

Heart Disease Prediction using Machine Learning

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Abstract - With the rise in cardiovascular diseases (CVDs), machine learning (ML) has become a key tool in early diagnosis using clinical data. Studies utilize algorithms like SVM, Random Forest, ANN, Decision Trees, and ensemble models, stressing the importance of data preprocessing and feature selection (e.g., PCA, RFE) for improved accuracy. Hybrid and ensemble techniques enhance predictive performance, while model interpretability remains vital for clinical use. Overall, ML models, when properly trained and tuned, significantly aid in the early detection of heart disease and support timely medical intervention.

Key Words: Heart Disease, Machine Learning, Feature Selection, Classification Algorithms, Data Preprocessing, SMOTE, Explainable AI (SHAP), Accuracy Metrics (e.g., AUC, Precision, Recall)

1. INTRODUCTION

Heart disease is the leading global cause of death, influenced by lifestyle, genetics, and chronic issues. While traditional tools like ECGs are useful, they can be costly and less accessible. Machine learning (ML) offers a better alternative by analyzing complex clinical data for improved prediction. Models like Random Forest, SVM, XGBoost, and Neural Networks, along with feature selection methods, boost accuracy. Explainable AI, ensemble models, and mobile apps enable real-time, interpretable support, making diagnosis faster and more affordable.

2. LITERATURE SURVEY

The reviewed studies highlight the growing use of machine learning (ML) in heart disease prediction using various techniques and datasets. Bhatt et al. achieved strong accuracy with real-world data but lacked clinical depth. Biswas et al. compared

feature selection methods effectively, though limited by a small sample. Hajiababi's meta-analysis emphasized methodology and diversity, while Ahire et al. offered a basic overview with minimal rigor. Common findings include the effectiveness of ensemble models, importance of feature selection, and challenges in generalizability. However, most studies lack clinical integration and interpretability. Future work must focus on validation, collaboration, and explainable AI for real-world impact.

1. Machine Learning for Heart Disease Prediction

This body of work highlights the increasing use of machine learning (ML) models in predicting heart disease using clinical datasets. Techniques such as Random Forest, Support Vector Machines (SVM), and ensemble models were commonly employed. Bhatt et al. used a large real-world dataset and advanced preprocessing to achieve high accuracy, although the study lacked clinical interpretability. Biswas et al. provided a comparative analysis of feature selection techniques, showing that Random Forest performed well despite a limited sample size. While these models demonstrated strong technical results, most lacked clinical integration or real-world validation, reducing their practical impact.

2. Meta-Analytical and Introductory Studies in Heart Disease Prediction

Hajiababi conducted a detailed meta-analysis focusing on methodology and dataset diversity, making it a strong academic reference. In contrast, Ahire et al. offered a basic, introductory overview of ML models for heart

disease, with minimal experimental evidence. Together, these studies emphasize the need for rigorous methodological approaches and deeper experimentation to support clinical applications. Despite the promising use of ML, the gap between research and clinical deployment remains a significant barrier.

3. Common Themes and Future Directions

Across the reviewed literature, common themes emerged: the effectiveness of ensemble models, the critical role of feature selection, and the ongoing challenge of generalizability. However, few studies employed interpretability tools like SHAP or addressed real-world clinical integration. The lack of explainability and domain collaboration hinders trust and adoption in healthcare settings. Future research must focus on validation, explainable AI, and integration with clinical workflows to translate ML advancements into practical, patient-centered solutions

Table -1: Summary of Heart disease dataset

Dataset Name	Framingham Heart Disease Dataset		
Class	Two (1-Absent; 2-Presence)		
Instances	4012 (Absent-2945; Presence-1067)		
Attribute	10		
	Minimum Value	Maximum Value	Missing Values
Age	32	70	-
Gender (0-Male; 1-Female)	0	1	-
Blood Pressure (Systolic)	83.5	295	-
Blood Pressure (Diastolic)	48	142.5	-
Heart Rate	44	143	✓
Blood Sugar	40	394	✓
BMI	15.54	56.8	✓
Cholesterol	107	696	✓
Smoking (0-No; 1-Yes)	0	1	-

Table -2: Comparison of Classification techniques

Author	Classification Technique	Test Technique Used	Model	Dataset	Accuracy
Vasanthakumari et al. [11]	ANN	Yes	WEKA	A dataset research institute in Chennai	98.408%
Mohammed et al. [12]	ANN Ensemble	Yes	Not mentioned	Cleveland (UCI)	86.90%
Dai et al. [13]	ANN Ensemble	Yes	SAS ensemble score v2	Cleveland (UCI)	86.81%
Chen et al. [14]	ANN (SG)	Yes	C and C#	Cleveland (UCI)	80%
Shang et al. [15]	ANN	Yes	WEKA	Cleveland and Statlog (UCI)	Nearly 100%
Subramanian and Sugumaran [16]	DT	100-fold feature selection	Not mentioned	A dataset with 300 records for testing and 120 for training	81%
Patel et al. [17]	ANN	Yes	WEKA	Cleveland (UCI)	86.70%
Sharma et al. [18]	ANN	Yes	Not mentioned	Cleveland (UCI)	97.4%
Wahedi et al. [19]	SVM	DTSSVM	Not mentioned	Cleveland (UCI)	81.80%
Chen et al. [20]	ML, KNN, DT and bagging technique	KNN	WEKA	Cleveland (UCI)	76.20%
Prasanna et al. [21]	ML, DT, ML, KNN, SVM, bagging, boosting and stacking	Boosting with SVM	Not mentioned	Cleveland (UCI)	84.81%
Arora et al. [22]	ANN and Genetic Algorithm hybrid system	Yes	MOGLAB	American Heart Association dataset	80%
Wagholikar and Patel [23]	ANN and Genetic Algorithm hybrid system	Yes	MOGLAB	American Heart Association dataset	80%
Arora [24]	ML and DT	Yes	Not mentioned	Cleveland (UCI)	94.37%
Wahedi and Wahedi [25]	ML and DT	Yes	WEKA	UCI	87.81%
Palanques and Wong [26]	DT, ML and ANN	Yes	EMK	Cleveland (UCI)	86.31%
Shang et al. [27]	ANN	Yes	WEKA	Cleveland and Statlog (UCI)	Nearly 100%
Zhang et al. [28]	ML, DT, Decision Trees, Random Forest, and SVM	DT	MOGLAB	Cleveland and Statlog (UCI)	99.03% for Cleveland and 98.19% for Statlog
Li et al. [29]	Ensemble and Weight Set (WFS) for feature selection, Ensemble using C4.5 for classification	Yes	MOGLAB	Statlog (UCI)	92.50%
Ghosh et al. [30]	SVM and Radial Basis Function	SVM	Not mentioned	Indian patients dataset of 114 records and 19 attributes	86.42%
Mahesh and Mahesh [31]	ML, SVM, RPART, Ensemble, Cart, and Boost	Simple Cart	WEKA	South African dataset containing 11 attributes	99.0311%

3. OBJECTIVES

- **Evaluate ML Models for Heart Disease Prediction**

Assess different machine learning models like Random Forest, SVM, and Neural Networks based on accuracy, precision, recall, and AUC. This helps determine which model performs best across diverse datasets for reliable heart disease prediction.

- **Compare Feature Selection Techniques**

Use methods like PCA, RFE, ANOVA, and Chi-Square to identify the most relevant features. Effective feature selection reduces complexity and improves model performance and training efficiency.

- **Analyze Strengths and Limitations of ML Approaches**

Evaluate how well models generalize beyond training data, especially with limited or imbalanced datasets. Understand their practical limitations, such as lack of clinical context or overfitting in small samples.

- **Highlight Model Interpretability and Explainability**

Emphasize tools like SHAP and LIME to explain predictions in a way clinicians can understand. Interpretability builds trust and helps integrate AI into clinical decision-making.

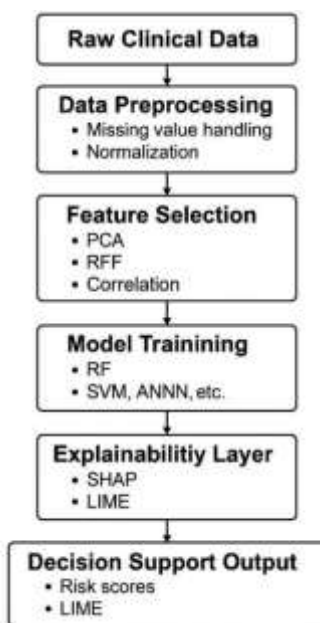
Identify Gaps in Clinical Integration

Most ML models are not validated in real clinical settings or connected to hospital systems like EHRs. Bridging this gap is crucial for real-world usability and adoption in healthcare workflows.

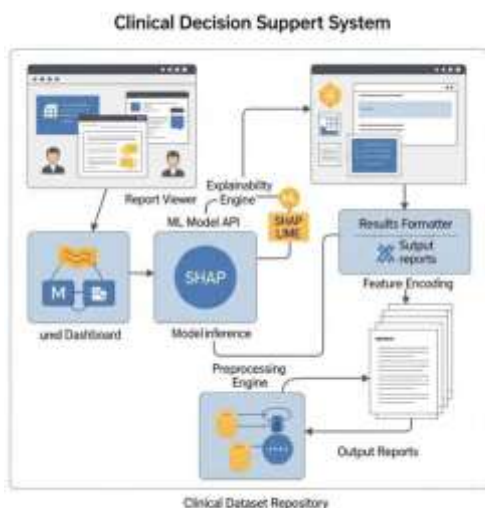
Provide Recommendations for Future Research

Encourage use of larger datasets, cross-disciplinary collaboration, and real-time AI tools. Future studies should focus on making ML models clinically interpretable, scalable, and deployable.

Workflow diagram:



System Architecture:



4. EXPECTED OUTCOME

Accurate Heart Disease Prediction Model:

Develop a machine learning model that can reliably predict the presence or risk of heart disease with high accuracy, precision, recall, and AUC by comparing multiple algorithms like Random Forest, SVM, and Neural Networks.

Effective Feature Selection:

Identify the most relevant clinical and demographic features that contribute significantly to heart disease prediction by applying various feature selection techniques.

Model Interpretability and Explainability:

Incorporate explainability tools such as SHAP or LIME to ensure the model's predictions are transparent and understandable, helping clinicians trust and adopt the model for real-world applications.

Comprehensive Evaluation:

Evaluate the models on diverse datasets to validate robustness and generalizability, including analysis of strengths and limitations of the applied ML techniques in the context of heart disease diagnosis.

Clinical Relevance:

Provide insights into how the model's predictions can aid early diagnosis and risk assessment in clinical settings, potentially supporting better patient outcomes through timely intervention.

3. CONCLUSION

The papers highlight the growing role of machine learning in heart disease prediction. Bhatt et al. use a large dataset with strong accuracy but lack clinical depth. Biswas et al. apply feature selection with good results but a small dataset. The literature review offers comparative insights without new

experiments. Hajiarbabi provides a detailed academic review by data type but is less practical. Ahire et al. focus on basic ML education. Random Forest and MLP models perform well overall, but most studies lack clinical validation, explainability, and integration with healthcare professionals, limiting real-world use. Future work should focus on bridging this gap.

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