62	Male	10.9	5.5	699	64	100
2	62	Male	7.3	4.1	490	60	68
3	58	Male	1.0	0.4	182	14	20
4	72	Male	3.9	2.0	195	27	59

```
data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 583 entries, 0 to 582
Data columns (total 11 columns):
 #   Column              Non-Null Count  Dtype  
---  -
 0   Age                 583 non-null   int64  
 1   Gender              583 non-null   object  
 2   Total_Bilirubin     583 non-null   float64 
 3   Direct_Bilirubin    583 non-null   float64 
 4   Alkaline_Phosphotase 583 non-null   int64  
 5   Alamine_Aminotransferase 583 non-null   int64  
 6   Aspartate_Aminotransferase 583 non-null   int64  
 7   Total_Protiens      583 non-null   float64 
 8   Albumin             583 non-null   float64 
 9   Albumin_and_Globulin_Ratio 579 non-null   float64 
10  Dataset             583 non-null   int64  
dtypes: float64(5), int64(5), object(1)
memory usage: 50.2+ KB
```

```
data.isnull().any()
```

```
Age                False
Gender             False
Total_Bilirubin    False
Direct_Bilirubin   False
Alkaline_Phosphotase False
Alamine_Aminotransferase False
Aspartate_Aminotransferase False
Total_Protiens     False
Albumin            False
Albumin_and_Globulin_Ratio True
Dataset            False
dtype: bool
```

```
data.isnull().sum()
```

```
Age                0
Gender             0
Total_Bilirubin    0
Direct_Bilirubin   0
Alkaline_Phosphotase 0
Alamine_Aminotransferase 0
Aspartate_Aminotransferase 0
Total_Protiens     0
Albumin            0
Albumin_and_Globulin_Ratio 4
Dataset            0
dtype: int64
```

```
#checking for the missing data after cleaning data
data['Albumin_and_Globulin_Ratio'] = data.fillna(data['Albumin_and_Globulin_Ratio'].mode()[0])
data.isnull().sum()

Age                0
Gender             0
Total_Bilirubin    0
Direct_Bilirubin   0
Alkaline_Phosphatase 0
Alamine_Aminotransferase 0
Aspartate_Aminotransferase 0
Total_Protiens     0
Albumin            0
Albumin_and_Globulin_Ratio 0
Dataset            0
dtype: int64
```



```
from flask import Flask, render_template, request
import numpy as np
import pickle
```

```
app=Flask(__name__) # our flask app

@app.route('/') # rendering the html template
def home():
    return render_template('home.html')
@app.route('/predict') # rendering the html template
def index() :
    return render_template("index.html")
```

```
@app.route('/data_predict', methods=['POST']) # route for our prediction
def predict():
    age = request.form['age'] # requesting for age data
    gender = request.form['gender'] # requesting for gender data
    tb = request.form['tb'] # requesting for Total_Bilirubin data
    db = request.form['db'] # requesting for Direct_Bilirubin data
    ap = request.form['ap'] # requesting for Alkaline_Phosphatase data
    aa1 = request.form['aa1'] # requesting for Alamine_Aminotransferase data
    aa2 = request.form['aa2'] # requesting for Aspartate_Aminotransferase data
    tp = request.form['tp'] # requesting for Total_Protiens data
    a = request.form['a'] # requesting for Albumin data
    agr = request.form['agr'] # requesting for Albumin_and_Globulin_Ratio data

    # coverting data into float format
    data = [[float(age), float(gender), float(tb), float(db), float(ap), float(aa1), float(aa2), float(tp),
    float(a), float(agr))]

    # Loading model which we saved
    model = pickle.load(open('liver_analysis.pkl', 'rb'))

    prediction= model.predict(data)[0]
    if (prediction == 1):
        return render_template('noChance.html', prediction='You have a liver disease problem, You must and :
    else:
        return render_template('chance.html', prediction='You dont have a liver disease problem')

if __name__ == '__main__':
    app.run()
```

```
base) D:\TheSmartBridge\Projects\2. DrugClassification\Drug c
* Serving Flask app "app" (lazy loading)
* Environment: production
  WARNING: This is a development server. Do not use it in a p
  Use a production WSGI server instead.
* Debug mode: off
* Running on http://127.0.0.1:5000/ (Press CTRL+C to quit)
```

## Liver Patient Prediction

You have a liver disease problem, You must and should consult a doctor. Take care