Basic Details:

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Unified Mentor ID: UIMD10042529348 **Project**: Machine Learning Project

Project Information:

Title: Liver Cirrhosis Stage Detection

Objective

Build a system that can output the level of liver damage (liver cirrhosis) given patient data.

Dataset

The data provided is sourced from a Mayo Clinic study on primary biliary cirrhosis (PBC) of the liver carried out from 1974 to 1984.

Description of columns:

- N_Days: Number of days between registration and the earlier of death, transplantation, or study analysis time in 1986
- Status: status of the patient C (censored), CL (censored due to liver tx), or D (death)
- Drug: type of drug D-penicillamine or placebo
- Age: age in days
- Sex: M (male) or F (female)
- Ascites: presence of ascites N (No) or Y (Yes)
- Hepatomegaly: presence of hepatomegaly N (No) or Y (Yes)
- Spiders: presence of spiders N (No) or Y (Yes)
- Edema: presence of edema N (no edema and no diuretic therapy for edema), S (edema present without diuretics, or edema resolved by diuretics), or Y (edema despite diuretic therapy)
- Bilirubin: serum bilirubin in [mg/dl]
- Cholesterol: serum cholesterol in [mg/dl]
- Albumin: albumin in [gm/dl]
- Copper: urine copper in [ug/day]
- Alk Phos: alkaline phosphatase in [U/liter]
- SGOT: SGOT in [U/ml] (a liver enzyme)
- Tryglicerides: triglicerides in [mg/dl]
- Platelets: platelets per cubic [ml/1000]
- Prothrombin: prothrombin time in seconds [s]
- Stage: histologic stage of disease (1, 2, or 3)

Project Link:

https://github.com/Alforeverything/UnifiedMentorInternshipProjects/blob/c86c2928100b9b567ee2 361675a7f402cc307a20/Liver%20Cirrhosis%20Stage%20Detection/liver_cirrhosis_stage.ipynb

https://github.com/AIforeverything/UnifiedMentorInternshipProjects/blob/c86c2928100b9b567ee2361675a7f402cc307a20/categorical/categorical model.py

Code

Steps Followed:

Step-1: Initially I have created a library for building a categorical machine learning model and used this library for building model.

categorical model.py

```
### Step-1: Common virtual environment was created and activated: myenv
# ## pip install virtualenv
### virtualenv myenv
###.\myenv\Scripts\activate.ps1
def greet(name):
  return f"good job {name}"
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import os
import sys
from pathlib import Path
import zipfile
import warnings
warnings.filterwarnings("ignore")
import sklearn
from sklearn.preprocessing import LabelEncoder, OneHotEncoder
from sklearn.preprocessing import MinMaxScaler
from sklearn.model selection import train test split
from sklearn.linear model import LogisticRegressionCV
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier, GradientBoostingClassifier
from sklearn.naive bayes import GaussianNB
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
from xgboost import XGBClassifier
from sklearn.model selection import GridSearchCV
from sklearn.metrics import accuracy score, classification report, confusion matrix
import joblib
# import tensorflow as tf
# from tensorflow import keras
# from tensorflow.keras.models import Sequential
# from tensorflow.keras.layers import Dense
class categorical Model:
```

```
def \_\_init\_\_(self,model, target\_column, df):
  Initializes the categorical Target class.
  Parameters:
  model (str): The name of the model to be used.
  target column (str): The name of the target column.
  df (pd.DataFrame): The DataFrame containing the data.
  self.df = df
  self.target column = target column
  self.model = model
# Importing data into a dataframe from csv file in the directory
def readingData():
 #checking the directory for .csv files
  directory = Path('./')
  # List all CSV files
  for csv file in directory.glob('*.csv'):
     print(csv file.name)
  df= pd.read csv(csv file)
  return df
## Data extraction from zipfile
def extractingZipFile(zipFilePath, extractTo):
  Extracts the contents of a zip file to a specified directory.
  Parameters:
  zipFilePath (str): The path to the zip file.
  extractTo (str): The directory to extract the contents to.
  with zipfile.ZipFile(zipFilePath, 'r') as zip ref:
     zip ref.extractall(extractTo)
# EDA (Exploratory Data Analysis)
# Checking missing values
def checkMissingValues(df):
  Checks for missing values in the DataFrame
  Parameters:
  df (pd.DataFrame): The DataFrame to check for missing values.
  Returns:
```

```
missing values
    return df.isnull().sum()
  # Removing duplicates
  ## function to check for duplicates and remove dupliates
  def checkDuplicates(df):
    Checks for duplicate rows in the DataFrame and removes them.
    Parameters:
    df (pd.DataFrame): The DataFrame to check for duplicates.
    Returns:
    pd.DataFrame: The DataFrame with duplicates removed.
    duplicates = df.duplicated().sum()
    if duplicates > 0:
       df = df.drop duplicates()
       print(f"Removed {duplicates} duplicate rows.")
    else:
       print("No duplicate rows found.")
    return df
  #Function for all columns
  def allColumns(df):
    return list(df.columns)
  # Function for categorical columns
  def catColumns(df):
    catCol=df.select dtypes(include='object').columns
    return catCol
  # Function for Non-categorical columns
  def nonCatColumns(df):
    numeric col=df.select dtypes(include='number').columns
    return numeric col
  ## function to check categorical columns and replacing them with numerical values
  def checkCategoricalColumnsAndReplacingWithLE(df):
    Checks for categorical columns in the DataFrame and replaces them with numerical
values.
```

df (pd.DataFrame): The DataFrame to check for categorical columns.

Parameters:

```
Returns:
    pd.DataFrame: The DataFrame with categorical columns replaced with numerical
values.
    categorical columns = df.select dtypes(include=['object']).columns
    print(f"Categorical columns: {categorical columns}")
    for col in categorical columns:
       print(f"col.unique(): {df[col].unique()}")
       print(f''col.value counts(): {df[col].value counts()}")
       le = LabelEncoder()
       df[col] = le.fit transform(df[col])
    return df
  # function to standardize Non Categorical columns
  def standardizeNonCategoricalColumns(df):
    minMax=MinMaxScaler()
    numeric col=df.select dtypes(include='number').columns
    df[numeric col]=minMax.fit transform(df[numeric col])
    return df
  ## function to removing the missing values
  def removeMissingValues(df):
    Removes rows with missing values from the DataFrame.
    Parameters:
    df (pd.DataFrame): The DataFrame to remove missing values from.
    Returns:
    pd.DataFrame: The DataFrame with missing values removed.
    df = df.dropna()
    return df
  #function to print the correlation matrix respect to the target column
  def printCorrelationMatrix(df, target column):
    Prints the correlation matrix of the DataFrame with respect to the target column.
    Parameters:
    df (pd.DataFrame): The DataFrame to print the correlation matrix for.
    target column (str): The name of the target column.
    Returns:
```

```
pd.DataFrame: The correlation matrix.
  # print the correlation matrix with respect to the target column
  print(f"Correlation matrix with respect to {target column}:")
  print(df.corr()[target column].sort values(ascending=False))
  corr text=df.corr()[target column].sort values(ascending=False)
  # .to string() provides a nicely formatted text version of the DataFrame.
  # This will produce a human-readable file.
  # If we want a machine-readable format instead, consider .to csv("file.txt", sep='\t').
  with open('correlation.txt', 'w') as f:
     f.write(corr text.to string())
  corr = df.corr()
  plt.figure(figsize=(12, 8))
  sns.heatmap(corr, annot=True, fmt=".2f", cmap='coolwarm')
  plt.title(f"Correlation Matrix with respect to {target column}")
  plt.show()
  return corr
#checking missing values of each column
def missing columns(df):
  return (df.isnull().sum())
#checking missing values of all columns
def missing columns total(df):
  return (df.isnull().sum().sum())
## function to split the data into X,y
def splitDataIntoXy(df, target column):
  Splits the DataFrame into X and y.
  retuns tuple
  X = df.drop(target column, axis=1)
  y = df[target column]
  return X,v
## function to split the data into train and test
def splitData(X,y):
  Splits the DataFrame into training and testing sets.
  Parameters:
  X,y
  Returns:
  tuple: The training and testing sets.
  X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
```

```
return X train, X test, y train, y test
```

function to train the model and compare the models and save the best model and the model report and the model performance

```
def trainModel(X_train, X_test, y_train, y_test):
```

Trains the model and compares the models and saves the best model and the model report and the model performance.

```
Parameters:
X train (pd.DataFrame): The training data.
X test (pd.DataFrame): The testing data.
y train (pd.Series): The training labels.
y test (pd.Series): The testing labels.
Returns:
None
** ** **
models = {
  "Logistic Regression": LogisticRegressionCV(max iter=10000),
  "Decision Tree": DecisionTreeClassifier(),
  "RandomForest": RandomForestClassifier(min samples split=5),
  "Gradient Boosting": GradientBoostingClassifier(),
  "Naive Bayes" : GaussianNB(),
  "KNN": KNeighborsClassifier(),
  "Support Vector Machines": SVC(),
  "XGBoost": XGBClassifier()
}
best model = None
best accuracy = 0
for name, model in models.items():
  model.fit(X train, y train)
  y pred = model.predict(X test)
  accuracy = accuracy score(y test, y pred)
  print(f"{name} Accuracy: {accuracy:.4f}")
  if accuracy > best accuracy:
    best accuracy = accuracy
    best model = model
    best model name = name
```

```
print(f"Best Model: {best_model.__class__._name__}) with accuracy:
{best accuracy:.2f}")
    # Save the best model
    joblib.dump(best model name, f'{best model name}.pkl')
    # Save the classification report
    report = classification report(y_test, y_pred)
    with open('classification report.txt', 'w') as f:
       f.write(f"Model: {best model name} \n\n")
       f.write(report)
    # Save the confusion matrix
    cm = confusion matrix(y test, y pred)
    np.savetxt('confusion matrix.txt', cm, fmt='%d')
  # function to load the model
  def loadModel(model path):
    Loads the model from the specified path.
    Parameters:
    model path (str): The path to the model.
    Returns:
    model: The loaded model.
    model = joblib.load(model path)
    return model
# making an object of the class to use the functions
def main():
  # Unzip the file
  file= categorical Model.extractingZipFile('./', "./")
  # Reading the data
  df = categorical Model.readingData()
  # Checking for missing values
  missing values = categorical Model.checkMissingValues(df)
  print(f"Missing values: {missing values}")
  # Checking for duplicates
  df = categorical Model.checkDuplicates(df)
  # Checking for categorical columns
```

```
df = categorical Model.checkCategoricalColumns(df)
  # Removing missing values
  df = categorical Model.removeMissingValues(df)
  # Choosing the target column
  target_column = input("Enter the target column name: ")
  if target column not in df.columns:
     print(f'Target column '{target column}' not found in DataFrame.")
  else:
     print(f"Target column '{target column}' found in DataFrame.")
  # Printing the correlation matrix
  corr matrix = categorical Model.printCorrelationMatrix(df, target column)
  # Splitting the data into train and test sets
  X train, X test, y train, y test = categorical Model.splitData(df, target column)
  # Training the model and saving the best model
  categorical Model.trainModel(X train, X test, y train, y test)
                               Step-2: Code for model:
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.preprocessing import MinMaxScaler
# In[2]:
# making a path to get the modules from categorical library
import os
os.chdir('..')
# In[3]:
from categorical.categorical_model import categorical_Model
# In[4]:
```

```
categorical_Model.extractingZipFile("./Liver Cirrhosis Stage
Detection/liver_cirrhosis_stage.zip",'./Liver Cirrhosis Stage Detection/')
# In[5]:
df=pd.read_csv("./Liver Cirrhosis Stage Detection/liver_cirrhosis_stage/liver_cirrhosis.csv")
df.head()
# ## EDA
# In[6]:
df.info()
#### Removing duplicates
# In[7]:
categorical_Model.checkDuplicates(df)
# ### Checking and Removing missing values
# In[8]:
# Checking missing values for each column
print(categorical_Model.missing_columns(df))
#checking missing values of all columns
print(categorical_Model.missing_columns_total(df))
df.dropna(inplace=True)
df.head()
# In[9]:
def showingUnique(x):
  return x.unique()
# In[10]:
c=list(df.columns)
```

```
for i in c:
  if df.dtypes[i]=='object':
    print(i,showingUnique(df[i]))
\# ### Splitting into X and y before standardize X
# In[11]:
X,y=categorical_Model.splitDataIntoXy(df,"Stage")
# ### Standardizing the numerical columns of X
# In[12]:
categorical\_Model.standardizeNonCategoricalColumns(X)
# ### converting categorical columns to numerical of X
# In[13]:
categorical\_Model. check Categorical Columns And Replacing With LE(X)
# In[14]:
X.info()
# In[15]:
y.info()
# In[16]:
y.unique()
# In[17]:
X_train, X_test, y_train, y_test=categorical_Model.splitData(X,y)
```

In[18]:

categorical_Model.trainModel(X_train, X_test, y_train, y_test)

Model Outcomes

Different models are built using the dataset and found

Logistic Regression Accuracy: 0.6012

Decision Tree Accuracy: 0.9186

KNN Accuracy: 0.8832

Best Model: DecisionTreeClassifier with accuracy: 0.92

classification report:

Model: Decision Tree

p	recision	recall	f1-sco	e sup	port
1	0.89	0.88	0.88	165	7
2	0.85	0.87	0.86	169	7
3	0.91	0.89	0.90	164	6
accurac	_y		0.88	5000)
macro a	vg = 0.8	88 0	.88 0	.88	5000
weighted avg (.88	0.88	0.88	5000