
Heart Disease Prediction using Machine Learning

Students

Rishabh Sahu - 1900300130083

Yash Tripathi - 1900300130126

Atul Kr Gautam- 1900300130025

Inderprastha Engineering College, Ghaziabad

Abstract-

Predicting heart disease is one of the most challenging tasks in medicine in recent years. Today about one person dies from a heart attack every minute. Data science plays an important role in processing large amounts of data in healthcare. Because the prediction of heart disease is a difficult task, it is necessary to complete the forecasting process to avoid the risks associated with it and to warn patients in advance. This article uses the cardiology dataset available in the uci machine learning repository. Functional planning uses different types of data, such as naive Bayes, decision trees, logistic regression, and random forests, to predict heart disease risk and categorize population risk levels. Therefore, this article conducts a comparative study by analyzing the effectiveness of different learning systems. Experimental results confirmed that the random forest algorithm achieved the highest accuracy of 90.16% compared to other ml algorithms.

Keywords— *Decision Tree, Naive Bayes, Logistic Regression, Random Forest, Heart Disease Prediction*

INTRODUCTION

The study presented in this paper focuses solely on various data used in cardiovascular disease prediction. The human mind is the most important part of the human body. Basically, it controls the blood flow in the body. An imbalance in the heart can cause discomfort in other parts of the body. Any interruption in the functioning of the heart can be classified as a heart attack. Heart disease is one of the leading causes of death in the world today. Poor lifestyle, smoking, drinking alcohol, and consuming too much fat can all lead to high blood pressure, which can lead to heart disease.

According to the World Health Organization, more than 10 million people worldwide die from heart disease each year. A healthy lifestyle and early detection are the only ways to prevent heart disease.

The greatest challenge in healthcare today is to provide the best care and accurate and accurate diagnosis. Although heart disease has become a leading cause of death worldwide in recent years, it is also a disease that can be effectively controlled and managed. The exact accuracy of disease control depends on the correct time of detection of the disease. The strategy tries to detect these heart conditions early enough to prevent serious damage.

Huge medical data produced by doctors can be analyzed and valuable information can be extracted from them. Data mining is the process of extracting important and confidential information from large amounts of data. Many medical records contain conflicting information. Therefore, it will be difficult and difficult to make a decision with inconsistent information. Machine learning (ML), a subfield of data mining, can process large, well-structured data.

In the pharmaceutical industry, machine learning can be used to diagnose, detect and predict many diseases. The main purpose of this article is to provide physicians with tools for early detection of heart disease. This will help provide patients with better treatment and greater remission. Machine learning plays an important role in the analysis of data provided by the analysis of nonuniform patterns. After analyzing the data, machine learning helps predict and early detect heart disease.

This article focuses on Naive Bayes, Decision Trees, Logistic Regression and Random Forests for the Early Detection of Cardiovascular Diseases.

I. RELATED WORK

Many studies have been done to predict heart diseases using the UCI machine learning dataset. Different levels of accuracy are achieved using various data mining techniques described below. Previously, several different ML algorithms available for heart disease classification: decision trees, KNN and KMeans algorithms that can be used for classification are examined and their accuracy is compared.

This study concludes that decision trees provide the highest accuracy and adds that performance can be improved by combining different methods and metrics. Some expert system using data mining techniques together with the MapReduce algorithm is proposed.

According to this article, the accuracy obtained for 45 test sets is higher than that obtained using fuzzy neural networks. Here, the accuracy of the algorithms used is increased thanks to the use of dynamic models and linear scaling. Fahd Saleh Alotaibi developed a machine-learning model that compares five different algorithms. Using Rapid Miner tool is more accurate compared to Matlab and Weka tools. In this study, the accuracy of the decision tree, logistic regression, random forest, naïve Bayes, and SVM classification algorithms were compared. The decision tree algorithm has the highest accuracy. In this, the experts proposed a method using NB (Naive Bayesian) methods to classify data and AES (Advanced Encryption Standard) algorithm for secure data transfer to predict viruses. Trisa Principe UAV. R et al conducted a study involving different classification techniques to predict heart disease. The classification methods used are Naive Bayes, KNN (KNearest Neighbor), decision tree, neural network, and the accuracy of the classification has been analyzed for various attributes. Nagaraj M Lutimath et al. Heart disease prediction using Naive Bayes classification and SVM (Support Vector Machine). The performance measures used in the analysis are mean error, sum of squared error, and root mean squared error, and it can be concluded that SVM outperforms Naive Bayes in terms of accuracy.

After reviewing the above information, the main idea of the proposed method is to generate a heart disease prediction based on the material shown in Table 1. We analyze decision trees, random forest, logistic regression and distribution algorithms. Naive Bayes determines the best available for heart disease prediction classification algorithm based on the accuracy, precision, recall, and f-measure scores of the classification algorithm.

II. PROPOSED MODEL

The proposed project predicts heart disease and performs performance evaluation by exploring the four classification algorithms above. The purpose of this study is to predict whether a patient has heart disease. Healthcare professionals access valuable information from patients' medical records. The data is fed into a model that predicts the probability of having a heart attack. As shown in the picture.

1. shows the whole process.

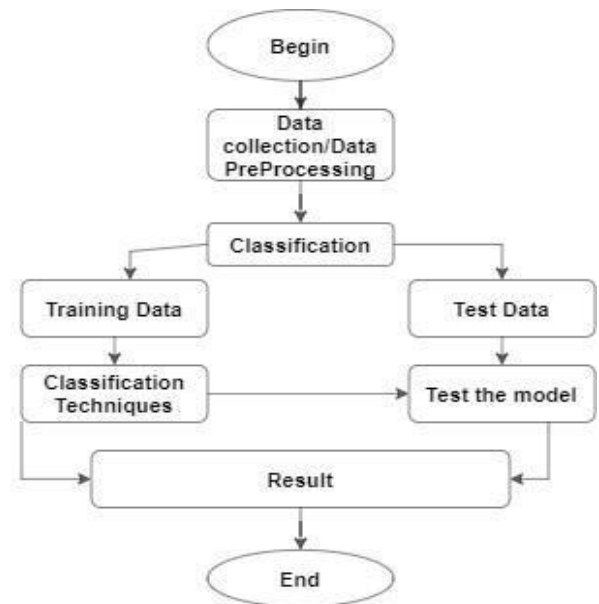


Fig. 1: Generic Model Predicting Heart Disease

A. Data Collection and Preprocessing

The data used by Kaggle dataset, which is a combination of 4 different data but uses only the UCI Cleveland dataset. The database contains a total of 76 features, but each published test uses only one of the 14 features. Therefore, we use the UCI Cleveland dataset already available on the Kaggle website for analysis. A full description of the 14 features used in the study is presented in Table 1 below.

TABLE I. FEATURES SELECTED FROM DATASET

Attribute	Description	Domain of value
Age	Age in years	29 to 77
Sex	Sex	Male (1) Female (0)
Cp	Chest pain type	Typical angina (1) Atypical angina (2) Non-anginal (3) Asymptomatic (4)
Trestbps	Resting blood sugar	94 to 200 mm Hg
Chol	Serum cholesterol	126 to 564 mg/dl
Fbs	Fasting blood sugar	>120 mg/dl True (1) False (0)
Restecg	Resting ECG result	Normal (0) ST-T wave abnormality (1) LV hypertrophy (2)
Thalach	Maximum heart rate achieved	71 to 202
Exang	Exercise induced angina	Yes (1) No (0)
Oldpeak	ST depression induced by exercise relative to rest	0 to 6.2
Slope	Slope of peak exercise ST segment	Upsloping (1) Flat (2) Downsloping (3)
Ca	Number of major vessels coloured by fluoroscopy	0–3
Thal	Defect type	Normal (3) Fixed defect (6) Reversible defect (7)
Num	Heart disease	0–4

B. Classification

The features specified in Table 1 are assigned to different machine learning algorithms such as random Forest, Decision Tree, Logistic Regression, and Naive Bayes classification techniques. The input data is divided into 80% of the training data and 20% of the test data. Training data is the data used to train the model. The test data algorithms reviewed in this article are listed below.

i).Random Forest

A random forest algorithm was used for regression and classification. It creates a tree for the data and makes predictions based on the tree; it can be used only on large data and can produce consistent results even on missing large datasets. It can save the structure created by decision trees so that it can be used in other files. There are two stages in the random forest, first it creates the random forest and then uses the random forest classifier created in the first stage to predict the next node.

ii).Logistic Regression

Logistic regression is a classification algorithm widely used in binary classification problems. In logistic regression, instead of fitting a line or hyperplane, the logistic regression algorithm uses a logistic function to compress the output of linear equations between 0 and 1.

iii)Decision Tree

A decision tree algorithm uses a flowchart where the middle row represents the dataset attributes and the outer branches represent the results. Decision trees were chosen because they are fast, reliable, easy to interpret and require little data preparation. In a decision tree, the prediction of the class list is taken from the root of the tree. The value of the base attribute is compared with the implicit attribute. Based on the ratio shown, follow the corresponding branch for that value and jump to the next link.

iv).Naive Bayes

Naive Bayes algorithm is based on Bayes' rule. The independence of the attributes of the dataset is a critical consideration, and is the most important factor when doing classification. Prediction is easy and fast and works best when the feeling of freedom persists. The Bayesian theorem calculates the final probability of event

(A), given the previous probability of event

B expressed by $P(A/B)$ [10], as shown in Equation 1,

$$P(A/B) = (P(B|A)P(A)) / P(B) \quad (1)$$

is used to check the effectiveness of the training model.

Performance for each algorithm is calculated and analyzed based on the different metrics used, such as accuracy, precision, regression, and F-test scores, as explained later. The different algorithms reviewed in this article are listed below

This section presents results using random forests, decision trees, pure Bayesian and logistic regression. The metrics used to drive the algorithm are accuracy score, precision (P), recall (R), and F-

measure. The precision (expressed in equation (2)) metric provides a measure of the accuracy of the analysis. Recall [expressed in Equation (3)] defines the measure of true quality. The F-

test [expressed in equation (4)] measures accuracy.

$$\text{Precision} = (TP) / (TP + FP) \quad (2)$$

$$\text{Recall} = (TP) / (TP + FN) \quad (3)$$

$$F\text{-Measure} = (2 * \text{Precision} * \text{Recall}) / (\text{Precision} + \text{Recall}) \quad (4)$$

- TP True Positive: The patient is infected and tested positive.
- FP False Positive: Patient has no disease but tests positive.
- TN True Negative: The patient has no disease and the test is negative.
- FN False Negative: The patient has disease but the test is negative.

In the experiment, previous data were used for testing and the above algorithms were investigated and applied. The above performance metrics are derived from the confusion matrix. The confusion matrix describes the performance of the model. The confusion matrices from the proposed models for the different algorithms are shown in Table 2 below. Accuracy scores obtained from random forest, decision trees, logistic regression and pure Bayesian classification method [12] are shown in Table 3 below.

TABLE II. VALUES OBTAINED FOR CONFUSION.MATRIX.USING DIFFERENT ALGORITHM

Algorithm	True Positive	False Positive	False Negative	True Negative
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III. RESULT AND ANALYSIS

Logistic Regression	22	5	4	30
Naive Bayes	21	6	3	31
Random Forest	22	5	6	28
Decision Tree	25	2	4	30

TABLE III. ANALYSIS OF MACHINE LEARNING ALGORITHM

Algorithm	Precision	Recall	Fmeasure	Accuracy
Decision Tree	0.845	0.823	0.835	81.97%
Logistic Regression	0.857	0.882	0.869	85.25%
Random Forest	0.937	0.882	0.909	90.16%
Naive Bayes	0.837	0.911	0.873	85.25%

IV. CONCLUSION

As the number of people dying from heart disease continues to rise, it has become more important to develop effective and accurate methods for predicting heart disease. The motivation for this work is to find the best ML algorithm for heart disease diagnosis. This study compares the accuracy scores of decision tree, logistic regression, random forest and Naive Bayes algorithms for cardiovascular disease prediction using the UCI Machine Learning Repository dataset. The results of this study showed that the random forest algorithm is the best algorithm for predicting heart diseases with an accuracy of 90.16%. This work can be improved in the future by creating a

website based on the Random Forest algorithm and using a larger dataset than used in this article. The analysis will help with better results and help doctors predict heart disease more effectively and efficiently.

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