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School of Computer Science and Engineering

J Component report

Programme : B.Tech

Course Title : Foundations of Data Analytics

Course Code : CSE3505

Slot : F2

Title: Machine Learning Based Diabetes Classification and Prediction for Healthcare Applications

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Date:

DECEMBER 2022

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DECLARATION

We hereby declare that the project report entitled **"Machine Learning Based Diabetes Classification and Prediction for Healthcare Applications"** undertaken by me under the supervision of **Dr. Trilok Nath Pandey**, School of Computer Science and Engineering, Vellore Institute of Technology, Chennai 600127 in partial fulfilment of the requirements for the award of the degree of **Bachelor of Technology – Computer Science and Engineering** is a record of bonafide work carried out by us**.** We further declare that the work reported in this report has not been submitted and will not be submitted, either in part or in full, for the award of any other degree or diploma of this institute or of any other institute or university.

Signature

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CERTIFICATE

This project report for the course **"Machine Learning Based Diabetes Classification and Prediction for Healthcare Applications"** is prepared and submitted by **Mayank Gupta (Register No: 20BCE1538) Fidal Mathew (Register No: 20BCE1538) Abhishek N.N. (Register No: 20BCE1538) Ayush Kapri (Register No: 20BCE1538)**. It has been found satisfactory in terms of scope, quality and presentation as partial fulfilment of the requirements for the award of the degree of **Bachelor of Technology – Computer Science and Engineering** in Vellore Institute of Technology, Chennai, India.

Examined by:

Examiner I **Examiner II**

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CONTENTS

ABSTRACT

One of the most important health issues in both industrialized and developing nations is diabetes mellitus. Consequently, the According to the International Diabetes Federation, 425 million people worldwide have diabetes. Within 20 years, this number is projected to increase to 380 million. Due to its significance, a classifier design for the early diagnosis of diabetes that is both affordable and effective is now necessary. Testing data mining methods to determine their predictive accuracy in the classification of diabetes data has become a standard at the UCI machine learning lab using the Pima Indian diabetic database.

The machine learning technique is focused on categorizing the diabetic illness from a large medical dataset into type 1 and type 2. The goal of this project is to create a model that can predict a patient's chance of developing diabetes with the highest degree of accuracy. As a result, this experiment uses Decision Tree, Naive Bayes, Random Forest, Kernel SVM and Logistic Regression five machine learning classification methods, to identify diabetes at an early stage. Confusion Matrix, Precision, Accuracy, F-Measure, and Recall are just a few of the metrics used to assess how well the three algorithms perform. Correctly and wrongly labelled examples are used to gauge accuracy. According to the results, Logistic Regression surpasses other algorithms with a highest accuracy of 79.6%. These findings are properly and methodically validated using Receiver Operating Characteristic (ROC) curves.

Keywords— Diabetes, Logistic Regression, Kernel SVM, Naïve Bayes, Decision Tree, Random Forest Accuracy

Introduction

In the medical industry, classification algorithms are frequently used to categorise data into different groups in accordance with specified constraints as opposed to using a single classifier. Diabetes is a condition that impairs the body's ability to produce the hormone insulin, which causes improper carbohydrate metabolism and raises blood glucose levels. A person with diabetes typically experiences elevated blood sugar. Increased hunger, increased thirst, and frequent urination are a few signs and symptoms of high blood sugar. Diabetes has a lot of side effects if it is not addressed. Diabetes-related ketoacidosis and nonketotic hyperosmolar be managed are two serious consequences. Diabetes is influenced by a number of variables, including height, weight, hereditary factors, and insulin, but the main component that is taken into considerationis sugar Machine Learning concerntration among all factors. The early identification is the only remedy to stay away from the complications. Diabetes is examined as a vital serious health matter during which the measure of sugar substance cannot parentheses, following the example. Some components, such as multi-leveled equations, graphics, and tables are not prescribed, although the various table text styles are provided. The formatter will need to create these components, incorporating the applicable criteria that follow.

Literature Survey

The study by Deepti Sisodia and Dilip Singh Sisodia, titled "Prediction of Diabetes using Classification Algorithms," had the greatest accuracy, 76.30%. In this experiment, they employed three machine learning classification algorithms—Decision Tree, SVM, and Naive Bayes—to identify diabetes at an early stage. Receiver Operating Characteristic (ROC) curves are used correctly and methodically to validate these findings. To predict blood sugar levels, an IEEE paper tested several machine learning techniques separately, including Support Vector Machine (SVM), Naive Bayes (NB), and Random Forest (RF). All of these techniques were effective, with a prediction accuracy of up to 75%. The suggested system has been compared, including feature extraction, support vector machine, naive Bayes, and random forest methods. The effectiveness of the three algorithms is then assessed using a variety of metrics, including accuracy, precision, F-measure, and recall.

Dataset Used

A. PIDD-Pima Indians Diabetes Dataset

The proposed methodology is evaluated on Diabetes Dataset namely (PIDD), which is taken from UCI Repository. This dataset comprises of medical detail of 768 instances which are female patients. The dataset also comprises numeric-valued 8 attributes where value of one class ’0’ treated as tested negative for diabetes and value of another class ’1’ is treated as tested positive for diabetes. Dataset description is defined by Table-4 and the Table-5 represents Attributes descriptions.

B. Correlation Matrix

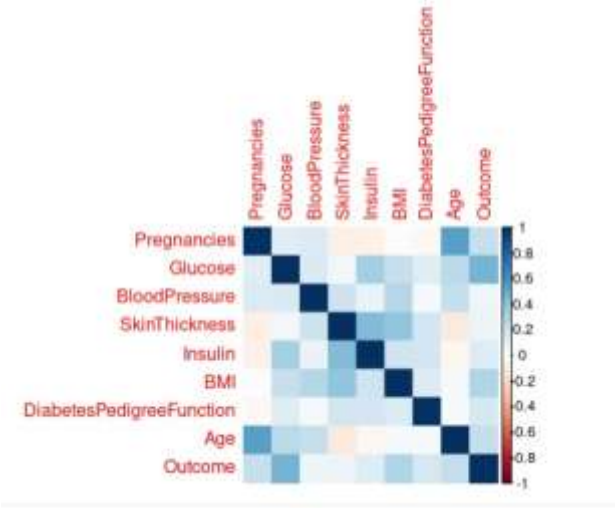

FIGURE 1: CORRELATION MATRIX

Data cleaning is **the process of correcting or eliminating data that is erroneous, corrupted, poorly formatted, duplicate, or incomplete within a dataset**. We have removed some unecessary features of our data by careful observations and experimentation. We implemented

correlation heatmaps. They help us to understand which variables are related to each other and the strength of this relationship. The PIC-2 depicts scatterplot matrix of the individual features of the PIDD dataset.

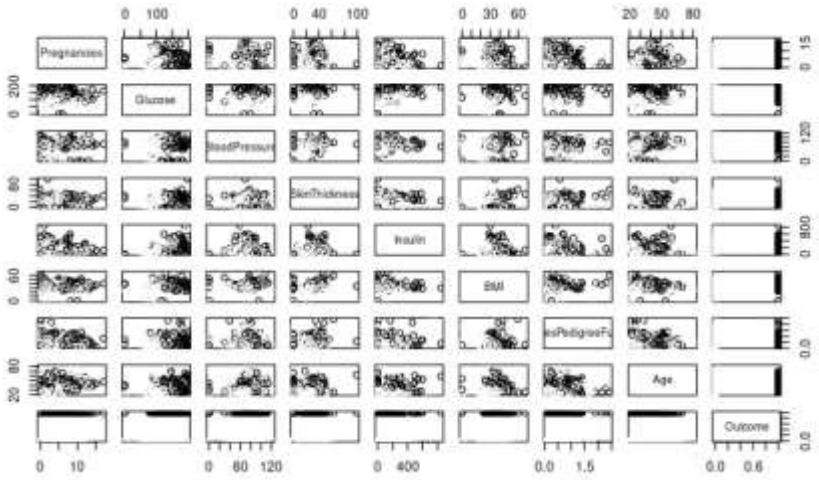

FIGURE 2: SCATTERPLOT MATRIX

By observation, we notice **Age** and **Pregnancies** features are very much correlated. To achieve a good machine learning model, we removed **Pregnancies** feature as **Age was more correlated to Outcome**. We also removed **Skin Thickness** from all the models as it has very slight correlation with the outcome. Having features which do not contribute to the final result makes the model more complex and inefficient.

Methodology used

Learning (determining) good values for all of the weights and the bias from labelled examples is all that is required to train a model. In this experiment, we will using different machine learning algorithms to predict whether the person is subjected to type-1 or type-2 diabetes.

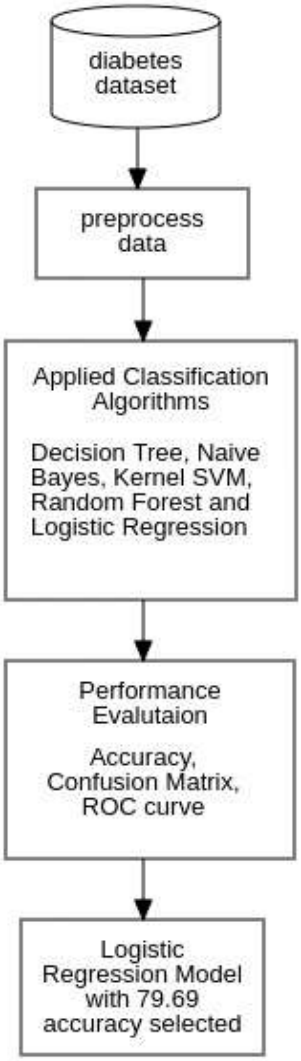

FIGURE 3: SYSTEM ARCHITECTURE

A. Logistic Regression

The Logistic regression helps to classify the concern person will get diabetes or not. Since we are using the logistic regression we have to mention that, family binomial. We are using all the attributes we have in the dataset. The Confusion Matrix of Logistic Regression is as follows:

	A	*B*
Test Positive	116	9
Test Negative	30	37

TABLE 1: CONFUSION MATRIX FOR LOGISTIC REGRESSION

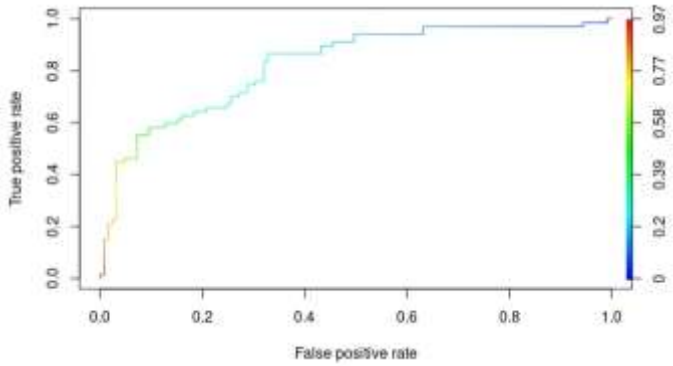

FIGURE 4: ROC CURVE FOR LOGISTIC REGRSSION

B. Naive Bayes Classifier

Naive Bayes is a classification technique based on the idea that all features are independent and unconnected to one another. It specifies that the status of one feature in a class does not impact the status of another. Because it is based on conditional probability, it is regarded as a powerful method used for classification. It works well with data that has imbalancing issues and missing values. The Bayes Theorem is used by Naive Bayes, a machine learning classifier. The posterior probability P(C|X) can be determined using the Bayes theorem from P(C), P(X), and P(X|C). Therefore, P(X|C) = P(X|C) P(C)/P (X).

Where P(C|X) is the posterior probability of the target class.

P(X|C) denotes the probability of the predictor class.

P(C) is the probability that class C is true.

P(X) denotes the predictor's prior probability.

The Naive Bayes algorithm's performance was tested using confusion matrix:

	A	*B*
Test Positive	117	8
Test Negative	33	34

TABLE 2: CONFUSION MATRIX FOR NAIVE BAYES

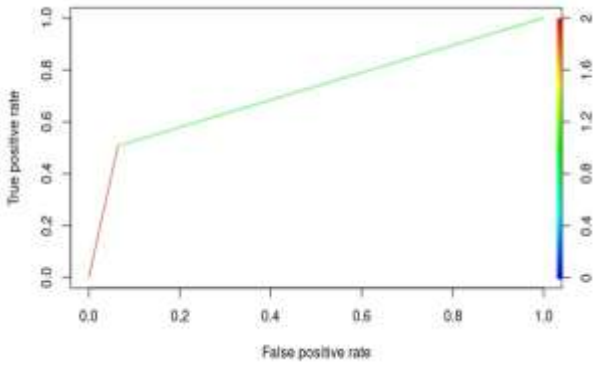

FIGURE 5: ROC CURVE FOR NAIVE BAYES

C. Decision Tree Classifier

A supervised machine learning algorithm used to tackle categorization problems is Decision Tree. The prediction of target class using decision rule drawn from prior data is one of the benefits of employing Decision Tree in this research endeavour. It predicts and classifies using nodes and internodes. Root nodes categorise instances based on various characteristics. The root node may have two or more branches, whereas the leaf node represents classification. At each level, the Decision Tree selects a node based on the maximum information gain among all qualities. The results of his evaluation of the Decision Tree technique using the Confusion Matrix are as follows:

	A	***B***
Test Positive	105	20
Test Negative	28	39

TABLE 3: CONFUSION MATRIX FOR DECISION TREE

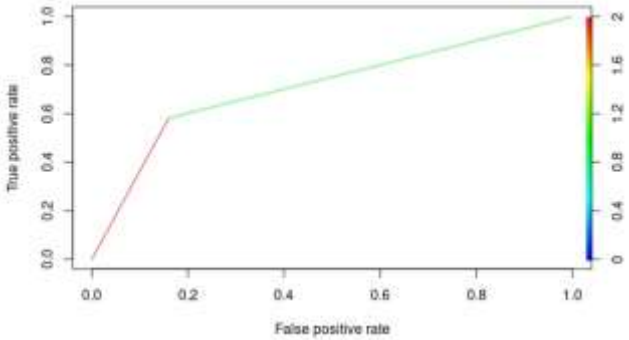

FIGURE 6: ROC CURVE FOR DECISION TREE

D. Kernel SVM

SVM is a supervised machine learning model that is commonly used in classification. A support vector machine's goal, given a two-class training sample, is to determine the optimal highest-margin separation hyperplane between the two classes. For greater generalisation, the hyperplane should not be located closer to data points from the other class. A hyperplane that is far from the data points in each category should be chosen. The support vectors are the

spots closest to the classifier's margin. The estimated performance of the SVM algorithm for diabetes prediction using the Confusion Matrix is as follows:

	A	***B***
Test Positive	114	11
Test Negative	30	37

TABLE 4: CONFUSION MATRIX FOR DECISION TREE

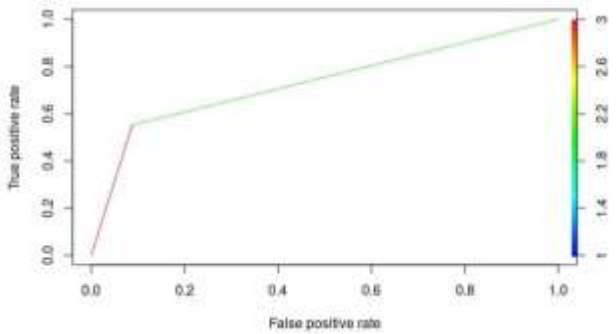

FIGURE 7: ROC CURVE FOR DECISION TREE

E. Random Forest

The method of supervised learning includes Random Forest. It can be used to solve ML problems involving both classification and regression. It is based on the concept of ensemble learning, which is a method of combining several classifiers to address tough difficulties and improve model performance. Following is an evaluation of the SVM algorithm's performance in predicting diabetes using the Confusion Matrix:

	A	***B***
Test Positive	111	14
Test Negative	27	40

TABLE 5: CONFUSION MATRIX FOR RANDOM FOREST

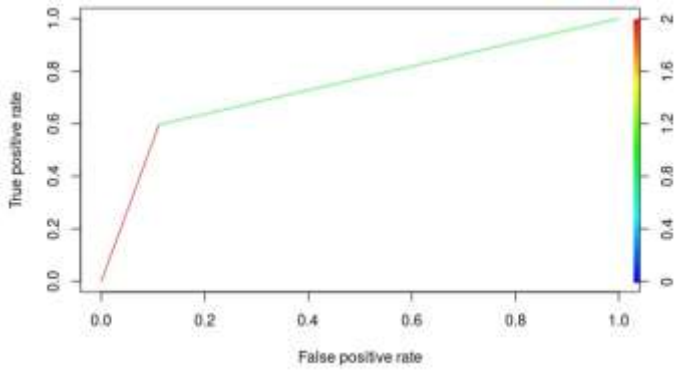

FIGURE 8: ROC CURVE FOR RANDOM FOREST

Accuracy Measures

In this study, the algorithms Naive Bayes, SVM, and Decision Tree are employed. Tenfold internal cross-validation is used throughout experiments. For the classification of this study, the metrics Accuracy, F-Measure, Recall, Precision, and ROC (Receiver Operating Curve) are employed. Below are the accuracy measures defined in TABLE 7:

I. Accuracy

The accuracy of a machine learning classification method can be used to determine how frequently it properly identifies a data point. Accuracy is defined as the fraction of correctly predicted data points among all data points.

II. Precision

Precision is defined as the proportion of correctly categorised positive samples (True Positive) to the total number of positively classified samples (either correctly or incorrectly).

III. F-Measure

The F-measure is calculated using the harmonic mean of precision and recall, with each given the same weight. It enables the evaluation of a model's precision and recall with a single score, which is valuable for discussing model performance and comparing models.

IV. Recall

The recall is calculated by dividing the proportion of Positive samples that were correctly recognised as Positive by the total number of Positive samples. The recall metric measures how successfully the model can detect positive samples. The greater the number of positive samples detected, the greater the recall.

V. ROC

A receiver operating characteristic curve, also known as a ROC curve, is a graphical representation of how the diagnostic capacity of a binary classifier system changes as the discrimination threshold is changed.

SN	***Measures***	***Formula***
1	Accuracy	A=(TP+TN) / (Total no of samples)
2	Precision	P = TP / (TP+ FP)
3	Recall	R =TP / (TP+FN)
4	FMeasure	F=2*(P*R) / (P+R)
5	ROC	Used to compare the usefulness of tests.

TABLE 6: ACCURACY MEASURES

SN	*Measures*	*OLD Accuracy*	*Accuracy with Feature Selection*
1	Logistic Regression	78.65	79.69
2	Naïve Bayes	77.08	78.65
3	Decision Tree	74.48	75.00
4	Kernel SVM	77.60	78.65
5	Random Forest	77.60	78.65

TABLE 7: COMPARATIVE ACCURACY GAIN USING FEATURE SELECTION

SN	**Performance Evaluation**				
	Machine Learning Models	***Accuracy***	***F Measure***	***Recall***	***Precision***
1	Logistic Regression	79.69	85.60	79.45	92.80
2	Naïve Bayes	78.65	85.09	78.00	93.60
3	Decision Tree	75.00	81.39	78.94	84.00
4	Kernel SVM	78.65	84.75	78.72	90.98
5	Random Forest	78.65	83.65	80.43	88.80

TABLE 8: COMPARATIVE PERFORMANCE OF CLASSIFICATION

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