Implications of Meta Classifiers for Onset Diabetes Prediction

Md. Ashaf Uddaula, Md. Al - Amin Hossain, Md. Khalid Hossen, Ahmed Al Marouf

Abstract: In the data mining area, the prophecy of human diseases initiates a research zone for researchers by applying various machine learning algorithms with various patterns. As a modern community disease, diabetes is becoming one of the fastest-progressive human diseases in the world because of eating heavily sugared foods and lack of proper diet knowledge. In this era, most of the middle age people have confusion about the presence of diabetes in their bodies. That’s why we choose to do research on diabetes. In this paper, we scrutinized the classification performance of six Meta Classifiers named as Multiclass Classifier Updatable, Attribute Selected Classifier, Ada Boost M1, Logit Boost, Bagging, and Filtered Classifier for forecasting diabetes through cross-validation and percentage split techniques using in WEKA whereas as a diabetes dataset we used Pima Indians Database. And finally, according to win-rate from the Win-Draw-Loss table, the highest performance comes from Multiclass Classifier Updatable which has an 80% win-rate. On the other hand, in the measurement of highest individual accuracy, 81.9923% comes from both Attribute Selected Classifier and Filtered Classifier. According to the measurement of the highest average performance, 66% Split as a percentage split technique and Attribute Selected Classifier show the highest performance.

Keywords: Data Mining, Meta Classifier, WEKA, Percentage Split Technique, Diabetes, Win Rate, Multiclass Classifier Updatable, Attribute Selected Classifier.

I. INTRODUCTION

Diabetes is a common chronic disease for our modern society. People getting used to diabetes day by day because of its availability in today's world. It can infect by genetically or, also by taking high sugared food. Nowadays it can be looked that, children are affected by diabetes increasingly due to the presence of diabetes to their parents. We know that we can't prevent diabetes but, we can control diabetes by controlling the sugar level in our blood circulation. But, most people don't know that they are affected by diabetes or, not yet. Even many people don't want to test the presence of diabetes because they don't want to lead their lives with a tight food schedule if they are affected. But, we need to checkup our body and maintain a disciplined food list in our life. Here, data mining helps people to check the presence of diabetes after analyzing some valid data.

Data mining is a result of some periodic processes like data purifying, integration of data, selection of data, transformation of data, mining the data, evaluate pattern and lastly representation of knowledge that are helped to invent ultimate patterns, relationships, insights of enterprises measuring and managing where we are now and predicting where we will be in the tomorrow from huge data sets.

![Fig. 1. A View of Data Mining [12]](image)

As a big asset for diabetes researchers, Data mining has performed a spontaneous role in diabetes research and also would be a beneficial way for our medical science. Actually, it builds a relationship with our medical healthcare resources. Data mining can identify clandestine knowledge from a large volume of diabetes-related data. Our belief is that data mining not only can significantly help in diabetes research but also it can ensure better quality health care for those patients who are affected by diabetes.

In diabetes research, data mining techniques are applied in some systematic ways. Machine learning algorithms used data mining techniques to build up the model and pattern to find out the accuracy rate of classification, prediction, relationship, and many others not only in the diabetes sector but also in diagnosis-related research sectors. Machine learning is an advanced study of mathematically proved algorithms and scientifically accepted statistical models that help computer-based hardware or, software systems to execute an appointed duty instead of using detail instructions, relying on models and hypothesis instead. It can be announced as a part of artificial intelligence.
Implications of Meta Classifiers for Onset Diabetes Prediction

And machine learning algorithms set up a mathematical pattern based on specimen datum which is named as "training data", according to make diagnostic or judgment instead of being in detail programmed to execute the task.

In Our paper, we scrutinized the comparative view of six Meta classifier algorithms named as Multiclass Classifier Updatable, Attribute Selected Classifier, Ada Boost M1, Logit Boost, Bagging, and Filtered Classifier for forecasting diabetes through cross-validation and splitting techniques using in WEKA whereas as a diabetes dataset we used the most renowned Pima Indians Diabetes Database. We also estimated our comparative views with so many mathematically calculated tables especially the win-draw-loss table, many curves like ROC curves and Precision-Recall Curves and also many statistical graphs.

We found so many results from different comparative views. On the basis of the Win-Draw-Loss table, Multiclass Classifier Updatable has performed the highest performance with an 80% win-rate. On another comparative view, Attribute Selected Classifier and Filtered Classifier have given the highest accuracy 81.9923% individually. Then, if we talk about the averagely, then the percentage split technique named 66% Split and Attribute Selected Classifier accomplished the maximum performance comparing others where 66% Split done 80.08% and Attribute Selected Classifier done 80.69%.

Actually Meta Classifier is habitually a proxy to the principle classifier, used to supply excessive data preprocessing. That's why we choose Meta classifier for our comparative analysis and we hope that we could make a better a comparison than others.

II. RELATED WORKS

Diabetes has been announced globally as an epidemic. This pestilence diseases could be hugely attributed to the quick growth in the rate of physical inactivity, fatness, and overweight. According to the survey of WHO, about 350 million diabetes affected people are suffering in today's world. Diabetes will rank as seventh of the leading cause of death global by 2030. It is expected that diabetes will be rising by 50% during the upcoming 10 years. In low rated and middle-income countries, 4 out of 5 people are leading lives with diabetes [11].

Sandeep Kumar Budhani et al., [7] has studied three Meta Classifier Algorithms: Adaboost M1, Stacking, and Bagging which has applied to diabetes dataset: Hyperplane1 and Hyperplan2 and WEKA was used here as a data mining tool for measuring performance. According to this paper, Bagging has shown the highest accuracy rate for both Hyperplane1 and Hyperplan2 datasets and that are respectively 84.54% and 83.83%.

Lujain AlThunayan et al., [5] used a diabetes dataset for comparing Bayesian, Naive Bayes, J48, Random Forest, Random Tree, REP Tree, CART, and SMO classification algorithms that has helped to find out the best classification algorithm among them by measuring the accuracy of those classifiers.

Mirza Shuja et al., [9] wanted to present a detailed survey of various techniques of data mining that have been used to design prognostic models which will be helpful for other data mining researchers to predict diabetes.

P. Suresh Kumar et al., [11] proposed a pattern to reduce the problems created in most useful data mining techniques like classification and clustering. That helps to apply those techniques easily to collected diabetes data. Especially it helps to predict the risk of gestational diabetes.

Sajida Perveen et al., [13] made some models with a better classification’s output of diabetes where the diabetes dataset is made with three age groups in the population of Canada, collected from Canadian Primary Care Sentinel Surveillance Network database. As a result, Adaboost has given better performance than bagging as well as J48 decision tree.

Lakshmi Devasena et al., [4] has scrutinized the proficiency of J48 Classifiers, Random Forest and REP Tree for the credit venture prognostics and compare their vigor through different measurements. As a dataset, the German Credit Dataset was used here. After final observation, Random Forest Classifier has performed best comparing respectively REP Tree Classifier and J48 Classifier. This paper didn't work with diabetes but, the scrutinized way in this paper was mind-blowing as comparing other's comparison works which will be helpful in diabetes research using data mining techniques.

Aiswarya Iyerv et al., [2] used Pima Indians Diabetes Dataset for analyzing the models using by Decision Tree and Naive Bayes classifiers, which help to build a most feasible model to search out the endemically and systematical dealing for diagnosing diabetes. This model will be helpful for flourishing the automation of diabetes scrutinize.

Razieh Asgarnezhad et al.,[10] has proposed a scheme with a proficient preprocessing technique together with absence value exploration(replace with mean) & optimize volition using the genetic algorithm on a diabetes dataset from Pima Indians database where this dataset has the lack of completeness. By using the SVM classifier which has predicted 84.35% accuracy rate which has the highest accuracy among the conferred comparison.

N. A. Nnamoko et al., [3] presented a diabetes prognostic model by investigating the way of predictions from different classification algorithms, repeating the task, could be utilized to output a greater performance comparing the highest separately learning algorithm. In this paper, RBF, RIPPER, SMO, Naive Bayes, and C5.4 have trained to build up to five populated models. After comparison, a Meta model with a Logistic Regression algorithm was used to train and make final prognostics using the output of the maximum and minimum performing algorithms as extra outputs. As a final result, C4.5 has performed the highest performance with a 77.9 % accuracy rate of classification and RBF has performed the lowest performance with a 73.6 % accuracy rate of classification. On the other hand, the Meta Model accomplished 77.0% accuracy rate of classification.

Nithya Settu et al.,[6] researched for diabetes and improved the performance of the filter algorithm by using Symmetrical Uncertainty Measure (SUM) and Novel Symmetrical Uncertainty Measure (NSUM) where SUM technique has reached 79.08% accuracy rate with0.06 sec run time and NSU technique has achieved 89.12% accuracy rate in 0.03 sec run time. Both techniques had applied through WEKA.

Retrieval Number: D2070029420/20200BEIESP
DOI: 10.35940/ijitee.D2070.039520
III. META CLASSIFIER OVERVIEW

Almost 20 Meta Classifiers are existing in WEKA. We choose only six of them for scrutinizing and comparing according to observing their accuracy rate in both cross-validation and splitting techniques. And they are

- Multiclass Classifier Updatable
- Attribute Selected Classifier
- Ada Boost M1
- Logit Boost
- Bagging
- Filtered Classifier

A. Ada Boost M1

Ada Boost M1 is an extensively executed boosting algorithm that advantage to known well. For boosting a multiclass basis classifier as if the multiclass classification is consists of a problem, this classifier is used. Because of the too much weakness in the base classifier, AdaBoost M1 won't work. But, after the interchange in Ada Boost M1 in one line only, it can be prepared as applicable. In our research, we used Hoeffding Tree as an internally used classifier.

B. Bagging

Bagging is also known as only Meta-Bagging also. Bagging is known as bootstrap aggregation. Bagging generates training data with bootstrap samples. It develops a distinct training set including numerous datasets. Various datasets are formed by unmethodical sampling happening with replacement. Each individual bootstrap specimen is used to train a classifier or, a regression function. Classification outputs are taken on the highest value of votes for classification intentions. For regression mean of prospective values are taken. Alternation is decreased and performance is developed for insecure classifiers that disagree meaningfully with tiny changes in the dataset. In the configuration of Bagging in our paper, REPTree is used as an internally used classifier.

C. Filtered Classifier

Restoring the architecture of the testing and training datum analogous to this classifier is used with different types of filters. Here in our paper, we configured Filtered Classifier where as a classifier we used Random Forest and as a filter we used Discretize.

D. Logit Boost

Logit Boost is the succession of the Ada Boost algorithm as it alternates the interpretative loss of the Ada Boost algorithm to temporary Bernoulli possibility loss. Logit Boost is used for the execution of preservative logistic regression. Here in our research work, we configured Logit Boost by determining Random Forest as internally used classifier.

E. Multiclass Classifier Updatable

Multiclass Classifier Updatable is an upgrade version of Multiclass Classifier. Error purification codes are modified with this classifier for achieving for much accuracy as this classifier is applied for categorizing events added to two classes. In our analysis, we used SGD as an internally used classifier in Multiclass Classifier Updatable.

F. Attribute Selected Classifier

The limit of the testing data and training data is reduced by Attribute Selected Classifier before being expired onto the classifier. Currently, researchers used base classifiers. So, the classifier is promoted several search ways are used during the stage of attribute selection. Here as an internally used classifier we used Hoeffding Tree, as evaluator we used cfsSubsetEval and as search, we used BFS.

IV. EXPERIMENTAL MODEL

We used 3 types of cross-validation techniques: 3-fold, 5-fold and 10-fold, and also 3-types of percentage split techniques: 66% Split, 75% Split and 80% Split. We collected our diabetes dataset from the Pima Indian Database and applied those techniques with declared Meta classifiers. Then, we scrutinized our output in a synchronized way. And finally, we presented our spontaneous opinion after some valid analyses and comparisons.

A. Dataset

These declared data mining techniques have been applied to the Pima Indians Diabetes Database of National Institute of Diabetes and Digestive and Kidney Diseases. This dataset is collected from the UCI Machine Learning Repository. This dataset is consists of 768 record samples with 9 attributes which are used to predict the presence of diabetes after analyzing deeply. It is known as a renowned diabetes dataset in the data mining research area. A large number of diabetes-related data mining research has completed by using this data set. That’s why we used this data set in our experiment. There is no doubt to accept its validity in the research area. Among 768 samples, 268 samples are tested positive and the rest of them are tested negative. All samples carried persons are Indian women with a minimum of 21 years old and live near Phoenix, Arizona, USA. The description of the attributes are listed in below.

<table>
<thead>
<tr>
<th>Attribute</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>Preg</td>
<td>Number of times pregnant</td>
</tr>
<tr>
<td>Plas</td>
<td>Plasma glucose concentration</td>
</tr>
<tr>
<td>Pres</td>
<td>Diastolic blood pressure</td>
</tr>
<tr>
<td>Skin</td>
<td>Triceps skin fold thickness</td>
</tr>
<tr>
<td>Insu</td>
<td>2-Hour serum insulin</td>
</tr>
<tr>
<td>Mass</td>
<td>Body mass index</td>
</tr>
<tr>
<td>Pedi</td>
<td>Diabetes pedigree function</td>
</tr>
<tr>
<td>Age</td>
<td>Patient Age (years)</td>
</tr>
<tr>
<td>Class</td>
<td>Class variable (0 or 1)</td>
</tr>
</tbody>
</table>

B. Internal Classifier

We have already declared in Meta Classifier Overview that in every Meta Classifier has to be used a main particular classifier with many various parameters which we said as an internally used classifier. Because, we know that Meta Classifier is habitually a proxy to the principle classifier, used to supply excessive data preprocessing. In our selected Meta Classifiers, we used the below classifiers as the internally used classifiers.

<table>
<thead>
<tr>
<th>Meta Classifier’s Names</th>
<th>Internally used Classifiers</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ada Boost M1</td>
<td>Hoeffding Tree</td>
</tr>
<tr>
<td>Bagging</td>
<td>REPTree</td>
</tr>
<tr>
<td>Filtered Classifier</td>
<td>Random Forest</td>
</tr>
</tbody>
</table>

Table- II: List of Internally used Classifiers in Meta Classifiers

Published By: 
Blue Eyes Intelligence Engineering & Sciences Publication

DOI: 10.35940/ijitee.D2070.039520
Implications of Meta Classifiers for Onset Diabetes Prediction

<table>
<thead>
<tr>
<th>Logit Boost</th>
<th>Random Forest</th>
</tr>
</thead>
<tbody>
<tr>
<td>Multiclass Classifier Updatable</td>
<td>SGD</td>
</tr>
<tr>
<td>Attribute Selected Classifier</td>
<td>Hoeffding Tree</td>
</tr>
</tbody>
</table>

- **Hoeffding Tree**: A Hoeffding tree known as a progressive decision tree which is able to learn from huge data drifts at any time with the assumption that the change according to time can't possible by distribution yield instances.

- **REPTree**: REPTree is known as an algorithm of quick decision tree because it can build a decision tree using variability or, obtaining information. It can prune in a fast way as it also can deal with absence values using splitting parts into shreds. The missing values operation is also similar like C4.5 Algorithm. This pruning process is called reduced-error-pruning [1].

- **Random Forest**: After so many combinations of tree predictors where every tree relies upon the values come from a random vector sampled automatically and with the equivalent dispensation for the whole tress in the forest [4]. Pruning is not needed here. Because, until every node holds simply very small number of monitoring, trees can be generated.

- **SGD**: SGD generally replaces all absence values and the nominal attributes transform in binary forms. The coefficients in the result are depending on t nominal attributes transform in binary forms. SGD holds simply very small number of monitoring, trees can from a random vector sampled automatically and with the equivalent algo predictors where every tree relies upon the values come from a random vector sampled automatically and with the equivalent algo.

- **Hoeffding Tree known as a**

C. **Performance Measured Used**

For measuring the performance of the declared Meta Classifier, we used numerous values that come from different sectors.

- **Accuracy Rate of Classification**: Accuracy Rate of Classification is computed as exactly classified samples divided by the entire number of samples multiplied by 100. Exact classified sample is the sum of True-Positive (TP) and True-Negative.

  \[
  \text{Accuracy Rate} = \frac{TP + TN}{Total} \times 100
  \]  

  (1)

- **Precision**: According to the Confusion Matrix, Precision is the ratio between true-positive samples and predicted yes samples.

  \[
  \text{Precision} = \frac{TP}{TP + FP}
  \]  

  (2)

Here, TP+FP = Predicted Yes

- **Recall**: Recall is also known as Sensitivity. According to the Confusion Matrix, Recall is the ratio true-positive samples and actual yes samples.

  \[
  \text{Recall} = \frac{TP}{TP + FN}
  \]  

  (3)

Here, TP+FN = Actual Yes

- **F-Score**: F-Score is also called F1-Score or, F-Measure. The F-Score can give a more feasible measurement of a test implementation using both recall and precision. When the value of F-Score becomes 1 that indicates the perfection of both recall and precision.

  \[
  \text{F-Score} = \frac{2 \times \text{Precision} \times \text{Recall}}{\text{Precision} + \text{Recall}}
  \]  

  (4)

- **Mean Absolute Error**: MAE calculates the average measurement of the errors in a set of prognostics, except considering their way.

  \[
  \text{MAE} = \frac{\sum{|p_i - a_i|}}{n}
  \]  

  (5)

- **Root Mean Squared Error**: RMSE is the square root of the mean of squared differences between prophecy and actual esplial.

  \[
  \text{RMSE} = \sqrt{\frac{\sum{(p_i - a_i)^2}}{n}}
  \]  

  (6)

- **Matthews Correlation Coefficient**: MCC calculates the quality of the classification which has two types. Actually, the value of MCC proposed the correlation coefficient among the predicted and noticed classification which is binary. According to Confusion Matrix, the formula of MCC will be,

  \[
  \text{MCC} = \frac{TP \times TN - FP \times FN}{(TP + FP)(TP + FN)(TN + FP)(TN + FN)}
  \]  

  (7)

- **Kappa Statistic**: Kappa Statistic is also known as Cohen's Kappa. Actually, it is used for quantifying the ability of reproduction of a distinct variable.

  \[
  K = \frac{p_o - p_e}{1 - p_e}
  \]  

  (8)

Here, \( p_o \) = Observed Agreement and \( p_e \) = Expected Agreement.

D. **Data Mining Techniques**

We have already said that we used cross-validation and percentage split techniques in numerous ways. And scrutinizing the results in different ways as Precision-Recall Curve, ROC curve, bar chart using MCC values and Win-Draw-Loss table.

- **Cross-Validation**: Cross-Validation is a heuristic works that arbitrarily classify the data into n-folds, each with nearly the similar number of records, makes n-models using the similar algorithms and training parameters where every model is trained with n-1 folds of the data and tested on the due fold, can be applied to search the best algorithm and its optimum training parameters.

- **Percentage Split**: Percentage Split is a process of re-sampling that reserves \( n \% \) of the rows as the training dataset for structuring the model and (n-100) \% of the rows reserved as the test dataset to test the model. The target classifier is trained as opposed to the trained data. On the other hand, the classification accuracy is measured on the test dataset.
Win-Draw-Loss Table: The win-draw-loss table represents the winning rate comparing other classifiers. Here, the value of a win, draw or, a loss will be equal or, less than the total numbers of comparison classifiers. Actually, it is the most applicable is biological research but, in the data mining research area as a comparison, it would be helpful. According to the value of the win, we can easily estimate the win-rate of a classifier.

\[
\text{Win Rate} = \frac{\text{Total Number of Win}}{\text{Total Number of Comparison}} \times 100 \quad (9)
\]

Receiver Operating Characteristic Curve: ROC curve is a graphical plot that representing the performance of a classifier at whole classification thresholds. According to confusion matrix, this curve consists of two parameters. One is called True-Positive Rate and another is called False-Positive Rate.

\[
\text{TPR} = \frac{TP}{TP + FN} \quad (10)
\]

\[
\text{FPR} = \frac{FP}{FP + TN} \quad (11)
\]

A classifier builds a model that has no skill if it illustrates at the point (0.5, 0.5) or, by a diagonal line that comes from the bottom left of the ROC Curve to the top right and contains an AUC of 0.5. A classifier builds a model that has that perfect skill if it illustrates by a line that comes from the bottom left of the ROC Curve to the top left of the ROC Curve and moved the top right of the ROC Curve.

Precision-Recall Curve: A precision-recall curve is a graphical plot for various thresholds where according to X-axis, the recall values are placed and according to Y-axis the precision values are placed. This curve is useful in applied data mining for estimating binary classification patterns. A classifier builds a model that has no skill if it illustrates by a diagonal line that comes from the top left (0, 1) of the precision-recall curve to the bottom right (1, 0). A skillful model is illustrated as a point at (1, 1). A classifier builds a model that has that perfect skill if it illustrates by a line that comes from the top left of the precision-recall curve to the top right of the precision-recall curve and moved to the bottom right of the precision-recall curve.

MCC Bar Chart: After plotting, the MCC values in the bar chart we can easily indicate the perfectly closed prediction because, if the value comes to close to +1 then, the result will be considered as close to perfect predictions. On the other hand, if the value comes to close to -1 then, it will be considered as close to worst predictions. And the value 0 indicates better than random prognostic.

V. COMPARATIVE ANALYSIS

For comparing, the below processes are followed in our study.

At first, we have applied numerous values with various internally used classifiers for both cross-validation and percentage split techniques. And we have selected some particular values and internally used classifiers (shown in table 01) for applying again in the cross-validation and percentage split techniques on the basis of accuracy variance and differences from each other. Here, we have selected 3-fold, 5-fold, and 10-fold for cross-validation and also selected 66%, 75% and 80% for percentage split.

Then we have recorded the value for precision, recall, f-score, accuracy, kappa statistics, MAE, RMSE, MCC and ROC area from outputs which are helped us to enhance our comparison in a further step. In table 03 and table 05, we have added two extra parameters named average accuracy and standard deviation of Accuracy in both row and column with completing their calculation. All records have shown in table 03, table 04, table 05 and table 06.
Implications of Meta Classifiers for Onset Diabetes Prediction

<table>
<thead>
<tr>
<th>Meta Classifier</th>
<th>3-Fold Cross Validation</th>
<th>5-Fold Cross Validation</th>
<th>10-Fold Cross Validation</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Kappa Statistic</td>
<td>MAE</td>
<td>RMSE</td>
</tr>
<tr>
<td>Ada Boost M1</td>
<td>0.4703</td>
<td>0.3023</td>
<td>0.408</td>
</tr>
<tr>
<td>Bagging</td>
<td>0.4576</td>
<td>0.307</td>
<td>0.3997</td>
</tr>
<tr>
<td>Filtered Classifier</td>
<td>0.4445</td>
<td>0.3179</td>
<td>0.4233</td>
</tr>
<tr>
<td>Logit Boost</td>
<td>0.4561</td>
<td>0.2505</td>
<td>0.4554</td>
</tr>
<tr>
<td>Multiclass Classifier Updatable</td>
<td>0.4693</td>
<td>0.2305</td>
<td>0.4801</td>
</tr>
<tr>
<td>Attribute Selected Classifier</td>
<td>0.4524</td>
<td>0.3007</td>
<td>0.4135</td>
</tr>
</tbody>
</table>

Table- IV: Kappa Statistic, MAE, RMSE, MCC & ROC Area of Meta Classifiers for Cross-Validation

After scrutinizing table 03 and table 05, Attribute Selected Classifier and Filtered Classifier both have given the highest accuracy of 81.9923% individually in the 66% split (Table 05). And on average, 66% split has achieved the highest accuracy of 80.08% as a technique and Attribute Selected Classifier has achieved the highest accuracy of 80.69% as a classifier (Table 05). Individually In table 03, as a classifier, Multiclass Classifier Updatable has achieved the highest accuracy of 80.69% as a classifier (Table 05). According to table 06, we have calculated the win rate for each Meta Classifier according to 10-fold cross-validation. The highest win rate has got the highest accuracy of 76.17%.

Here, we have produced a win-draw-loss shown in table 06. We have calculated the win rate for each Meta Classifier shown in table 06. According to table 06, we have identified that Multiclass Classifier Updatable has achieved the highest win rate.

For the view of comparison, from ROC area values of table 04 and table 06, we have made ROC curves for each selected Meta Classifier according to 10-fold cross-validation (Fig.02) and 66% split (Fig.03).

For comparing another angle of view, from precision and recall values of table 03 and table 05, we have made precision-recall graphs for both 10-fold cross-validation (Fig.04) and 66% split (Fig.05) and also have shown the precision-recall curves (Fig.06 & Fig.07) for every declared Meta Classifier.

Lastly, we have represented two bar charts (Fig.08 & Fig.09) from MCC values of table 04 and table 06 for calculating the qualities of prediction.

Table- V: Percentage Split wise performance metrics for Meta-Classifiers
Fig. 2. Receiver Operating Characteristics (ROC) Curves of Meta Classifiers for Tested Positive Class according to 10-Fold Cross Validation
Implications of Meta Classifiers for Onset Diabetes Prediction

Fig. 3. Receiver Operating Characteristics (ROC) Curves of Meta Classifiers for Tested Positive Class according to 66% Split

Fig. 4. Precision-Recall Graph for 10-Fold Cross Validation

Fig. 5. Precision-Recall Graph for 66% Split

Fig. 6. Precision-Recall (PR) Curves of Meta Classifiers for Tested Positive Class according to 10-Fold Cross Validation

Fig. 7. Precision-Recall (PR) Curves of Meta Classifiers for Tested Positive Class according to 66% Split

Fig. 8. MCC Value for 10-Fold Cross Validation

Fig. 9. MCC Value for 66% Split
VI. CONCLUSION

This comparison investigated the overall efficiency of the six Meta Classifiers namely, Multiclass Classifier Updatable, Attribute Selected Classifier, Ada Boost M1, Logit Boost, Bagging, Filtered Classifier for forecasting diabetes. And finally, Attribute Selected Classifier and Filtered Classifier perform better than others in the individual platform. On average, 66% split as a technique and Attribute Selected Classifier as a classifier give the best performance. But, most importantly according to the win rate, Multiclass Classifier Updatable takes place over all of them.

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AUTHORS PROFILE

Md. Ashaf Udduula, is with the Department of Computer Science and Engineering (CSE) of Daffodil International University. His research interests are Data Mining, Machine Learning, Artificial Intelligence and Embedded System.

Md. Al - Amin Hossain, is with the Department of Computer Science and Engineering (CSE) of Daffodil International University. His research interests are Data Mining, Machine Learning, Artificial Intelligence and IoT.

Md. Khalid Hossen, is with the Department of Computer Science and Engineering (CSE) of Daffodil International University. His research interests are Data Mining, Machine Learning and Image Processing.

Ahmed Al Marouf, is with the Department of Computer Science and Engineering (CSE) of Daffodil International University (DIU). He received his Bachelor degree from the Department of Computer Science and Engineering (CSE), IUT in 2014 and M.Sc. Engg. in CSE from the same university in 2019. He was a graduate researcher of Systems and Software Lab (SSL) in the CSE department of IUT. His research interest lies within Computational social science, Data science and Machine learning He is currently working as a lecturer in Department of Computer Science and Engineering (CSE) of Daffodil International University (DIU), Dhaka, Bangladesh. He also acts as the Technical Lead of DIU HCI (Human Computer Interaction) Research Lab.