# Heart Disease Prediction Using Machine Learning

Article in International Journal of Advanced Research in Science Communication and Technology · May 2021

DOI: 10.48175/IJARSCT-1131

CITATIONS

READS

19

28,331

5 authors, including:



Baban Uttamrao Rindhe

K. C. College of Engineering and Mangememt Studies and Research, Kopri, Thane (E)

55 PUBLICATIONS 115 CITATIONS

SEE PROFILE



International Journal of Advanced Research in Science, Communication and Technology (IJARSCT)

Volume 5, Issue 1, May 2021

# **Heart Disease Prediction Using Machine Learning**

Baban.U. Rindhe<sup>1</sup>, Nikita Ahire<sup>2</sup>, Rupali Patil<sup>3</sup>, Shweta Gagare<sup>4</sup>, Manisha Darade<sup>5</sup>
HOD and Professor, Department of Electronics and Telecommunication<sup>1</sup>
Students, Department of Electronics and Telecommunication<sup>2,3,4,5</sup>
K.C. College of Engineering and Management Studies & Research Thane, Maharashtra, India

Abstract: Heart-related diseases or Cardiovascular Diseases (CVDs) are the main reason for a huge number of death in the world over the last few decades and has emerged as the most life-threatening disease, not only in India but in the whole world. So, there is a need fora reliable, accurate, and feasible system to diagnose such diseases in time for proper treatment. Machine Learning algorithms and techniques have been applied to various medical datasets to automate the analysis of large and complex data. Many researchers, in recent times, have been using several machine learning techniques to help the health care industry and the professionals in the diagnosis of heart-related diseases. Heart is the next major organ comparing to the brain which has more priority in the Human body. It pumps the blood and supplies it to all organs of the whole body. Prediction of occurrences of heart diseases in the medical field is significant work. Data analytics is useful for prediction from more information and it helps the medical center to predict various diseases. A huge amount of patient-related data is maintained on monthly basis. The stored data can be useful for the source of predicting the occurrence of future diseases. Some of the data mining and machine learning techniques are used to predict heart diseases, such as Artificial Neural Network (ANN), Random Forest, and Support Vector Machine (SVM). Prediction and diagnosing of heart disease become a challenging factor faced by doctors and hospitals both in India and abroad. To reduce the large scale of deaths from heart diseases, a quick and efficient detection technique is to be discovered. Data mining techniques and machine learning algorithms play a very important role in this area. The researchers accelerating their research works to develop software with thehelp of machine learning algorithms which can help doctors to decide both prediction and diagnosing of heart disease. The main objective of this research project is to predict the heart disease of a patient using machine learning algorithms.

**Keywords**: Neural Network, Machine Learning, Supervised learning, Support vector machine, Random forest.

#### I. INTRODUCTION

Heart is an important organ of the human body. It pumps blood to every part of our anatomy. If it fails to function correctly, then the brain and various other organs will stop working, and within few minutes, the person will die. Change in lifestyle, work related stress and bad food habits contribute to the increase in the rate of several heart-related diseases. Heart diseases have emerged as one of the most prominent causes of death all around the world. According to World Health Organisation, heart related diseases are responsible for taking 17.7 million lives every year, 31% of all global deaths. In India too, heart-related diseases have become the leading cause of mortality. Heart diseases have killed 1.7 million Indians in 2016, according to the 2016 Global Burden of Disease Report, released on September 15,2017. Heart-related diseases increase the spending on health care and also reduce the productivity of an individual. Estimates made by the World Health Organisation (WHO), suggest that India has lost up to \$237 billion, from 2005-2015, due to heart-related or Cardiovascular diseases. Thus, feasible and accurate prediction of heart-related diseases is very important.

Medical organizations, all around the world, collect data on various health-related issues. These data can be exploited using various machine learning techniques to gain useful insights. But the data collected is very massive and, many times, this data can be very noisy. These datasets, which are too overwhelming for human minds to comprehend, can be

Copyright to IJARSCT DOI: 10.48175/IJARSCT-1131 267
www.ijarsct.co.in



# International Journal of Advanced Research in Science, Communication and Technology (IJARSCT)

Volume 5, Issue 1, May 2021

easily explored using various machine learning techniques. Thus, these algorithms have become very useful, in recent times, to predict the presence or absence of heart-related diseases accurately

The usage of information technology in the health care industry is increasing day by day to aid doctors in decision-making activities. It helps doctors and physicians in disease management, medications, and discovery of patterns and relationships among diagnosis data. Current approaches to predict cardiovascular risk fail to identify many people who would benefit from preventive treatment, while others receive unnecessary intervention. Machine-learning offers an opportunity to improve accuracy by exploiting complex interactions between risk factors. We assessed whether machine-learning can improve cardiovascular risk prediction

# II. LITERATURE SURVEY

ChalaBeyene et al[1], recommended Prediction and Analysis of the occurrence of Heart Disease Using Data Mining Techniques. The main objective is to predict the occurrence of heart disease for early automatic diagnosis of the disease within result in a short time. The proposed methodology is also critical in a healthcare organization with experts that have no more knowledge and skill. It uses different medical attributes such as blood sugar and heart rate, age, sex are some of the attributes are included to identify if the person has heart disease or not. Analyses of the dataset are computed using WEKA software.

Senthilkumar Mohan et al[2], implemented hybrid machine learning for heart disease prediction. The data set used is Cleveland data set. The first step is data pre-processing step. In this the tuples are removed from the data set which has missed the values. Attributes age and sex from data set are also not used as the authors think that it's personal information and has no impact on predication. The remaining 11 attributes are considered important as they contain vital clinical records. They have proposed their own Hybrid Random Forest Linear Method (HRFLM) which is the combination of Random Forest (RF) and Linear method (LM). In the HRFLM algorithm, the authors have used four algorithms. First algorithm deals with partitioning the input dataset. It is based on a decision tree which is executed for each sample of the dataset. After identifying the feature space, the dataset is split into the leaf nodes. Output of first algorithm is Partition of data set. After that in second algorithm they apply rules to the data set and output here is the classification of data with those rules. In third algorithm features are extracted using Less Error Classifier. This algorithm deals with finding the minimum and maximum error rate from the classifier. Output of this algorithm is the features with classified attributes. In forth algorithm they apply Classifier which is hybrid method based on the error rate on the Extracted Features. Finally they have compared the results obtained after applying HRFLM with other classification algorithms such a decision tree and support vector machine. In result as RF and LM are giving better results than other, both the algorithms are put together and new unique algorithm HRFLM is created. The authors suggest further improvement in accuracy by using combination of various machine learning algorithms.

Ali, Liaqat, et al[3],propose a system containing two models based on linear Support Vector Machine (SVM). The first one is called L1 regularized and the second one is called L2 regularized. First model is used for removing unnecessary features by making coefficient of those features zero. The second model is used for prediction. Predication of disease is done in this part. To optimize both models they proposed a hybrid grid search algorithm. This algorithm optimizes two models based on metrics: accuracy, sensitivity, septicity, the Matthews correlation coefficient, ROC chart and area under the curve. They used Cleveland data set. Data splits into 70% training and 30% testing used holdout validation. There are two experiments carried out and each experiment is carried out for various values of C1, C2 and k where C1 is hyperparameter of L1 regularized model, C2 is hyperparameter of L2 regularized model and k is the size of selected subset of features. First experiment is L1-linear SVM model stacked with L2-linear SVM model which is giving maximum testing accuracy of 91.11% and training accuracy of 84.05%. The second experiment is L1-linear SVM model cascaded with L2-linear SVM model with RBF kernel. This is giving maximum testing accuracy of 92.22% and training accuracy of 85.02. They have obtained an improvement in accuracy over conventional SVM models by 3.3%.

Singh, Yeshvendra K. et al[4],deal with various supervised machine learning algorithms such as Random Forest, Support Vector Machine, Logistic Regression, Linear Regression, Decision Tree with 3 fold, 5 fold and 10 fold cross-



# International Journal of Advanced Research in Science, Communication and Technology (IJARSCT)

Volume 5, Issue 1, May 2021

validation techniques. They have used Cleveland data set having 303 tuples, with some tuples having missing attributes. In the preprocessing of data they just removed the missing value tuple from the data set which are six in number and then from the remaining 297 tuples, they divided the data as training 70% and testing 30%. First algorithm applied is Linear Regression. In this, they have defined the dependency of one attribute over others which can be linearly separated from each other. Basically the classification takes place with the help of the group of attributes used for binary classification. They have obtained best results in 10 fold which is 83.82%. Logistic regression classification is done using a sigmoid function. This algorithm applied for heart disease prediction shows maximum accuracy with 3 and 5 fold cross-validation and it is 83.83%. Support Vector Machine is the classification algorithm in supervised machine learning. In this the classification is done by hyperplane.

The maximum accuracy achieved by SVM in 3 fold cross-validation is 83.17%. For Decision Tree in this paper, the authors have used different number splits and different number of leaf nodes to find the maximum accuracy. With 37 number splits and 6 leaf nodes maximum accuracy is achieved which is 79.12%. When used with cross-validation, accuracy achieved by the decision tree 79.54% with 5 fold. Random forest algorithm used on nonlinear data set gives better results as compared to the decision tree. Random forest is the group of decision tree created by the different root nodes. From this group of decision tree, voting can be done first and then classification can be done from the one getting maximum votes. Authors have used different number splits, different number of tree per observation and different number of folds for cross-validation. For random forest, 85.81% accuracy is achieved by 20 Number of splits, 75 Number of trees and 10 number of folds.

#### III. DATASET

We performed computer simulation on one dataset. Dataset is a Heart dataset. The dataset is available in UCI Machine Learning Repository [10]. Dataset contains 303 samples and 14 input features as well as 1 output feature. The features describe financial, personal, and social feature of loan applicants. The output feature is the decision class which has value 1 for Good credit and 2 for Bad credit. The dataset-1 contains 700 instances shown as Good credit while 300 instances as bad credit. The dataset contains features expressed on nominal, ordinal, or interval scales. A list of all those features is given in Table

S. no.	Feature name	Feature code	Description
1	Age	AGE	Age in years
2	Sex	SEX	Male = 1 Female = 0
3	Type of chest pain	CPT	a typical angina     typical angina     asymptomatic
4	Resting blood pressure Serum cholesterol	RBP SCH	4 — nonanginal pain mm Hg admitted at the hospital In mg/dl
6	Fasting blood sugar >120 mg/dl	FBS	Fasting blood sugar > 120 mg/dl (1 = true; 0 false)
7	Resting electrocardiog aphic results	RES	0 – normal 1 – having ST-T 2 – hypertrophy
8	Maximum heart rate achieved	MHR	
9	Exercise-induced angina	EIA	1 - yes 0 - no
10	Old peak = ST depression induced by exercise relative to rest	OPK	-
11	Slope of the peak exerc se ST segment	PES	1 – up sloping 2 – flat 3 – down sloping
12	Number of major vessels (0-3) colored by fluoroscopy	VCA	
13	Thallium scan	THA	3 – normal 6 – fixed defect



# International Journal of Advanced Research in Science, Communication and Technology (IJARSCT)

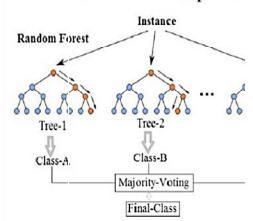
Volume 5, Issue 1, May 2021

# Attribute Selection Preprocessing Classification techniques SVM RF ANN Disease Prediction Accuracy Measure

#### 4.1 Random Forest

Random Forest is a supervised machine learning algorithm. This technique can be used for both regression and classification tasks but generally performs better in classification tasks. As the name suggests, Random Forest technique considers multiple decision trees before giving an output. So, it is basically an ensemble of decision trees. This technique is based on the belief that more number of trees would converge to the right decision. For classification, it uses a voting system and then decides the class whereas in regression it takes the mean of all the outputs of each of the decision trees. It works well with large datasets with high dimensionality

# Random Forest Simplified



# 4.2 Support Vector Machines (SVMs)

Support vector machines exist in different forms, linear and non-linear. A support vector machine is a supervised classifier. What is usual in this context, two different datasets are involved with SVM, training and a test set. In the ideal situation the classes are linearly separable. In such situation a line can be found, which splits the two classes perfectly. However not only one line splits the dataset perfectly, but a whole bunch of lines do. From these lines the best is selected as the "separating line".

A SVM can make some errors to avoid over-fitting. It tries to minimize the number of errors that will be made. Support vector machines classifiers are applied in many applications. They are very popular in recent research. This popularity is due to the good overall empirical performance. Comparing the naive Bayes and the SVM classifier, the SVM has been applied the most



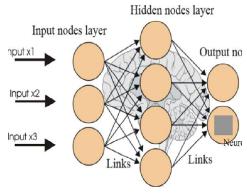
# International Journal of Advanced Research in Science, Communication and Technology (IJARSCT)

Volume 5, Issue 1, May 2021

#### 4.3 Artificial Neural Network

These are used to model/simulate the distribution, functions or mappings among variables as modules of a dynamic system associated with a learning rule or a learning algorithm. The modules here simulate neurons in nervous system and hence ANN collectively refers to the neuron simulators and their synapsis simulating interconnections between these modules in different layers.

Neural Network is built by stacking together multiple neurons in layers to produce a final output. First layer is the input layer and the last is the output layer. All the layers in between is called hidden layers. Each neuron has an activation function. Some of the popular Activation functions are Sigmoid, ReLU, tanh etc. The parameters of the network are the weights and biases of each layer. The goal of the neural network is to learn the network parameters such that the predicted outcome is the same as the ground truth. Back-propagation along loss-function is used to learn the network parameters.



V. SOFTWARE USED

#### 5.1 Python

To collect data a web scraper programmed in Python was used. According to Wikipedia Python's syntax allows programmers to express concepts in fewer lines of codes. Guido van Rossum at CWI in the Netherlands started Python's implementation in December 1989. Python 2.0 was released on October 16th 2000 and Python 3.0 was released December 3rd 2008.

Why use Python for web scraping and not another thing? Python offers a module called 'urllib2', which has suitable functions to open websites and extract information easily. Python is used to program the web scraper that is in charge of collecting the weather data for the model.

#### 5.2 MS Excel

Microsoft Excel is a spreadsheet application developed by Microsoft for Windows and Mac OS X. It features calculation, graphing tools, pivot tables and a macro-programming language. The first version was released in 1987. Why choose MS Excel versus another similar type of software? MS Excel is a very complete spreadsheet application tool, which supports almost any kind of file extension, and it has a lot of features. Its user-friendly interface helps you most of the time. However, if this doesn't seem enough, I will say that, apart from the typical things a normal user would do in Excel (Charts, Calculation...), it enables you to use the VBA language to create functions to use on the spreadsheets you've created. Excel can also be used as if it were an SQL database as was explained in a previous chapter. Having said this, for me it is the perfect program.MS Excel is used a lot throughout the project, to visualize the data and perform cleaning tasks on it.

# VI. RESULT AND DISCUSSION

This project aims to know whether the patient has heart disease or not [15]. The records in the dataset are divided into the training set and test sets. After preprocessing the data. The data classification technique namely support vector

Copyright to IJARSCT www.ijarsct.co.in

271



# International Journal of Advanced Research in Science, Communication and Technology (IJARSCT)

Volume 5, Issue 1, May 2021

machine, artificial neural network, random forest were applied. The project involved analysis of the heart disease patient dataset with proper data processing. Then, 3 models were trained and tested with maximum scores as follows:

1. Support Vector Classifier: 84.0 %

2. Neural Network: 83.5 %

3. Random Forest Classifier: 80.0 %

#### VII. CONCLUSION

This project provides the deep insight into machine learning techniques for classification of heart diseases. The role of classifier is crucial in healthcare industry so that the results can be used for predicting the treatment which can be provided to patients. The existing techniques are studied and compared for finding the efficient and accurate systems. Machine learning techniques significantly improves accuracy of cardiovascular risk prediction through which patients can be identified during an early stage of disease and can be benefitted by preventive treatment. It can be concluded that there is a huge scope for machine learning algorithms in predicting cardiovascular diseases or heart related diseases. Each of the above-mentioned algorithms have performed extremely well in some cases but poorly in some other cases.

#### ACKNOWLEDGEMENTS

We would like to express special thanks of gratitude to our guide Baban U. Rindhe as well as our project coordinator Ms. Sushma Kore, who gave us the golden opportunity to do this wonderful project on the topic Heart Disease Prediction Using Machine Learning, which also helped us in doing a lot of research and we came to know about so many new things. We would also like to thank our H.o.D. of EXTC – Baban U. Rindhe and Principal Vilas Nitnaware for providing us the opportunity to implement our project. We are thankful to both of them. Finally, we would also like to thank our department staff members and our parents & friends who helped us a lot in finalizing this project within the limited time frame.

#### REFERENCES

- [1]. Mr. ChalaBeyene, Prof. Pooja Kamat, "Survey on Prediction and Analysis the Occurrence of Heart Disease Using Data Mining Technique", International Journal of Pure and Applied Mathematics, 2018.
- [2]. Mohan, Senthilkumar, Chandrasegar Thirumalai, and Gautam Srivastava, "Effective heart disease prediction using hybrid machine learning techniques" IEEE Access 7 (2019): 81542-81554.
- [3]. Ali, Liaqat, et al, "An optimized stacked support vector machines based expert system for the effective prediction of heart failure" IEEE Access 7 (2019): 54007-54014.
- [4]. Singh Yeshvendra K., Nikhil Sinha, and Sanjay K. Singh, "Heart Disease Prediction System Using Random Forest", International Conference on Advances in Computing and Data Sciences. Springer, Singapore, 2016.
- [5]. Prerana T H M1, Shivaprakash N C2, Swetha N3 "Prediction of Heart Disease Using Machine Learning ,Algorithms- Naïve Bayes, Introduction to PAC Algorithm, Comparison of Algorithms and HDPS" International Journal of Science and Engineering Volume 3, Number 2 2015 PP: 90-99
- [6]. B.L DeekshatuluaPriti Chandra "Classification of Heart Disease Using K- Nearest Neighbor and Genetic Algorithm" International Conference on Computational Intelligence: Modeling Techniques and Applications (CIMTA) 2013.
- [7]. Michael W.Berryet.al, Lecture notes in data mining, World Scientific(2006)
- [8]. S. Shilaskar and A.Ghatol, "Feature selection for medical diagnosis :Evaluation for cardiovascular diseases," Expert Syst. Appl., vol. 40, no. 10, pp. 4146–4153, Aug. 2013.
- [9]. C.-L. Chang and C.-H. Chen, "Applying decision tree and neural network to increase quality of dermatologic diagnosis," Expert Syst. Appl., vol. 36, no. 2, Part 2, pp. 4035–4041, Mar. 2009.
- [10]. T. Azar and S. M. El-Metwally, "Decision tree classifiers for automated medical diagnosis," Neural Comput. Appl., vol. 23, no. 7–8, pp. 2387–2403, Dec. 2013. [10] Y. C. T. Bo Jin, "Support vector machines with



# International Journal of Advanced Research in Science, Communication and Technology (IJARSCT)

Volume 5, Issue 1, May 2021

genetic fuzzy feature transformation for biomedical data classification.," Inf Sci, vol. 177, no. 2, pp. 476–489, 2007.

- [11]. N. Esfandiari, M. R. Babavalian, A.-M. E. Moghadam, and V. K. Tabar, "Knowledge discovery in medicine: Current issue and future trend," Expert Syst. Appl., vol. 41, no. 9, pp. 4434–4463, Jul. 2014.
- [12]. E. Hassanien and T. Kim, "Breast cancer MRI diagnosis approach using support vector machine and pulse coupled neural networks," J. Appl. Log., vol. 10, no. 4, pp. 277–284, Dec. 2012.
- [13]. Sanjay Kumar Sen 1, Dr. Sujata Dash 21Asst. Professor, Orissa Engineering College, Bhubaneswar, Odisha India
- [14]. Domingos P and Pazzani M. "Beyond Independence: Conditions for the Optimality of the Simple Bayesian Classifier", in Proceedings of the 13th Conference on Machine Learning, Bari, Italy, pp 105-112,1996.
- [15]. Elkan C. "Naive Bayesian Learning, Technical Report CS97-557", Department of Computer Science and Engineering, University of California, San Diego, USA, 1997.
- [16]. B.L Deekshatulua Priti Chandra "Reader, PG Dept. Of Computer Application North Orissa University, Baripada, Odisha India. Empirical Evaluation of Classifiers Performance Using Data Mining Algorithm"