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PAPER

Optimizing Machine Learning Algorithms for Heart Disease Classification and Prediction

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ABSTRACT

According to the World Health Organization (WHO), cardiovascular disease is one of the leading causes of death worldwide. Thus, the prevention of this kind of illness is considered as a huge human health challenge. Additionally, the diagnostic process often involves a combination of clinical examination, laboratory tests, and other diagnostic procedures, which can be complex and time-consuming. However, advances in medical technology and research have led to improved methods for diagnosing heart disease, which can help to improve patient outcomes. Furthermore, Machine Learning (ML) methods have shown promise in helping to improve the diagnosis of heart disease. Each method requires specific parameters to produce good results. In this paper, we propose a diagnosis support system based on optimized Machine Learning algorithms, which is Artificial Neural Network (ANN), Support Vector Machine (SVM), K_Nearest Neighbour (KNN), Naive Bayes (NB), and Decision Tree (DT) to analyze the major cardiovascular risk factors, such as age, gender, high blood pressure, etc. To train and validate the ML models, a medical dataset of 558 patients with atherosclerosis is used. In this work, we achieved a 96.67% as promising accuracy level for the atherosclerosis prediction with ANN.

KEYWORDS

machine learning, atherosclerosis, cardiovascular disease, optimization

1 INTRODUCTION

Atherosclerosis, which is a type of heart disease, is largely caused by the accumulation of cholesterol deposits, called plaques, on the walls of the arteries. These plaques can restrict or block the flow of blood, leading to a range of serious health problems [1]. When the plaques build up in the coronary arteries, which supply blood to the heart muscle, it can cause angina (chest pain) or a heart attack. When plaques build up in the arteries that supply blood to the brain, it can cause a stroke. When plaques build up in the peripheral arteries, it can cause Peripheral Arterial Disease (PAD), which can lead to pain, numbness, and even amputation of the

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affected limb. Atherosclerosis is a chronic disease that develops over many years, and it is one of the leading causes of death globally [2]. The prevention of cardiovascular disease has become a major human health challenge in recent years. Early treatment of risk factors will reduce cardiovascular disease and serious long-term complications, as recommended by physicians [3].

Extracting useful information from patient records can be a time-consuming and challenging task, especially when dealing with large amounts of unstructured data, such as electronic health records. The manual process of extracting and analyzing this data can lead to errors and inconsistencies, which can affect the accuracy of the diagnosis. Artificial Intelligence (AI) techniques, particularly Machine Learning (ML) [4], have been widely used in medical diagnosis and treatment systems to automate the classification of patients and the prediction of cardiovascular diseases. These techniques have been applied to a wide range of domains, beyond healthcare [5]–[7], including: Medical image segmentation [8], [9], COVID-19 Disease [10], Natural Language Processing (NLP) [11], Image processing [12], Industrial automation, Predictive modelling and Diagnosis of Neurological disorders [13]. With the help of ML, we can process large datasets, identify patterns, and make predictions that would be difficult or impossible to do manually [14]. Early detection of heart disease can help patients make lifestyle changes that may reduce the likelihood of developing heart disease. Moreover, these changes allow for timely medical intervention, which can prevent or delay the onset of heart disease and its complications [15], [16].

In this work, we proposed an ML-based diagnostic system based on five optimized ML methods. The goal is to classify patients and predict heart diseases using Cleveland and Hungarian heart disease databases. These datasets are databases that use both controllable and uncontrollable risk factors, such as obesity, hypertension, smoking, diabetes, cholesterol, age, family history, and gender. The main contributions of this research can be summarized as follows:

- We developed and implemented an automatic medical diagnosis support system for classifying patients and predicting heart disease;
- We adjusted the parameters of the used ML algorithms to obtain optimal performance;
- The structure of the proposed system allows its use to be extended to the prediction of other diseases;
- In comparison with state of the art, the proposed system tested on two wellknown datasets reach high performance.

The rest of the paper is organized as follows. In Section 2, we provide a general review of the related work, discussing previous studies and approaches in the field of ML-based diagnostic systems for heart disease. Section 3 presents the details of the proposed ML-based diagnostic system including its process and implementation, as well as the performance measures used for its evaluation. In Section 4, we afford a detailed analysis of the results obtained from the implementation of the proposed methods. Section 5 is dedicated to comparing the results obtained by this system with the results of related work. This section provides a comparison of the performance of the proposed system with other existing ML-based diagnostic systems for heart disease detection. Finally, in Section 6, we conclude this work by summarizing the main findings and discussing the potential future work.

2 RELATED WORK

In recent years, many researchers have proposed various methods for the diagnosis of atherosclerosis disease. In [17], authors used a set of fuzzy rules. The weighting of these rules used to give more importance to certain rules over others, and this method has been found to have an accuracy of 57.85%. In another work published in [18], the researchers used neural network ensemble method to build new models by combining the predicted values of several predecessor models. The obtained accuracy by this method was 89.01%. Furthermore, another proposed method in [19], called hybrid-genetic consists of increasing the performance of the neural network by improving its initial weights by genetic algorithm. The accuracy obtained by this method was 93%. In [20], the authors applied the Fast Decision Tree and c4.5 methods, and used an integration of the outcomes of the ML methods implemented on different datasets targeting CAD. The accuracy obtained by this approach was 78.06%. In [21], the authors proposed a new Intelligent Heart Disease Prediction System based on three ML Methods to predict patients affected by atherosclerosis. The highest accuracy achieved was 89%. Similarly, in [22], the authors proposed a decision support system based on the integration of the ANN and the Fuzzy-AHP methods. The accuracy achieved by this approach was 91.10%. Moreover in [23], the authors implemented five data mining algorithms in prediction heart diseases. The best performance was achieved by DT method, where the accuracy was 93.02%. Recently, the authors in [24] have proposed an MDSS for heart disease. This system is based on Fuzzy Analytic Hierarchy Process (AHP) and Fuzzy Inference System (FIS). The results show the efficiency and accuracy of this approach. In [25], the authors have proposed hybrid random forest with a linear model (HRFLM) for atherosclerosis early prediction. The accuracy reached was 88.7%.

Artificial intelligence techniques are widely used in the healthcare field, where they are used to diagnose many diseases other than heart diseases such as retinopathy [26] and Alzheimer's Disease [27]. ML techniques are effective in many fields, such as healthcare, for example, to recognize human images and their abnormal activity [28], [29].

The main objective of this work is to propose a new medical diagnosis aid system based on five optimized machine learning methods, to classify patients and predict cardiovascular diseases with an optimal accuracy rate. The methods we used are Artificial Neural Network (ANN), K-nearest neighbor (KNN), Naive Bayes (NB), Support Vector Machine (SVM) and Decision Tree (DT). The role of this system is to compute the performance of each algorithm and choose the model that gives the best performance by optimizing the main parameters.

3 THE PROPOSED METHODOLOGY AND IMPLEMENTATION

3.1 Overview of the proposed system

The proposed ML based diagnostic system is built using several steps, which are: (i) Understanding problems and objectives. (ii) Data mining and analysis. (ii) Preparation of data sets and choice of parameters. (iv) Execution of algorithms and search for optimal parameters. (v) Calculation of performance and selection of best algorithm. (vi) Presentation of results. The flowchart shown in Figure 1 provides a visual representation of the steps and how they are interconnected.

Fig. 1. Proposed system architecture

3.2 Machine learning algorithms

In this work, we have chosen to use five ML algorithms to classify patients and predict heart diseases. For our choice, we relied on the results of previous studies. The analysis of these results allowed us to see that the algorithms chosen to give good results when dealing with databases have a similar structure to the databases that we used. The chosen algorithms are ANN, KNN, NB, DT, and SVM. Then, we evaluate the performance of each algorithm using some evaluation metrics and choose the model that gives the best performance.

Artificial neural network (ANN). The artificial neural network is a biologically inspired ML method that attempts to simulate the tasks of neurons in the human brain [30]. Figure 2 presents an ANN sample. A neural network-based algorithm learns to make predictions by adjusting the weights and biases of its

neurons based on the input and output data. Where x_i design the Input, w_i is the weights associated with *xⁱ* . After several training operations and using several optimization methods, we obtained improved results with the parameters shown in the Table 1.

Fig. 2. Artificial neuron model

Parameter	Cleveland Dataset		Hungarian Dataset	
Learning Rate	0.0001		0.0001	
Layers number	5		7	
Neuron number	Input	64	Input	128
	Hidden 1	32	Hidden 1	68
	Hidden 2	13	Hidden 2	34
	Hidden 3	8	Hidden 3	17
	Output	$\mathbf{1}$	Hidden 4	16
			Hidden 5	$\overline{9}$
			Output	$1\,$
Activation functions	Input	Relu	Input	Relu
	Hidden 1	Relu	Hidden 1	Relu
	Hidden 2	Relu	Hidden 2	Relu
	Hidden 3	Relu	Hidden 3	Relu
	Output	Sigmoid	Hidden 4	Relu
			Hidden 5	Relu
			Output	Sigmoid
Training percentage	90%			
Testing percentage	10%			
Weights and bias	Random			
Loss function	Binary crossentropy			
Epochs	383		950	
Optimazer	Adam		RMSprop	

Table 1. Neural network hyperparameters

Support vector machine (SVM). Support Vector Machine (SVM) is a supervised ML algorithm originally introduced for the classification problems and later extended to several different situations. A linear or non-linear algorithm that finds the hyperplane maximally separates different classes in the feature space, called the optimal hyperplane. In the implementation step we use 90% for training and 10% for testing, the hyperparameters for this algorithm are shown in Table 2.

Decision tree (DT). The decision tree (DT) is a tree-based algorithm that recursively partitions the feature space into smaller regions, to make predictions based on the majority class in each region. This method was applied for classification and regression cases. A DT structure includes a node corresponding to a feature (attribute), a link (branch) corresponding to a decision (rule) and a leaf corresponding to a result. The algorithm works by dividing the information into two parts or subsets that support the values of the input variables. This algorithm is represented by a tree which contains nodes and leaves. This algorithm was trained with 90% of data, the features are split by the Gini impurity and the maximum depth chosen is 3.

The naïve bayes algorithm. A probabilistic algorithm that makes predictions based on the probability of each class given the input features. This is a supervised algorithm which is used for classification, using the equation:

$$
\hat{p}(Y = i | X_1, X_2, ..., X_3) = \frac{\pi(Y = i) \prod_{j=1}^{q} P(X_j | Y = i)}{\sum_{j=1}^{K} \pi(Y = i) \prod_{j=1}^{q} P(X_j | Y = i)}
$$
(1)

K-Nearest neighbors (KNN) algorithm. The KNN is a supervised learning algorithm that uses a distance metric to classify data points based on the proximity of the k nearest training examples [31]. The best results are obtained when we use the Euclidean distance for both datasets, and the odd k value $K = 23$ for the Cleveland dataset, while the k value for Hungarian dataset was $k = 18$.

3.3 Description of datasets

Cleveland dataset. David Aha collected this database, which contains 303 samplers and 76 attributes but only 298 samplers and 14 attributes are used due to missing data. This database contains 53.9% of healthy cases and 46.1% of coronary cases. The patients who are healthy are marked with a label of "0", while patients who are sick are marked with a different label, such as "1". This is a common approach in binary classification problems, where the goal is to predict one of two outcomes (in this case, healthy or sick). Figure 3 shows the distribution percentage of positive and negative cases for Cleveland dataset.

Fig. 3. Distribution of the Cleveland database

Hungarian dataset. This database is collected by Andras Janosi at the Institute of Cardiology, Budapest. It contains 294 patients and 10 attributes, only 262 members are used due to missing data. This database consists of 62.5% sick patients and 37.5% healthy patients. Figure 4 shows the distribution percentage of positive and negative cases for Hungarian dataset.

Fig. 4. Distribution of the Hungarian database

3.4 Selection of characteristics

After cleaning the databases by ignoring entries with missing data, the inputs data for each database are risk factors that have been identified by medical experts as having a clear influence on heart disease. In this work, we relied on the literature mentioned above that used the Cleveland or Hungary datasets for the selection of these features, and we use the correlation coefficient method to verify the importance of these features. Figures 5 and 6 show the correlation results with the target in the Cleveland and Hungary datasets.

Fig. 5. Correlation with target in Cleveland dataset

The corresponding output used in these two databases for classification and prediction are binary labels indicating the real status of the patient. There are two classes: a patient with heart disease, which signifies a diameter narrowing of more than 50%, or a patient without heart disease, which signifies a diameter narrowing of less than 50%, based on the UCI Cleveland and Hungary databases. The goal is to classify patients as having or not having heart disease based on the input risk factors and the binary labels. Table 3 shows the characteristics selected for each database.

3.5 Performance evaluation metrics

Confusion matrix: A binary confusion matrix is an array used to assess the performance of a binary classifier model on a set of test data for which the true values are known. The matrix is of the dimension 2x2 that contains the following four elements: True positive (TP), False positive (FP), True negative (TN), False negative (FN).

The binary confusion matrix is widely used to evaluate the performance of binary classification models and it is used to calculate several performance metrics such as Specificity (SP), Sensibility (SS), Accuracy (ACC), F1-score (FS) and Matthews's correlation coefficient (MCC).

$$
\text{ACC} = \frac{TP + TN}{TP + TN + FP + FN} \tag{2}
$$

$$
SS = \frac{TP}{TP + FN}
$$
 (3)

$$
SP = \frac{TN}{TN + FP}
$$
 (4)

$$
FS = \frac{2*TP}{2*TP + FP + FN}
$$
\n⁽⁵⁾

$$
MCC = \frac{TP * TN - FP * FN}{\sqrt{(TP + FP) * (TP + FN) * (TN + FP) * (TN + GN)}}
$$
(6)

4 RESULTS OF PERFORMANCE MEASURES

To determine which hyperparameter of each algorithm performs best, we use commonly used ML technologies such as Manual Search, Grid Search and Randomized Search. Then, we trained and evaluated the models with different combinations of hyperparameter values. The goal is to find the combination of hyperparameter values that result in the best performance of the model, as measured by the ACC, SS, SP, MCC, and F1-score.

Splitting data is a common technique used in ML to divide a dataset into two or more subsets, such as a training set and a test set. The goal of splitting the data is to use the training set to train the model and the test set to evaluate the model's performance. In the case we mentioned, the Cleveland and Hungarian datasets are each

split into two parts, 90% for training and 10% for testing. The training set is used to train the model and optimize the hyperparameters of each algorithm, while the test set is used to evaluate the model's performance. After the training process, we evaluate each method using the performance metrics mentioned above by the test data. Table 4 shows the confusion matrix results. the remainder of the performance metrics are provided in Table 5. As we can see from the results presented in the Tables 4 and 5, the ANN model gives the best results compared to the other MLs techniques. Most importantly, we have achieved excellent results indicating the quality of the proposed system.

Furthermore, the results of the ROC curve metric are consistent with the above results, this metric uses threshold values for every classifier over the interval [0, 1] to the output field. Two major values are then calculated for each threshold, the TPR and the FPR. The result is shown in the Figures 7 and 8 for this parameter.

Fig. 7. ROC analysis of Cleveland dataset

Fig. 8. ROC for Hungarian dataset

Additionally, the area under the ROC curve (AUC) is a commonly used scalar performance metric for binary classification models. A model with an AUC of 1.0 has a perfect prediction, while a model with an AUC of 0.5 has no discrimination ability. An analytical reading of Figures 7 and 8 shows clearly that the ANN model has a better sensitivity and specificity than the other models.

Presumably, the performance metrics (Tables 4 and 5) and ROC curve (Figures 7 and 8) can provide valuable insights into the performance of the different models used in the proposed system. Particularly, the ANN model appears to have the better SS, SP and ACC than the other models. A high sensitivity indicates that the model can correctly identify a large proportion of actual positive cases, while a high specificity indicates that the model is able to correctly identify a large proportion of actual negative cases. Therefore, a model with a high sensitivity and specificity is considered a good classifier.

5 COMPARISON WITH RELATED WORK

 In literature several methods are implemented to classify patients and predict heart disease. The aim of each method is always to achieve the best results to the classify and predict patients with heart disease. Researchers have often compared the performance of different methods on the same dataset to determine which one gives the best results. The choice of method will also depend on the specific features of the dataset and the research question being addressed. Additionally, the interpretability of the model and the ability to use it for clinical decision-making are also important aspects to consider. Therefore, we compared our proposed system with some existing methods that deal with the same database, the results of the comparison are shown in the Table 6.

Table 6. The accuracy obtained in some approach for Cleveland and Hungarian datasets

As mentioned in the table, several methods have been used by researchers to classify and predict heart disease. This comparison shows that our proposed system gave the best performance results compared to other existing methods.

6 CONCLUSION

In this work, we proposed a new ML based diagnostic support system based on five algorithms known for their efficiency in data classification, which are ANN, KNN, NB, SVM and DT algorithms where we obtained an accuracy of 96.67% in the classification of heart diseases utilizing these famous datasets, Cleveland and Hungarian. The aim of this work was to find the hyperparameters of each algorithm that give the best results in terms of performance. The experimental results obtained in the test phase show that all the algorithms give better classification and prediction performance, especially the ANN algorithm which gives good results after being optimized. All this shows that these methods can produce better results if we continue to optimize them. Consequently, the challenge for future research will be to develop methods capable of optimizing these algorithms according to datasets to achieve the maximum possible performance for each algorithm. These results also show that the ANN method has a better ability to classify and predict diseases, which can be explained by the weighting of each input according to its impact on the patient's condition. All this confirms the ability of the ML methods mentioned above to predict heart disease. Finally, we compared the results obtained using this model with some previous methods, which shows that this system has added value in the classification and the prediction of patients with cardiovascular diseases compared with the methods available in the literature. In view of the promising results obtained, we recommend using this model to predict other diseases.

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