Lecture Notes in Networks and Systems 868

Sandeep Kumar Balachandran K. Joong Hoon Kim Jagdish Chand Bansal *Editors*

Fourth Congress on Intelligent Systems CIS 2023, Volume 1



Lecture Notes in Networks and Systems

Volume 868

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Fourth Congress on Intelligent Systems

CIS 2023, Volume 1



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Preface

This book contains outstanding research papers as the proceedings of the 4th Congress on Intelligent Systems (CIS 2023), held on September 4–5, 2023, at CHRIST (Deemed to be University), Bengaluru, India, under the technical sponsorship of the Soft Computing Research Society, India. The conference is conceived as a platform for disseminating and exchanging ideas, concepts, and results from academia and industry researchers to develop a comprehensive understanding of the challenges of intelligence advancements in computational viewpoints. This book will help in strengthening congenial networking between academia and industry. We have tried our best to enrich the quality of the CIS 2023 through the stringent and careful peer-review process. This book presents novel contributions to Intelligent Systems and is a reference material for advanced research in medical imaging and health informatics, agriculture, and engineering applications.

We have tried our best to enrich the quality of the CIS 2023 through a stringent and careful peer-review process. CIS 2023 received many technical contributed articles from distinguished participants from home and abroad. CIS 2023 received 870 research submissions from different countries. After a very stringent peer-reviewing process, only 104 high-quality papers were finally accepted for presentation and the final proceedings.

This book presents the first volume of 35 data science and applications research papers and serves as a reference material for advanced research.

Bengaluru, India Bengaluru, India Seoul, Korea (Republic of) New Delhi, India Sandeep Kumar Balachandran K. Joong Hoon Kim Jagdish Chand Bansal

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A Simple Way to Predict Heart Disease Using AI



Soumen Kanrar^(D), Suman Shit, and Subhadeep Chakrarbarti

Abstract Early diagnosis of cardiovascular diseases in high-risk patients will help them make decisions about lifestyle changes and, in turn, minimise their complications. Due to asymptomatic illnesses like cardiovascular diseases, healthcare costs are exceeding the average national medical treatment cost and corporate budgets. The need for early identification and treatment of such diseases is critical. One of the developments in machine learning is the technology that has been used for disease prediction in many fields around the world, including the healthcare industry. Analysis has been attempted to classify the most influential heart disease causes and to reliably predict the overall risk using homogeneous techniques of data mining. This paper uses machine learning algorithms and selects the best one based on its classification report to find a simple way to predict heart disease. It helps to develop cost-effective software to predict heart disease for the betterment of mankind.

Keywords Heart disease \cdot Chest pain \cdot Cardiovascular \cdot Machine learning \cdot AdaBoost \cdot Random forest

1 Introduction

Recent research has delved into amalgamating these techniques using methods such as hybrid data mining algorithms. To provide an approximate approach for the heart disease prediction, this paper has considered a rule-based model to compare the performance of applying regulations to the individual effects of classification and regression machine learning techniques. This work will follow these steps in order to develop a heart disease classifier. The well-known UCI heart disease dataset is used to develop the heart disease classifier. The purpose of this study is to classify the most important cardiac disease predictors and use classification and regression algorithms to predict the overall risks [1]. The classification algorithms in machine learning are thus used to classify the predictors [2]. Furthermore, to verify models,

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researchers conduct data analysis in Python using Jupyter Notebook [3]. Health policies and programs are frequently recommended based on the doctor's instructions rather than knowledge-rich data. Sometimes, this results in errors and high expenses, both of which have an impact on the quality of medical treatments. Clinical choices can be improved by using analytic tools and data modelling. As a result, the goal is to create a web application that will guide doctors in diagnosing cardiac disorders [4]. The main advantage is that if the doctors can make a proper diagnosis at the exact time, they can provide proper treatment at a considerable cost. This paper presents the exhibitor's analysis of the predictions of heart disease using algorithms for classification. These secret trends in medical data can be used for health diagnosis. Healthcare management may use the knowledge to get better treatment. For big data concerns and data science projects, the data science lifecycle has to be constructed first. Problem formulation, data collection, data preparation, data exploration, data modelling, model evaluation, and model deployment are the seven major steps normally used in data science. The human heart beats one hundred thousand times every day, pumping 7570.824 L of blood into the body. There are 96,560.64 km of blood veins inside the body. Men are at higher risk of heart attack than women, based on the basic symptoms. Generally, women may feel squeezing, pressure, fullness, or discomfort in the middle of their chest during a heart attack. It can also cause arm, back, neck, jaw, or stomach pain, as well as shortness of breath, nausea, and other symptoms. The basic symptoms observed for a heart attack in humans are chest pain, filling uneasiness, and mental stress. Sometimes, it is observed that humans have pain in their hands, arms, and shoulders, as well as in their neck, back, and jaw, as well as shortness of breath. To make the classification process more effective, the heart disease database is pre-processed. Classification and regression categories are the pre-processed data. In order to diagnose heart diseases early, it is important that health complications be reduced. In the current healthcare industry, machine learning has been commonly used to diagnose and predict the existence of diseases using data models. One of these relatively common machine learning algorithms for studies involving risk assessment is the classification technique for complex illnesses. Therefore, the research aims to classify the most relevant cardiovascular disease predictors and predict the overall risk using classification algorithms. On the Kaggle website, the dataset used for support vector machine, random forest, AdaBoost, and gradient boost analysis is available for various projects. To develop a classifier for the early stage of heart disease prediction, the researcher must collect and pre-process the data as prerequisites at the initial steps. Researchers used the UCI heart disease dataset for this purpose. The aim is to decide if the patient has a risk of future heart disease. The UCI dataset consists of 442 patient data records and 14 attributes. We have conducted the test through the Python program, which is run in a Jupyter Notebook. The Jupyter Notebook is equipped with a very powerful data science software package. For the last two decades, predicting heart attacks using data mining methods has gained importance. Much of the article has applied techniques to various patient datasets from around the world, such as SVM, neural networks, regression, decision trees, Naive Bayesian classifiers. The multiple regression model was also suggested in the research paper to forecast heart disease [5].

Maarten et al. (2022) addressed the issue of artificial intelligence needed to solve the targeted medical problem of cardiovascular disease [6]. It has been observed from the research papers that various linear regression models present the perfect conditions for predicting the chance of heart disease. The classification accuracy of the regression algorithm is good compared to other algorithms [7]. S. Das et al. have proposed the local discovery method of frequent disease prediction in the medical dataset [4]. S. Das et al. focus on examining information mining processes that are needed to classify for medical information mining, such as heart disease, lung malignancy, and bosom disease. The prediction algorithm used Naive Bayes analytic model, which is developed based on Baye's theorem [8]. Therefore, Naive Bayes could make independent assumptions [9]. Huang et al. have applied artificial intelligence to Wearable Sensor Data to identify and forecast cardiovascular disease [10]. V. Jackins et al. used a random forest classifier and Naive Bayes in the efficient prediction of clinical disease in the heart [11]. Big data tools such as Hadoop Distributed File System, MapReduce, and support vector machine are also considered for heart disease prediction [12]. Maarten et al. used the concept of data mining to forecast heart disease [6]. It is recommended to use HDFS on various nodes for storing a large volume of data and performing the prediction algorithm using SVM simultaneously. Concurrently, these results are obtained in faster computation time compared to other machine learning models. Another repository from which we can collect the sourced data is one of the leading institutes for diabetic research in Chennai, Tamil Nadu, India [13]. The dataset consists of more than 500 patients' data. In another dataset, the instrument implemented in Weka considers a seventy percent break for classification. Here, 86.419% of the data is the precision provided by Naive Bayes. Neurological computing is a little bit helpful for analysing the electric signal from the heart to determine the cardiovascular issue [14]. Ghrabat et al. [15] discussed the use of a nonlinear classification algorithm for the estimation of heart disease [16]. On the Kaggle website, the dataset uses support vector machine, random forest, AdaBoost, and gradient boost analysis, which are available from ongoing Kaggle projects. The aim of our study is to determine if the patient has a risk of future heart disease. We have conducted the study considering a training dataset consisting of 3000 cases with thirteen unique attributes. The dataset is separated into two clusters. One cluster contains seventy percent of the data that is used for training purposes, and another cluster contains thirty percent of the data that is considered for testing purposes.

2 Materials and Methods

In the body, the heart is like any other muscle. For the muscle to contract and pump blood to the rest of the body, it requires enough blood flow to deliver oxygen. The heart pumps blood into the coronary arteries. Arteries derive from the aorta (the largest artery of the body, which contains the heart's oxygenated blood) and then spread out along the heart wall. Table 1 presents the different types of heart disease with their common symptoms and causes. If humans change certain habits in their lives, they can reduce the chance of heart disease.

These include:

- Consume fibre-rich, nutritious diets.
- Regular exercise: These will help to improve the heart condition and circulatory system, decrease cholesterol, and maintain blood pressure.
- Control the body weight with respect to height.
- Control alcohol consumption: Per day may be consumed according to the prescription of the doctor.
- Managing underlying issues: Finding care for problems such as elevated blood pressure, obesity, and diabetes that impair heart well-being.
- Quitting or avoiding smoking: Smoke is an important risk factor for cardiovascular and heart disease.

Machine learning is commonly used in nearly all sectors of the healthcare industry around the world. ML technology helps to adopt new formations from experience. In comparison, the process of considering various methodologies to predict, forecast, and explore something in the universe based on machine learning is relatively simple [17]. Sometimes the complex heart disease problem can be predicted by machine learning, i.e. regression and classification. Regression algorithms are primarily used for numeric results and binary and multi-category problems. Algorithms for machine learning were further split into two groups, such as supervised learning and unsupervised learning. Supervised learning does not possess fixed tags; consequently, it infers the spontaneous build-up in the dataset. Therefore, machine learning algorithm selection needs to be properly analysed. AdaBoost is an ensemble strategy in which a group of weak learners are joined to generate a powerful learner. The AdaBoost model is present in Fig. 1. Typically, each weak learner is created as a "decision stump" (a tree with only one major split). In the random forest procedure, different

S. No.	Disease types								
	Heart disease types	Symptoms	Reason						
1	Coronary artery disease	Chest pain or discomfort (angina)	Plaque build-up in the arteries						
2	Congenital heart defects	Rapid heartbeat and breathing	Several genetic health conditions						
3	Arrhythmia	A fluttering in the chest and a racing heartbeat	Low supply of oxygen and nutrients to the heart						
4	Heart failure	Out of breath. Feel tired always. Swollen ankles and legs	High blood pressure and high cholesterol						
5	Myocardial infarction	Chest pain, fatigue, and heartburn	Smoking and high cholesterol						

Table I Heart	disease
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weights are assigned to increase efficiency. When we need high accuracy, we need to assign more weight. Tree is created from the consecutive training set. AdaBoost can improve its efficiency by anticipating the updated weights. Gradient Boosting Machine (GBM), like AdaBoost, combines a number of weak learners to generate a powerful learner. The classifier's residual is used as the input for the next classifier, on which the trees are created, making this an additive model. The residuals are acquired by the classifiers in a step-by-step manner to bag major variation in the data, which is accomplished through massive learning. The Gradient Boosting Machine is present in Fig. 2. It is gradually moving in the correct direction towards improved prediction using this strategy. As a result, based on the number of classifiers, we arrive at a prediction value that is very close to the observed value. Initially, a single-node tree is constructed that predicts the aggregated value of Y in the case of regression or the log (odds) of Y in the case of classification issues, following which trees with greater depth are developed on the residuals of the preceding classifier. Olsen et al. [18] provide a good overview of machine learning procedures for the prediction of heart failure [18]. Joon-myoung et al. (July 2019) used a deep learning-based artificial intelligence algorithm to predict acute heart failure [19]. Akbilgic et al. [20] developed an electrocardiographic artificial intelligence model to predict heart failure [20]. They have used the electrocardiogram artificial intelligence model for the prediction of heart failure. Manimurugan, S., et al. used the Two-Stage Classification model for the identification of heart disease [21].

In GBM, each tree has 8–32 terminal nodes, and approximately, 100 trees are produced. For each tree, learning rates are set to a constant so that the model may take modest steps in the appropriate direction to capture variance and train the classifier. Unlike in AdaBoost, where all trees are given equal weights. The dataset used



Fig. 1 AdaBoost model



Fig. 2 Gradient boosting machine model

in the study of classification and regression algorithms is from an ongoing UCI cardiovascular study and is available on the Kaggle website.

The object of this analysis for classification is to predict that the patient has a chance of potential heart failure from the infections. With 442 pieces of information, the Framingham dataset consists of patient data and 14 attributes. Reviewing the Framingham data set and using Jupyter Notebook in Python programming, which is a more flexible and efficient technology tool for data science. Figure 3 indicates the steps followed to build the classification and regression models in machine learning.

In the data pre-processing stage, the initial process involves cleaning the data first to remove noise, identify inconsistent data, and obtain missing data. The initial dataset attributes are presented in Table 2. The regression model is created using the UCI Machine Learning Repository.



Fig. 3 Workflow of classification and regression model

Attribute name	Attribute description, type, value								
	Description	Type/value							
Age	Age in years	Integer							
СР	Chest pain	Common angina	a	Value	= 0				
		Anomalous ang	ina	Value	= 1				
		Non-angina pair	n	Value	= 2				
		No symptoms		Value	= 3				
trestbps	Resting blood pressure	In mm Hg							
chol	Serum cholesterol	In mg/dl							
fbs	Fasting blood sugar	More than	True		Value $= 1$				
		120 mg/dl	False		Value $= 0$				
restecg	Resting	Normal			Value $= 0$				
	electrocardiographic	ST-T wave abno	ormality	Value $= 1$					
		Left ventricular	hypertr	ophy Value = 2					
thalach	Heart rate	Heart rate/minir	num						
Exang	Angina	Yes		Value	= 1				
		No		Value = 0					
oldpeak	Depression	Exercise relative	e to rest						
slope	Slope of the peak	Upsloping		Value $= 0$					
	exercise	Flat		Value $= 1$					
		Downsloping		Value $= 2$					
ca	Number of major vessels	Coloured by fluoroscopy		Value	range (0–3)				
thal	Thalassemia	Normal		Value	= 3				
		Fixed defect		Value = 6					
		Reversible defe	et	Value = 7					
Gender	Male	Value = 1							
	Female	Value = 0							

Table 2 Dataset attributes

3 Data Handle

The regression model is created using the UCI Machine Learning Repository. We have developed Python language code and tested it on the Jupiter notebook platform after removing some unnecessary fields from the collected data. In addition, the number of missed values for cleaning up current databases has been identified. A sample snapshot of the data header is presented in Table 3. The dataset is error-free and contains all of the data required for each variable. There are no issues regarding missing values or inconsistencies that are identified using the info (), describe (), and isnull () functions as defined in the Pandas Library. As a result, the dataset is

fairly balanced. From the collected data, the proposed techniques approximate the (0.56) probability portion of the human population suffering from heart disease in the sample dataset presented in Table 3. Figure 4 presents the heart disease of the people with respect to the small sample dataset present in Table 3.

To build the attribute correlation, we consider the heatmap, which depicts the relationships between the dataset's attributes as well as how they interact. Attribute correlation is presented in Fig. 5.

Generally, from the observed prediction, we draw the inference that older people suffer from heart disease. The heart rate becomes lower at old age as he or she suffers from heart disease. The correlation between age and heart rate is presented in Fig. 6. The heatmap shows that the types of chest pain (CP), exercise-induced angina (Exang), exercise-induced ST depression, atypical, and non-anginal are all strongly linked to heart disease (target). We also see that cardiac illness and maximum heart rate have an inverse relationship (thalamus). It is presented in Fig. 7.

	Age	Sex	CP	trestbps	chol	fbs	restecg	thalach	Exang	oldpeak	slope	ca	thal	Target
0	70	1	4	130	322	0	2	109	0	2.4	2	3	3	2
1	67	0	3	115	564	0	2	160	0	1.6	2	0	7	1
2	57	1	2	124	261	0	0	141	0	0.3	1	0	7	2
3	64	1	4	128	263	0	0	105	1	0.2	2	1	7	1
4	74	0	2	120	269	0	2	121	1	0.2	1	1	3	1

 Table 3
 Sample dataset





ar l	1	-0.094	0.097	0.27	0.22	0.12	0.13	-0.4	0.098	0.19	0.16	0.36	0.11	0.21
ă.	-0.094	x	0.035	-0.063	-0.2	0.042	0.039	-0.076	0.18	0.097	0.051	0.087	0.39	
	0.097	0.035	.1	-0.043	0.09	-0.099	0.074	-0.32		0.17	0.14	0.23	0.26	
-		-0.063	-0.043	1	0.17	0.16	0.12	-0.039	0.083	0.22	0.14	0.086	0.13	0.16
	0.22	-0.2	0.09	0.17	1	0.025	0.17	-0.019	0.078	0.028	-0.0058	0.13	0.029	0.12
	0.12	0.042	-0.099	0.16	0.025	1	0.053	0.022	-0.0041	-0.026	0.044	0.12	0.049	-0.016
R.u.	0.13	0.039	0.074	0.12	0.17	0.053	1	-0.075	0.095	0.12	0.16	0.11	0.0073	0.18
	-0.4	-0.076	-0.32	-0.039	-0.019	0.022	-0.075	-1	-0.38	-0.35	-0.39	-0.27	-0.25	-0.42
	0.098	0.18	0.35	0.083	0.078	-0.0041	0.095	-0.38	4	0.27	0.26	0.15	0.32	0.42
	0 19	0.097	0.17	0.22	0.028	-0.026	0.12	-0.35	0.27	1	061	0.26		
	0.16	0.051	0.14	0.14	-0.0058	0.044	0.16	-0.39	0.26	0 6 1	1	0.11		
		0.087	0.23	0.086	0.13	0.12	011	-0.27	0.15	0.26	0.11	1	0.26	
-	0.11	0.39	0.26	0.13	0.029	0.049	0.0073	-0.25				0.26	1	
	0.21			0.16	0.12	-0.016	0.18	-0.42					0.53	1
	age	sex	æ	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	target

Fig. 5 Attributes correlation



Fig. 6 Correlation between age and heart rate



Fig. 7 Heart disease based on chest pain

4 Experiment and Results

From the result analysis, we get four different types of chest pain as the main cause of heart disease. The majority of heart disease patients have asymptomatic difficulty breathing.

The majority of patients with heart disease are elderly and have one or more main arteries coloured by Fluoroscopy. It is presented in Fig. 8.

We have logically subdivided the dataset. As some columns treat it as the "target" column, it is considered to serve as the class. A training set and a test set are created from the data. Seventy percent of the data is used for training purposes, and the remaining amount is used for testing purposes. Finally, the test results for the models for support vector machine, random forest, AdaBoost, and gradient boosting are exhibited in Figs. 9, 10, 11 and 12. Inference can be considered based on the best classification report.

Based on the aforementioned obtained results, with component-wise comparison based on precision, we can consider that gradient boosting is one of the best classifiers for heart disease prediction. The correlation between number of major vessels colored by flourosopy and age





classificatio	on_report :			
	precision	recall	f1-score	support
1	0.71	0.90	0.79	30
2	0.81	0.54	0.65	24
accuracy			0.74	54
macro avg	0.76	0.72	0.72	54
weighted avg	0.76	0.74	0.73	54

Fig. 9 Snap of the support vector machine used in the dataset

RandomForestClassifier

classificatio	n_report :			
	precision	recall	f1-score	support
1	0.76	0.73	0.75	30
2	0.68	0.71	0.69	24
accuracy			0.72	54
macro avg	0.72	0.72	0.72	54
weighted avg	0.72	0.72	0.72	54

Fig. 10 Snap of the random forest classifier used in the dataset

classificati	on_report : precision	recall	f1-score	support
1	0.75	0.80	0.77	30
2	0.73	0.67	0.70	24
accuracy	•		0.74	54
macro ave	0.74	0.73	0.73	54
weighted ave	0.74	0.74	0.74	54

Fig.11 Snap of the AdaBoost classifier used in the dataset

AdaBoostClassifier

GradientBoostingClassifier

classificatio	n_report :			
	precision	recall	f1-score	support
1	0.79	0.73	0.76	30
2	0.69	0.75	0.72	24
accuracy			0.74	54
macro avg	0.74	0.74	0.74	54
weighted avg	0.74	0.74	0.74	54

Fig.12 Snap of the gradient boosting classifier used in the dataset

5 Conclusion

In conclusion, we learned how to build a suitable application for drawing predictions about heart diseases from medical datasets. We began with the definition of the problem and the moderate amount of accumulated data. Then we worked on data preparation, investigation, and building models. To create a classification model, we used a standardised dataset of heart disease patients. We began with data exploration and then moved on to data preparation. After that, we tested four heart disease classification models. Finally, we chose the finest model and saved it for later usage. The limitations of these works focus primarily on the use of classification strategies and algorithms for the prediction of heart disease. In this domain, researchers use different mechanisms for data cleaning and plan to construct a dataset that is suitable for their desired algorithms.