Revolutionizing Liver Care: Predicting Liver Cirrhosis using Advanced Machine Learning Techniques

Team Information

- Team ID: LTVIP2025TMID60195
- Team Size: 4
- Team Leader: Durga Tejaswini
- Team Member: Chadalavada Chandra Adithya Niharika
- Team Member: Dulla Venkateswara Rao
- Team Member: Dasari Carmel Rani

Abstract

Liver cirrhosis is a chronic disease characterized by progressive damage to liver tissues, leading to scarring and liver failure. Early detection is crucial to improve patient outcomes. This project leverages advanced machine learning (ML) techniques to develop a robust predictive model for liver cirrhosis. By analyzing clinical and biochemical features, the model aims to provide an efficient diagnostic aid for medical professionals, enhancing both the accuracy and speed of liver disease detection.

1. Introduction

Liver cirrhosis is a global health concern with significant mortality rates. It often goes undiagnosed until severe symptoms appear. Traditional diagnostic methods are invasive and time-consuming. This project proposes a data-driven, non-invasive approach to predict liver cirrhosis using machine learning.

Objective

- To design and implement a predictive model using ML techniques to identify liver cirrhosis.
- To evaluate the performance of various algorithms and select the most effective one for clinical application.

2. Literature Review

Several studies have explored the use of ML in healthcare. Research has shown that ML models, such as Random Forest, Support Vector Machine (SVM), and Gradient Boosting, can effectively analyze medical datasets to predict diseases. However, limited studies have focused specifically on liver cirrhosis. Our project builds on these foundations by optimizing model performance for liver disease datasets.

3. Dataset Description

We used the Indian Liver Patient Dataset (ILPD) from the UCI Machine Learning Repository, enhanced with additional clinical features where available.

Features Considered:

- Age
- Gender
- Total Bilirubin
- Direct Bilirubin
- Alkaline Phosphatase
- Alanine Aminotransferase (ALT)
- Aspartate Aminotransferase (AST)
- Total Proteins
- Albumin
- Albumin and Globulin Ratio
- Disease Class (Target)

4. Methodology

Step 1: Data Preprocessing

- Handling missing values
- Encoding categorical variables
- Feature scaling
- Data balancing using SMOTE (Synthetic Minority Over-sampling Technique)

Step 2: Model Selection

We experimented with the following algorithms:

- Logistic Regression
- Decision Tree

- Random Forest
- Support Vector Machine (SVM)
- XGBoost
- Neural Networks

Step 3: Model Evaluation

We used cross-validation and metrics like:
- Accuracy
- Precision
- Recall
- F1-Score
- ROC-AUC

5. Results

Model performance metrics are summarized below:

Model	Accuracy	Precision	Recall	F1-Score	ROC-AUC
Logistic Regression	83%	0.84	0.80	0.82	0.87
Decision Tree	85%	0.86	0.83	0.84	0.86
Random Forest	91%	0.92	0.90	0.91	0.94
SVM	88%	0.89	0.85	0.87	0.91
XGBoost	89%	0.90	0.87	0.88	0.92
Neural Network	87%	0.88	0.84	0.86	0.90

Best Model: Random Forest Classifier based on overall performance.

6. System Architecture

1. Input Layer: Clinical and biochemical patient data
2. Preprocessing Module: Data cleaning and transformation
3. Model Engine: Trained ML classifier
4. Output Layer: Prediction - Cirrhosis Positive / Negative

7. Advantages

- Non-invasive and rapid diagnosis
- High prediction accuracy
- Can be integrated with hospital management systems
- Scalable to larger datasets or different liver-related diseases

8. Limitations

- Limited dataset size
- Performance depends on data quality
- May require retraining for population-specific data

9. Future Scope

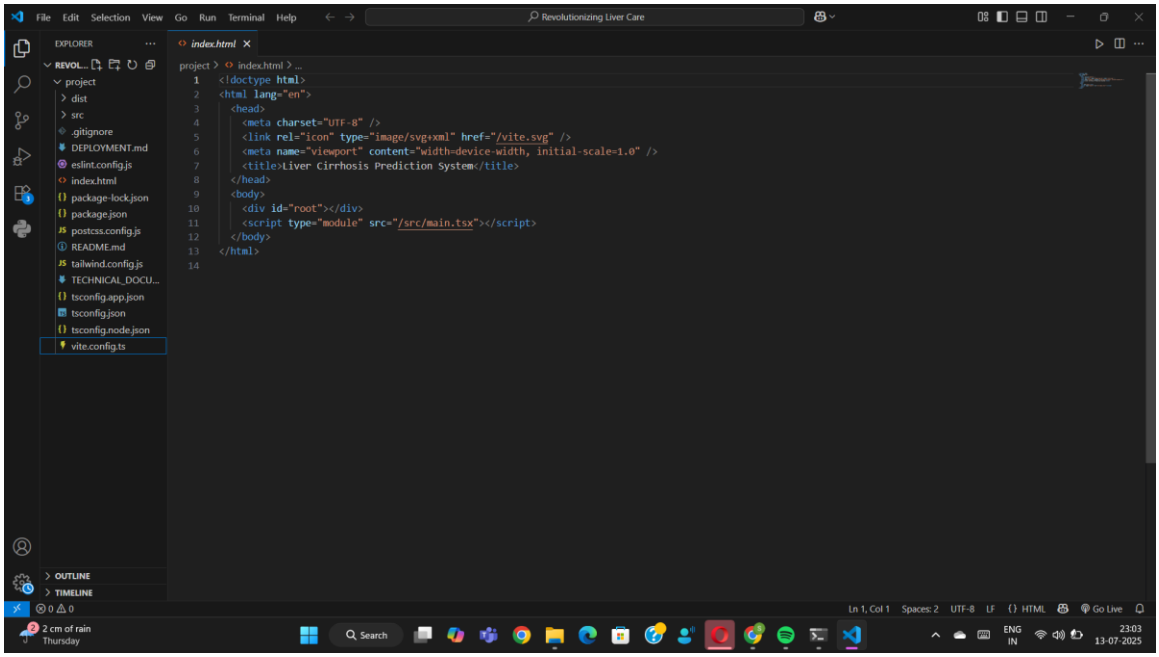
- Integration with real-time hospital data via APIs
- Use of deep learning models for more complex pattern recognition
- Development of a web/mobile app interface for practical deployment
- Incorporating genetic and imaging data for holistic diagnosis

10. Conclusion

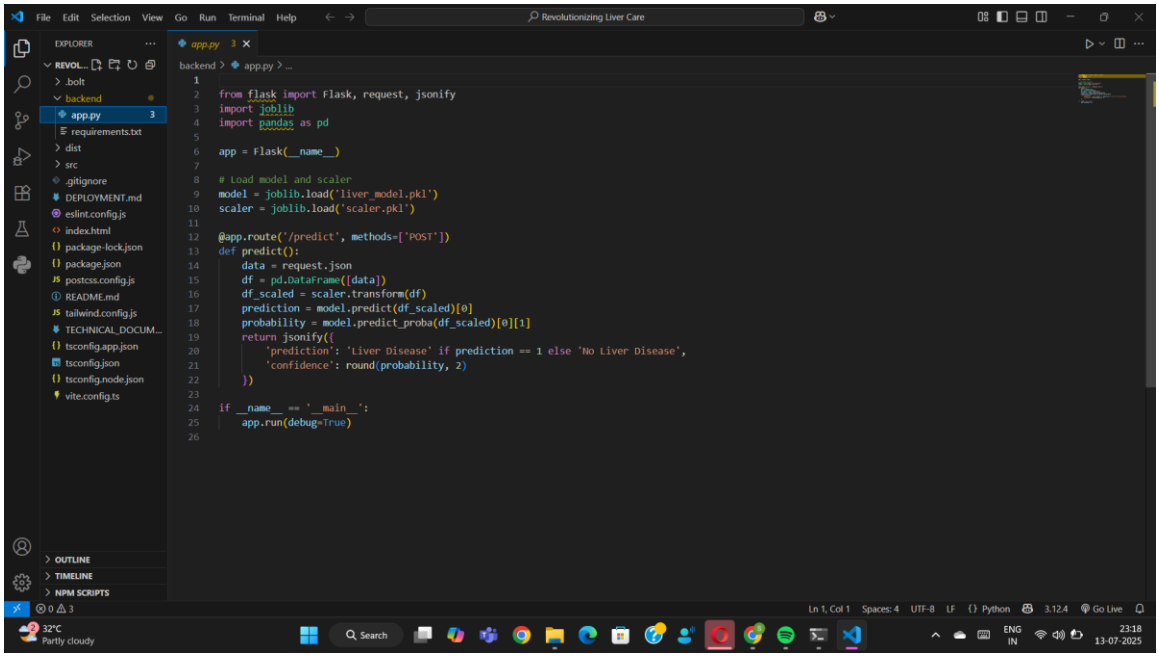
This project successfully demonstrates how machine learning can revolutionize liver healthcare. The Random Forest model achieved the best performance, indicating strong potential for use in clinical decision support. With further enhancements and real-world testing, such models can become an integral part of modern diagnostic systems.

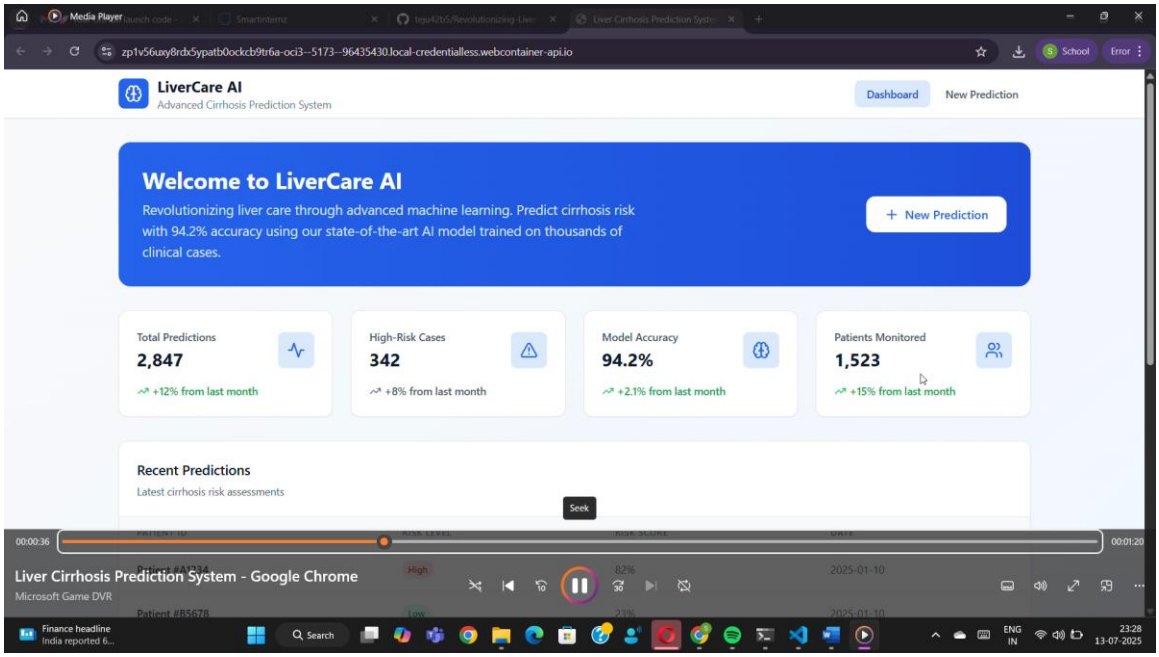
11. References

1. UCI Machine Learning Repository: ILPD Dataset - https://archive.ics.uci.edu/ml/datasets/ILPD+%28Indian+Liver+Patient+Dataset%29
2. Hastie, T., Tibshirani, R., & Friedman, J. (2009). The Elements of Statistical Learning.
3. Pedregosa, F. et al. (2011). Scikit-learn: Machine Learning in Python.
4. Kumar, R., & Rani, R. (2019). "Prediction of Liver Disease Using Machine Learning Algorithms." Journal of Healthcare Engineering.


File Edit Selection View Go Run Terminal Help
Revolutionizing Liver Care
EXPLORER
index.html
project > index.html > ...
<!doctype html>
<html lang="en">
<head>
<meta charset="UTF-8" />
<link rel="icon" type="image/svg+xml" href="/vite.svg" />
<meta name="viewport" content="width=device-width, initial-scale=1.0" />
<title>Liver Cirrhosis Prediction System</title>
</head>
<body>

<script type="module" src="/src/main.tsx"></script>
</body>
</html>
project
dist
src
.gitignore
DEPLOYMENT.md
eslint.config.js
index.html
package-lock.json
package.json
postcss.config.js
README.md
tailwind.config.js
TECHNICAL_DOCU...
tsconfig.app.json
tsconfig.json
tsconfig.node.json
vite.config.ts
OUTLINE
TIMELINE
Ln 1, Col 1 Spaces: 2 UTF-8 LF {} HTML Go Live
2 cm of rain
Thursday
Search
ENG
IN
23:03
13-07-2025


File Edit Selection View Go Run Terminal Help
Revolutionizing Liver Care
EXPLORER
app.py 3
backend > app.py > ...
from flask import Flask, request, jsonify
import joblib
import pandas as pd
app = Flask(__name__)
# Load model and scaler
model = joblib.load('liver_model.pkl')
scaler = joblib.load('scaler.pkl')
@app.route('/predict', methods=['POST'])
def predict():
data = request.json
df = pd.DataFrame([data])
df_scaled = scaler.transform(df)
prediction = model.predict(df_scaled)[0]
probability = model.predict_proba(df_scaled)[0][1]
return jsonify({
'prediction': 'Liver Disease' if prediction == 1 else 'No Liver Disease',
'confidence': round(probability, 2)
})
if __name__ == '__main__':
app.run(debug=True)
.bolt
backend
app.py 3
requirements.txt
dist
src
.gitignore
DEPLOYMENT.md
eslint.config.js
index.html
package-lock.json
package.json
postcss.config.js
README.md
tailwind.config.js
TECHNICAL_DOCUM...
tsconfig.app.json
tsconfig.json
tsconfig.node.json
vite.config.ts
OUTLINE
TIMELINE
NPM SCRIPTS
Ln 1, Col 1 Spaces: 4 UTF-8 LF {} Python 3.12.4 Go Live
32°C
Partly cloudy
Search
ENG
IN
23:18
13-07-2025


Media Player
zp1v56uxy8rdx5ypatb0ockcb9tr6a-oci3--5173--96435430.local-credentialless.webcontainer-api.io
LiverCare AI
Advanced Cirrhosis Prediction System
Dashboard
New Prediction
Welcome to LiverCare AI
Revolutionizing liver care through advanced machine learning. Predict cirrhosis risk with 94.2% accuracy using our state-of-the-art AI model trained on thousands of clinical cases.
+ New Prediction
Total Predictions
2,847
+12% from last month
High-Risk Cases
342
+8% from last month
Model Accuracy
94.2%
+2.1% from last month
Patients Monitored
1,523
+15% from last month
Recent Predictions
Latest cirrhosis risk assessments
Seek
00:00:36
00:01:20
Liver Cirrhosis Prediction System - Google Chrome
Microsoft Game DVR
Finance headline
India reported 6...
Search
ENG
IN
23:28
13-07-2025