



Heart Disease Prediction System Using Convolutional Neural Networks

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Abstract

Now a days, based on different reasons heart diseases are increasing rapidly. If we find out or identify the heart diseases in human beings at an early stage, it is easy to prevent the disease and help the patients. Even though cardiologists and health centers gather relevant data and information every day, but, not applying the knowledge of machine learning algorithms to retrieve valuable of prediction. The main objective of this research is to predict and classify heart diseases by using proposed convolutional neural network classifier. In this classification of evaluation process, feed forward process and back propagation methods will be applied in between the hidden layers. Due to this, the proposed CNN classifier gives best accuracy. By applying this trained classifier has identified the given data, which are either normal or abnormal. So, the entire research has been implemented in Python which produced good results.

Keywords: Deep Learning; Classification; Convolutional Neural Network; Heart Disease

Introduction

Classification is one of the basic approaches in computer science and other research technologies. It can evaluate the bases on many other computer science technology tasks such as object identification, image segmentation and object prediction. The task of categorizing object into one of several predefined classes known as object classification. By using machine learning techniques, data set can be classified, and falls under the category of deep learning. Deep learning is a type of neural network algorithms in which each layer is responsible for extracting one or more features of the data sets.

A neural network is a computational model that is similar to a human brain. It is a collection of nodes known as neurons. These nodes are organized into layers where each neuron in the layer takes some input processes and passes the output to the neuron in the next layer. Different layers may perform different kinds of transformations. Data transfers from the input layer (first layer)

to the output layer (last layer) by traversing various hidden layers. The Convolutional Neural Network (CNN) classifier used for improving the accuracy of image data set for classification [1]. Figure 1 show the structure of an artificial neural network.

The neural network can generate efficient classification rules. Convolution neural network classifier is a multilayer perceptron that is the special design for identification of 2-dimensional data information. Always have more layers: input layer, convolution layer, sample layer, and output layer. Deep learning refers to the shining branch of machine learning that is based on learning levels of representations.

The CNN is one kind of deep neural network. To perform classification task of heart disease dataset, the neural network is trained using convolutions algorithm. The experiment is conducted with heart disease dataset by considering the single and multilayer neural network modes. The proposed algorithm gives detailed analysis of the process of CNN both the forward process and back propagation.

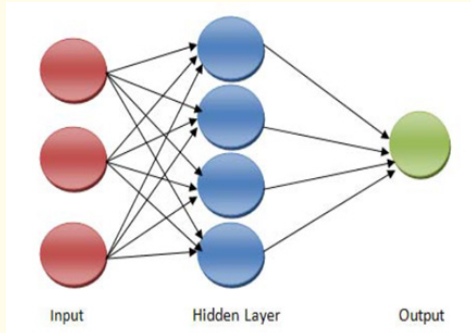


Figure 1: Neural Network.

In this work, building a Convolutional Neural Network based on object classifier which can identify and separate the heart disease data records of normal from that of abnormal. The CNN classifier performed well for performance measures of Mean Absolute Error, Sensitivity, Specificity, F-Measure, False Positive Rate, False Negative Rate, and Overall Error Rate, and compared with existing data mining techniques for better accuracy.

The remainder of this article has been organized as follows: Related work has been briefly introduced in Section 2. The data source of heart disease data sets has been presented in Section 3. Proposed methodology of CNN and its architecture has been elaborated in Section 4. Results of CNN with existing classifiers using heart disease data sets mentioned in Section 5. Finally, concluding remarks.

Related work

In [2], certain fusion methods have been built to diagnose CVDs sideways with its sternness. Machine learning (ML) techniques like artificial neural network, SVM, Logistic Regression, Decision Tree, Random Forest, and AdaBoost have been used to the heart disease dataset to predict diagnose. Random ended tester was executed as of the class inequity in multiclass classification. To increase the performance of classification, a weighted score fusion method was occupied. Initially, the methods were trained. After training, two techniques choice was joint applying a weighted sum rule. Some of these fusion methods have been established from the six ML techniques. The output was likely in the performance parameter. The proposed method has been investigated with dissimilar test training ratios for binary and multiclass classification problems, and for both of them, the fusion methods performed well. The

maximum accuracy for multiclass classification was found as 75%.

In [3], projected k-nearest neighbor founded heart disease prediction method, and author calculated the performance of the proposed method. Further, the output has been calculated for predictive performance of the proposed method. Heart disease data sets are obtained from Kaggle machine learning data repository. The dataset contains of 1025 observations of which 499 or 48.68% is heart disease negative and 526 or 51.32% is heart disease positive. Lastly, the performance of KNN is examined on the test set. The output of performance examination on the experimental outputs on the Kaggle heart disease data repository displays that the accuracy of the KNN is 91.99%.

Correlative examination of numerous diseases has been completed by applying Machine Learning Techniques [5-9]. In [4], a correlative examination was established by applying ML techniques. Together cardiovascular and diabetes disease datasets were applied for classification. Dissimilar ML techniques similar XGBoost, Random Forest, and weighted ensemble methods applied to predict disease. The vital features that fund maximum to the dataset were recognized. Finally, the performance parameters had been exposed to detect the outputs. Ensemble methods expressed marginally well accuracy than further methods. The academics also anticipated to use the method in a real-life situation to verify the risk of the disease happening. Author used ten-fold cross-validation, and its accuracy was marginally less. Two data mining techniques have been applied [10] for the diagnosis of cardiovascular disease. Between them, the Support Vector Machine (SVM) exhibited a robust competence. The accuracy of SVM, and Artificial Neural Network was 74.42% and 73.64%, correspondingly [11-14].

It [15], depends on the test bang values, diagnose a possible solutions. Over assessing the obtainable data gathering, predicted whether the patient has heart disease or diabetes using the model. In this work, projected Naive Bayes technique applied for classifying the data set, and it reached accurate output.

In [16], proposed Machine Learning technique of SVM. This technique predicts the heart disease depends on the assumed features like sex, age, pulse rate etc. Here, SVM reached the greatest accuracy and reliable outputs then compared to the other techniques. And also evaluated perfor-

mance measure with better accuracy, specificity, and sensitivity then compared to the other machine learning techniques.

Data source

The data sets of Cleveland and Statlog are utilized. Cleveland data set comprises 303 patient records of heart disease data [17], and the Statlog data set comprises 270 patient records of heart disease data [18]. A total of 573 patient records are chosen for valuation of the proposed prediction scheme. The data from the patient records will possess 13 input attributes, namely, age, sex, cp, t-rest-bps, chol, Rest-ecg, fbs, thalach, exang, old-peak, solpe, Ca, thal etc. The detail descriptions of these attributes are provided in given below table 1.

Data sets and attributes

The data set was categorized with 3 attributes: Key, Predictable, and Input attributes as given below:

- Key Attribute: Patient Id: Patient’s Recognition Number.
- Predictable Attribute: Diagnosis: Value 1 ≤ 50 % (no heart disease); Value 0 ≥ 50 % (has heart disease).
- Input Aattributes (Heart Disease): Table 1.

F8	Thalach	Maximum heart rate attained	Continuous value
F9	Chol	Exercise make angina	0 = no; 1 = yes
F10	Oldpeak	ST despair make by exercise virtual to rest	Continuous value
F11	Slop	Slope of the crest exercise ST fragment	1 = un sloping 2 = fat 3 = down sloping
F12	Ca	Number of main vessels tinted by fluoroscopy	0 - 3 value
F13	Thal	Defect type	3 = normal 6 = fixed 7 = reversible defect

Table 1: Input Attributes for Prediction System.

Variable Name	Attribute	Descriptions	Values
F1	Age	Age in years	Continuous
F2	Sex	Male or Female	1 = male; 0 = female
F3	Cp	Chest Pain Type	1 = typical type I 2 = typical type angina 3 = non-angina pain 4 = asymptomatic
F4	T rest bps	Resting blood pressure	Continuous value in mm hg.
F5	Chol	Serum cholesterol	Continuous value in mg/dl.
F6	Restecg	Resting electrographic results	0 = normal 1 = containing ST_T sign odd 2 = left ventricular hypertrophy
F7	Fbs	Fasting blood sugar	1 ≥ 120 mg/dl. 0 ≤ 120 mg/dl.

Here, a key attribute of a patient record may be patient identification number (Patient ID), which is specified for every patient and can be easily distinguishable.

The above 13 attributes of data of a patient are said to be as input attributes which may have some personal information. Along with these attributes there is one more attribute for every record and can be helpful to identify it even among a number of similar records also termed as key attributes. This database contains 76 attributes, but all published experiments refer to using a subset of 13 of them.

In particular, the Cleveland, Statlog databases are used by Machine Learning researchers. The “goal” field refers to the presence of heart disease in the patient. It is integer valued from 0 (no presence) to 4. Experiments with this database have concentrated on simply attempting to distinguish presence (values 1, 2, 3, 4) from absence (value 0). The names and social security numbers of the patients were recently removed from the database, replaced with dummy values.

Proposed Methodology

Convolutional Neural Network (CNN) has input layers, output layers and hidden layers. The hidden layers consist of con-

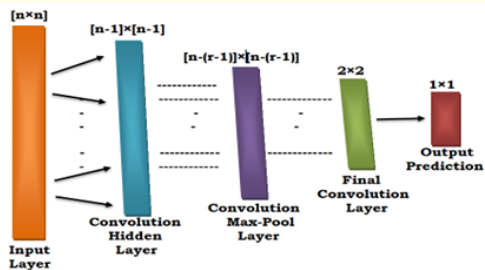


Figure 2: Proposed CNN Classifier Architecture for Heart Disease Data Sets.

convolutional layer, flattened layer and a fully connected layer. The Figure 2 shows the architecture of the proposed CNN.

Convolutional Neural Networks (CNN) is one kind of deep neural network. It can study concurrently. The proposed detailed analysis of the process of Convolutional Neural Networks algorithm both the forward process and back propagation. Then applied the particular Convolutional Neural Network to implement the typical heart disease dataset problems. The figure 3 shows general architecture of the CNN.

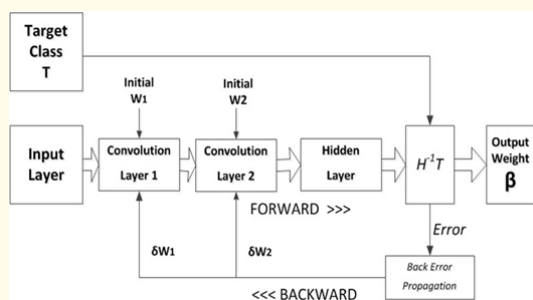


Figure 3: General Convolutional Neural Network Architecture.

In addition, by measuring the actual time of forward and backward evaluation, analyse the maximal speed up and parallel efficiency theoretically.

Role of convolutional neural networks

In general, the structure of CNN includes two layers, one is feature extraction layer, and the input of every neuron is connected to the local except fields of the previous layer, and extracts the local feature. Once the local features are extracted, the position

relationship between it and other features will be evaluated. The other layer is feature map layer; all calculation layers of the network is mapped a multitude in feature map. Every feature is a level, and the weights of the neurons in the level are close.

The construction of feature applies the sigmoid function as stimulation function of the convolution network, which makes the feature shift-in-variance. The numeral of free parameters of the network is reduced. All convolution layers in CNN by calculating the layer which is applied to calculate the local mean and the second extract. This particular two feature extraction constructions reduce the record's matrix level size. Multi-dimensional input vector of heart disease data sets can exit the network, which avoids the quality of data reconstruction in feature extraction and classification procedure.

Feature selection algorithm

Several feature ranking and feature selection algorithms have been projected in the machine learning study. The purpose of these algorithms identifying and locate the unfit or unnecessary of from the vector. For the implementation of this research work, it applied feature ranking and selection modes by two initial steps of overall architecture: subset creation and subset calculation for the ranking of all features in every data set. Filter mode was applied to measure all subsets.

Information gain

The proposed feature choice both class membership and the presence/absence of a specific period are observed as random variables and one evaluates how more information around the class membership is increased by finding the presence/absence statistics as applied in decision tree induction. So, if the class membership is taken as a random variable C with two values, positive and negative, and a word is similarly observed as a random variable T with two values, present and absent, it is by applying the information practical statement of ordinary information specified as the following equation:

$$IG(T) = H(C) H(C/T) = \sum cP(C=c, T=\tau) \ln [cP(C=c, T=\tau)/P(C=c).P(T=\tau)] \dots\dots\dots(1)$$

Here, τ ranges over {present, absent} and c ranges over { $c+$, $c-$ }. As pointed out above, this is the measure of information about C (the class label) that increases by finding T (presence or absence of a word).

Back propagation algorithm

Convolutional Neural Networks technique is a multilayer perceptron, which is the specific system for recognition of two-dimensional data. It has many layers: input layer, convolution layer, sample layer and output layer. The CNN has two primary procedures: convolution and sampling. Convolution procedure applies a predictable filter F_x , re-convolution of the input data (the initial phase is the input data, the input of the later convolution is the feature data of every layer, namely, Feature Compose). Then, add a bias b_x , get convolution b_x layer C_x . A selecting procedure: n points of each neighborhood by pooling steps, get a point, and then by scalar weighting W_x+1 weighted, add bias b_x+1 , and then by an activity function, make a narrow n times feature S_x+1 .

The central engineering of CNN is the local tract, jointing of weights, sub selecting by time or space, and hence the training parameters extract feature and reduce the size. The benefit of CNN technique is prevention of explicit feature extraction, and learning from the training data. The neuron weights on the surface of the feature composing, thus, the network can see parallels, and reduce the multilevel of the network, Adapting sub sampling structure by the time or space, can attain a few degrees of robustness, scale and modification replacement. Input information and network topology can be a very good match, It has specific benefits in speech identification and data sets processing.

$$O_{x,y}^{1,k} = \tanh \sum_{t=0}^{f-1} \sum_{r=0}^{k_h} \sum_{c=0}^{k_w} W_{(r,c)}^{(k,t)} O_{(x+r,x+c)}^{(l-1,x)} + Bias^{(l,k)} \quad \text{-----}(2)$$

Among them, f is the numeral of convolution cores in a feature pattern, output of neuron of row x, column y in the lth sub sample layer and kth feature pattern:

$$O_{x,y}^{1,k} = \tanh (W^k \sum_{r=0}^{s_h} \sum_{c=0}^{s_w} O_{(x \times s_h + r, y \times s_w + c)}^{l-1,t} + Bias^{(l,k)}) \quad \text{-----}(3)$$

The output of the jth neuron in lth hides layer H:

$$O_{(i,j)} = \tanh (W^k \sum_{k=0}^{s-1} \sum_{x=0}^s \sum_{y=0}^{s_w} W_{(x,y)}^{(j,k)} O_{(x,y)}^{(l-1,k)} + Bias^{(l,k)}) \quad \text{-----}(4)$$

Among them, s is the number of feature patterns in sample layer output of the ith neuron lth output layer F:

$$O_{(i,j)} = \tanh (\sum_{j=0}^H O_{(l-1,j)} W_{(i,j)}^l + Bias^{(l,i)}) \quad \text{-----}(5)$$

Modified back-propagation

A speed matrix depends on the technique to evaluate the output from a Neural Network (NN). Especially, it is an excellent mode of acquiring the notation used in back-propagation. Back-propagation is a Neural Network learning algorithm. The neural networks field was addressed by psychologists and neuro-biologists who wanted to create and test evaluation analogy of neurons. An NN is a set of input/output units in which every attachment has a weight joined by it. During the learning stage, the network learns with changing the weights so far capable to predict the exact class label of the input tuples.

Output deviation of the kth neuron in output layer O:

$$d(O_k^0) = y^k - \tau^k \quad \text{-----}(6)$$

Input deviation of the kth neuron in output layer:

$$d(I_k^0) = (y^k - \tau^k) \varphi(v_k) d(O_k^0) \quad \text{-----}(7)$$

Weight and bias variation of kth neuron in output O:

$$\Delta W_{k,x}^0 = d(I_k^0) y_{k,x} \quad \text{-----}(8)$$

$$\Delta Bias_k^0 = d(I_k^0) \quad \text{-----}(9)$$

Output bias of kth neuron in hide layer H:

$$d(O_k^H) = \sum_{i=0}^{i++} d(I_k^0) W_{i,k} \quad \text{-----}(10)$$

Input bias of kth neuron in hide layer H:

$$d(I_k^H) = \varphi(v_k) d(O_k^H) \quad \text{-----}(11)$$

$$\Delta W_{m,x,y}^{H,k} = d(I_k^H) y_{x,y}^m \quad \text{-----}(12)$$

$$\Delta Bias_k^H = d(I_k^H) \quad \text{-----}(13)$$

Output bias of row x, column y in mth feature pattern, sub sample layer S

$$d(O_{x,y}^{S,m}) = \sum_{k=0}^{i++} d(I_{m,x,y}^H) W_{m,x,y}^{H,k} \quad \text{-----}(14)$$

Input bias of row x, column y, in mth feature pattern, sub sample layer S:

$$d(I_{x,y}^{S,m}) = \varphi(v_k) d(O_{x,y}^{S,m}) \quad \text{-----}(15)$$

Weight and bias variation of row x, column y, in mth feature of sub sample

$$\Delta W_{x,y}^{S,m} = \sum_{x=0}^{fh} \sum_{y=0}^{sw} d(I_{x,y}^{S,m}) O_{x,y}^{C,m} \quad \text{-----}(16)$$

Among them, C represents convolution layer:

$$\Delta \text{Bias}^{S,m} = \sum_{x=0}^{fh} \sum_{y=0}^{fw} d(O_{x,y}^{S,m}) \dots\dots\dots (17)$$

Output bias of row x, column y in kth feature pattern, convolution layer C:

$$d(O_{x,y}^{C,k}) = d(I_{x,y}^{S,k}) W^k \dots\dots\dots (18)$$

Input bias of row x, column y in kth feature pattern, convolution layer C:

$$d(I_{x,y}^{C,k}) = \varphi(v_k) d(O_{x,y}^{C,k}) \dots\dots\dots (19)$$

Weight variation of row r, column c in mth convolution core, corresponding to kth feature pattern in lth layer, convolution C:

$$\Delta W_{r,c}^{k,m} = \sum_{x=0}^{fh} \sum_{y=0}^{fw} d(I_{x,y}^{C,k}) O_{x+r,y+c}^{l-1,m} \dots\dots\dots (20)$$

Total bias variation of the convolution core:

$$\Delta \text{Bias}^{C,k} = \sum_{x=0}^{fh} \sum_{y=0}^{fw} d(I_{x,y}^{C,k}) \dots\dots\dots (21)$$

CNN algorithm process steps

- Step 1: Take the arff data set.
- Step 2: Feature selection using information gain and ranking.
- Step 3: Classification algorithm.
- Step 4: Every Feature calculates f_x value of input layer.
- Step 5: Bias class of every feature calculation.
- Step 6: After giving the feature map, it goes to forward pass input layer.
- Step 7: Evaluate the convolution cores in a feature pattern.
- Step 8: Gives sub sample layer and feature value.
- Step 9: Back-propagation input deviation of the kth neuron in output layer.
- Step 10: Lastly, produce the chosen feature and classification results of (1*1) matrix output (i.e. Normal or Abnormal of heart disease data set records).

Results

The dataset of all records having a set of 13 similar attributes. During the collection of a few missing values are found because of manual write down by doctors. By means of data missing Weka 3.6.6 tool the missing values are replaced using mean, mode method by filter option available in the tool. A confusion matrix is obtained to compute the accuracy of classification. In this work, confusion matrix shows how many instances have been assigned to every class. Table 2 shows the confusion matrix for evaluation of performance measures.

	A (has heart disease)	B (no heart disease)
A (has heart disease)	True Positive (TP)	False Negative (FN)
B (no heart disease)	False Positive (FP)	True Negative (TN)

Table 2: Class A = YES (has heart disease);

The data set made of 573 records and it has been divided into on 25 classes, where each class consists of 23 records. Training and Testing sets are divided into equal amount to predict the exactness of the system. To measured certainty of 13 attributes are chosen to classify the system by a symbolic learning approach by interval method, i.e., $(\mu\sigma, \mu+\sigma)$ as indicate in the Table 1.

In this work trained the classifiers to classify the heart diseases are either “has heart disease” or “no heart disease”. The general and particular of confusion matrixes are two classes (i.e. normal and abnormal) of five classifiers. Table 3 shows for the evaluation of True Positive, True Negative, False Positive, and False Negative values in confusion matrix.

	A	B
A	257	05
B	08	271

Table 3: Confusion matrix accèss from CNN classifier.

The CNN algorithm compared with the classifiers such as K-NN, Naïve Bayes, SVM, and ANN, and evaluated their individual Confusion Matrix respectively.

Accuracy, mean value error

The classifiers of Accuracy and Mean Value Error performance measures are evaluated by using below formulae and results are shown in the Table 4, and the comparison chart is specified in the figure 4.

$$\text{Accuracy} = \frac{TP+TN}{TP+TN+FP+FN}, \text{ Mean Absolute Error} = \frac{1}{n} \sum |y_i - x_i|$$

Where: TP=True Positive, TN=True Negative, FP=False Positive, FN=False Negative

2nd equation MAE: is actual value of ith observation, calculated value of ith observation; n is total number of observations.

Classifiers	Accuracy	Mean Absolute Error
K-NN	54.42	0.046
Naive Bayes	75.68	0.003
SVM	79.58	0.009
ANN	87.87	0.0028
CNN	97.43	0.0014

Table 4: Accuracy, and Mean Value Error comparative of five classifiers.

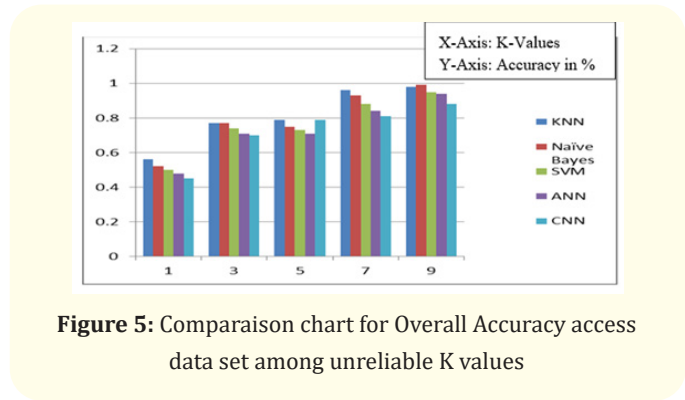


Figure 5: Comparison chart for Overall Accuracy access data set among unreliable K values

Postoperative Pain Control In Femto-Lasik With The Use Of Therapeutic Contact Lenses Soaked In Ketorolac Trometamol 0.45%

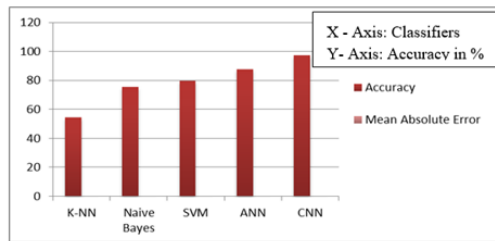


Figure 4: Accuracy and Mean Absolute Error Comparison Chart.

Similarly, the Overall accuracy performance measure evaluated for data set among unreliable K values shows in the table 5, and the comparison chart is specified in the figure 5.

K	K-NN	Naive Bayes	SVM	ANN	CNN
1	0.56	0.77	0.79	0.96	0.98
3	0.52	0.77	0.75	.93	0.99
5	0.50	0.74	0.73	0.88	0.95
7	0.48	0.71	0.71	0.84	0.94
9	0.45	0.70	0.79	0.81	0.88

Table 5: Overall Accuracy accèss for data set among unreliable K values.

Sensitivity, Specificity, and F-Measure

The classifiers of Sensitivity, Specificity, and F-Measure performance measures are evaluated by using below formulae and results are shown in the Table 6, and their comparison chart is specified in the Figure 6.

$$\text{Sensitivity} = \frac{TP}{TP+FN}, \text{ Specificity} = \frac{TN}{TN+FP}$$

$$\text{F-Measure} = \frac{2 \times \text{Sensitivity} \times \text{Specificity}}{\text{Sensitivity} + \text{Specificity}}$$

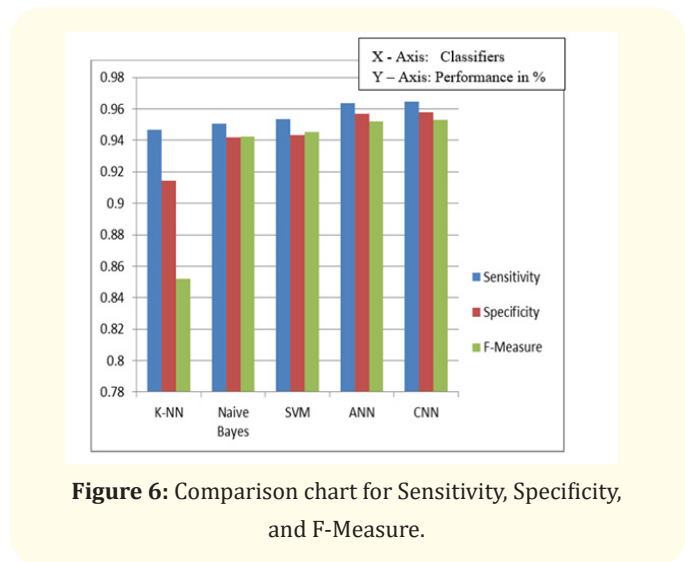


Figure 6: Comparison chart for Sensitivity, Specificity, and F-Measure.

False positive rate, false negative rate and overall error rate

The classifiers of False Positive Rate, False Negative Rate and Overall Error Rate performance measures are evaluated by using below formulae and results are shown in the table 7, and their comparison chart is specified in the figure 7.

$$\text{FPR} = \frac{\text{FP}}{\text{FP}+\text{TN}}, \text{FNR} = \frac{\text{FN}}{\text{FN}+\text{TP}}, \text{Overall Error Rate} = \frac{\text{FP}+\text{FN}}{\text{FP}+\text{FN}+\text{TP}+\text{TN}}$$

Classifiers	FPR	FNR	Overall Error Rate
K-NN	0.0631	0.0596	0.0678
Naive Bayes	0.0568	0.0382	0.0453
SVM	0.0487	0.0356	0.0417
ANN	0.0321	0.0267	0.0345
CNN	0.0398	0.0298	0.0343

Table 7: Comparison Table for FPR, FNR, and Overall Error Rate.

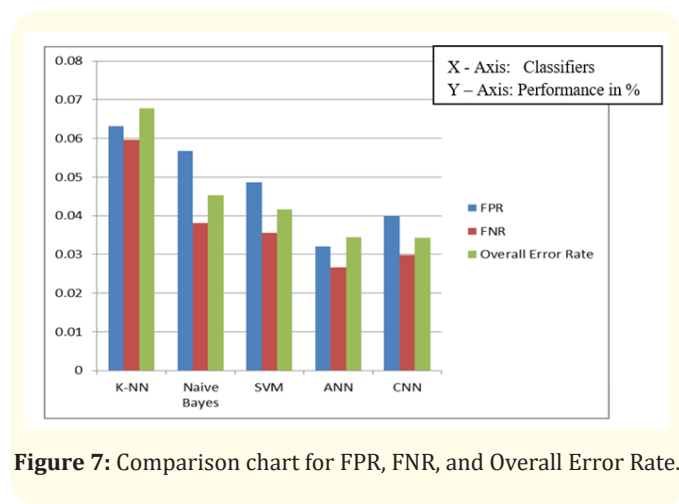


Figure 7: Comparison chart for FPR, FNR, and Overall Error Rate.

The evaluated measurement of the performance based on 13 attributes is accurate in natural history, and the accuracy of Convolution neural network using heart disease data set is reached up to 99%. CNN is improved performance than other classifiers, while K-NN shows weak performance than other classifiers.

Conclusion

In this proposed system, neural network classification methodology is made efficiently. To perform classification task of medical data, the neural network is trained using Convolution technique. The experimentation is shown with heart disease dataset by considering the single and multilayer neural network methods. Convolution neural network technique is a multilayer perceptron that is the su-

perior design for reorganization of two-dimensional heart disease data information. In this so many layers are observed, i.e.: input layer, convolution layer, sample layer and output layer. In addition, in deep neural network architecture the convolution layer and sample layer can have multiple layers in between. The overall objective of this work is to predict the heart diseases with more accuracy. This work can be enhanced by increasing the more number of records with attributes to provide better accuracy for the existing system.

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Data Availability Statement

- Cleve land data base: <http://archive.ics.uci.edu/ml/datasets/Heart+disease>
- Stalogdatabase: [https://archive.ics.uci.edu/ml/datasets/statlog+\(heart\)](https://archive.ics.uci.edu/ml/datasets/statlog+(heart))

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Conflict of Interest

No conflict of interest.

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