

# Revolution Liver care: Predicting Liver Cirrhosis using Advanced Machine Learning techniques

## Introduction:

Liver cirrhosis is a progressive and irreversible liver disease often resulting from chronic liver conditions such as hepatitis, alcohol abuse, and fatty liver disease. Early detection is crucial for effective treatment and patient survival. Traditional diagnostic methods are often invasive, expensive, or too late in detecting the onset of cirrhosis.

In this project, we harness the power of **machine learning** to build a predictive model capable of identifying potential liver cirrhosis cases using clinical and biochemical data. This approach aims to aid medical professionals in early diagnosis, minimize the need for invasive tests, and ultimately improve patient outcomes.

## Description:

Objective: To develop a machine learning model that can accurately predict liver cirrhosis from patient medical data.

Dataset Used:

The Indian Liver Patient Dataset (ILPD) from the UCI Machine Learning Repository.

Attributes include:

- Age
- Gender
- Total Bilirubin
- Direct Bilirubin
- Alkaline Phosphatase
- Alamine Aminotransferase (ALT)
- Aspartate Aminotransferase (AST)
- Total Proteins
- Albumin

- Albumin and Globulin Ratio
- Class (1 for patients with liver disease, 2 for no disease)

📋 Steps Involved:

1. Data Preprocessing & Cleaning
2. Feature Engineering
3. Splitting dataset
4. Model training using algorithms like Random Forest, XGBoost, or SVM
5. Model Evaluation (Accuracy, Precision, Recall, F1-Score, AUC)
6. Visualization of results
7. Deployment (optional)

## Code:

```
# Upload the CSV file to Colab from local machine
```

```
from google.colab import files
```

```
uploaded = files.upload()
```

```
# Data manipulation
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Data visualization
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
# For better table display
```

```
from IPython.display import display
```

```
# Set Seaborn style
```

```
sns.set(style="whitegrid")
```

```
# Make sure plots display inline
```

```
%matplotlib inline
```

```
# Read the uploaded dataset
```

```
df = pd.read_excel("HealthCareData.xlsx")
```

```
# Display first 5 rows
```

```
display(df.head())
```

```
# Print info about the dataset
```

```
print("\nDataset Info:\n")
```

```
df.info()
```

```
# Print shape
```

```
print("\nDataset Shape:", df.shape)
```

<

```
# Check the shape of the dataset
```

```
print("Dataset Shape:", df.shape)
```

```
# Check data types and non-null counts

print("\nDataset Info:\n")

df.info()


# Check for missing values

print("\nMissing Values per Column:\n")

print(df.isnull().sum())

# List numerical columns (excluding the target if needed)

numerical_cols = df.select_dtypes(include=['int64', 'float64']).columns.tolist()

import matplotlib.pyplot as plt

import seaborn as sns


plt.figure(figsize=(15, 10))

for i, col in enumerate(numerical_cols):

    plt.subplot((len(numerical_cols)+2)//3, 3, i+1)

    sns.boxplot(x=df[col], color='lightblue')

    plt.title(col)

    plt.tight_layout()

plt.show()

from google.colab import drive

drive.mount('/content/drive')
```

```
print("\nMissing Values per Column:\n")
print(df.isnull().sum())
```

Dataset Shape: (950, 42)

Dataset Info:

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 950 entries, 0 to 949
Data columns (total 42 columns):
```

#	Column	Non-Null Count	Dtype
0	S.NO	950 non-null	int64
1	Age	950 non-null	int64
2	Gender	950 non-null	object
3	Place(location where the patient lives)	816 non-null	object
4	Duration of alcohol consumption(years)	950 non-null	int64
5	Quantity of alcohol consumption (quarters/day)	950 non-null	int64
6	Type of alcohol consumed	950 non-null	object
7	Hepatitis B infection	950 non-null	object
8	Hepatitis C infection	950 non-null	object
9	Diabetes Result	950 non-null	object
10	Blood pressure (mmhg)	950 non-null	object
11	Obesity	950 non-null	object
12	Family history of cirrhosis/ hereditary	950 non-null	object
13	TCH	591 non-null	float64
14	TG	591 non-null	object
15	LDL	591 non-null	object
16	HDL	582 non-null	float64
17	Hemoglobin (g/dl)	950 non-null	float64
18	PCV (%)	920 non-null	float64
19	RBC (million cells/microliter)	398 non-null	float64
20	MCV (femtoliters/cell)	641 non-null	float64

## Conclusion:

This project demonstrates the potential of machine learning in revolutionizing liver care. By leveraging clinical data, healthcare systems can detect liver cirrhosis early, improve decision-making, and enhance patient outcomes. This initiative shows how data science can save lives—transforming raw data into actionable medical intelligence.