



*Research article*

## **Enhancing cardiovascular disease prediction: A hybrid machine learning approach integrating oversampling and adaptive boosting techniques**

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### **Appendix 1 (All Algorithm)**

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#### **Algorithm 1 SmoteR(D,tE,o.u,k)**

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1. Initialize empty sets:
  - o rareHD to store instances with high relevance of heart disease ( $\phi(y) > tE$ ) and  $y = 1$
  - o rareNHD to store instances with high relevance of no heart disease ( $\phi(y) > tE$ ) and  $y = 0$
  - o newCasesHD to store synthetic cases for rareHD
  - o newCasesNHD to store synthetic cases for rareNHD o newCases as the concatenation of newCasesHD and newCasesNHD o normCases as the set for undersampling
2. For each instance  $(x, y)$  in D:

- o If  $\phi(y) > tE$  and  $y = 1$ , add  $(x, y)$  to rareHD o If  $\phi(y) > tE$  and  $y = 0$ , add  $(x, y)$  to rareNHD

3. Generate synthetic cases for rareHD: o newCasesHD = genSynthCases(rareHD, %o, k)
4. Generate synthetic cases for rareNHD:
  - o newCasesNHD = genSynthCases(rareNHD, %o, k)
5. Concatenate newCasesHD and newCasesNHD: o newCases = newCasesHD newCasesNHD
6. Determine the number of cases for under-sampling:
  - o nrNorm = %u of |newCases|
7. Randomly select cases from  $D\{\text{rareHD} \text{ rareNHD}\}$  for under-sampling:
  - o normCases = random sample of nrNorm cases from  $D\{\text{rareHD} \text{ rareNHD}\}$
8. Return the concatenated synthetic and under-sampled cases:
  - o Return newCases normCases

end

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## Algorithm 2 for XGB

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1. Import the essential libraries:
  - o pandas as pd o numpy as np o xgboost as xgb
  - o train\_test\_split from sklearn.model\_selection o accuracy\_score from sklearn.metrics
2. Load the heart disease dataset into a pandas DataFrame.
3. Pre-process the data:
  - o Perform data cleaning and handle missing values. o Conduct feature selection based on domain knowledge or statistical techniques.
  - o Normalize or standardize the features if necessary. o Divided the data into training with testing sets using train\_test\_split() function, considering stratification if needed.
4. Define the XGBoost model:
  - o Set the hyperparameters for the XGBoost model, such as the number of trees, learning rate, and maximum depth.
  - o Optionally, perform cross-validation or grid search for hyperparameter tuning.
5. Train the model: o Fit the XGBoost model using the training data.
6. Evaluate the model:
  - o Make predictions on the testing data using the trained model. o Compute evaluation metrics specific to heart disease prediction, such as accuracy, precision, recall, and F1-score. o Analyze the performance of the model and consider any issues, such as overfitting or underfitting.
7. Interpret the results:

- o Investigate the importance of features in predicting heart disease. o Identify any patterns or relationships between features and the target variable.
- o Consider the impact of individual features on the model's predictions.

8. Iterate and refine:

- o Based on the results, iterate and refine the model by adjusting hyperparameters, modifying feature engineering techniques, or exploring different algorithms.
- o Consider additional techniques like ensemble methods or addressing class imbalance if necessary.

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Algorithm 3 for ET

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1. Import the necessary libraries:
  - o pandas as pd o numpy as np
  - o ExtraTreesClassifier starting sklearn.ensemble o train\_test\_split after sklearn.model\_selection o accuracy\_score, confusion\_matrix from sklearn.metrics
2. Load the heart disease dataset into a pandas DataFrame:
  - o data = pd.read\_csv('heart\_disease\_data.csv')
3. Pre-process the data:
  - o Separate the features (X) from the target variable (y). o Splitting data's into training with testing sets use train\_test\_split() function, considering stratification if needed.
4. Define the Extra Trees model:
  - o Initialize the ExtraTreesClassifier model.
5. Train the model: o Fit the Extra Trees model using the training data.
6. Make predictions: o Generate predictions for the testing data using the trained model.
7. Evaluate the model:
  - o Compute the accurate model by comparing predicted labels with the actual labels.
  - o Generate the confusion matrix to assess the outcome of the model.
8. Print the accuracy and confusion matrix:
  - o Print the accuracy score.
  - o Print the confusion matrix.

End

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#### Algorithm 4 for RF

1. Import the necessary libraries:
  - o pandas as pd o numpy as np
  - o RandomForestClassifier after sklearn.ensemble o train\_test\_split after sklearn.model\_selection o accuracy\_score, confusion\_matrix from sklearn.metrics
2. Load the heart disease dataset into a pandas DataFrame:
  - o data = pd.read\_csv('heart\_disease\_data.csv')
3. Preprocess the data:
  - o Separate the features (X) from the target variable (y).
  - o Divide the data into training with testing sets use train\_test\_split() function, considering stratification if needed.
4. Define the Random Forest model:
  - o Initialize the RandomForestClassifier model.
5. Train the model:
  - o Fit the Random Forest model using the training data.
6. Make predictions:
  - o Generate predictions for the testing data using the trained model.
7. Evaluate the model:
  - o Compute the accurate model through comparing the predicted labels with the actual labels.
  - o Generate the confusion matrix to assess the performance of the model.
8. Print the accuracy and confusion matrix:
  - o Print the accuracy score.
  - o Print the confusion matrix.

End

### Algorithm 5 for AdaBoost

1. Input:
  - o Training input data:  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$ , where  $x_i$  represents the features of the  $i$ th patient and  $y_i$  is the corresponding label (+1 for heart disease present, -1 for heart disease not present).
  - o Number of iterations:  $T$
2. Output:
  - o Boosted hypothesis for heart disease prediction:  $H(x)$
3. Step 1: Initialize weights for training samples
  - o Set  $D_1(n) = 1/N$  for all  $n$ , where  $N$  is the total number of training samples.
4. Step 2: Iterate  $T$  times
  - o For  $t = 1$  to  $T$ :

- Train a weak learner  $h_t(x)$  using the weighted training data.
- $h_t(x)$  predicts the presence or absence of heart disease (+1 or -1) based on the patient's features.

5. Step 3: Calculate weighted error rate and weight of the weak learner o      Compute the weighted error rate  $\epsilon_t$  using the equation:  $\epsilon_t = \sum_{n=1}^N D_t(n) \cdot 1\{h_t(x_n) \neq y_n\} / \sum_{n=1}^N D_t(n)$  o      Calculate the weight  $\alpha_t$  for the weak learner using the equation:  $\alpha_t = 0.5 * \ln((1 - \epsilon_t) / \epsilon_t)$

6. Step 4: Update the weights of the training samples o      For  $n = 1$  to  $N$ :

If  $h_t(x_n) = y_n$ , then:  $D_{t+1}(n) = D_t(n) * \exp(-\alpha_t)$

If  $h_t(x_n) \neq y_n$ , then:  $D_{t+1}(n) = D_t(n) * \exp(\alpha_t)$

7. Step 5: Normalize the updated weights o      Normalize the updated weights  $D_{t+1}(n)$  by dividing them by the sum of all updated weights:  $D_{t+1}(n) = D_{t+1}(n) / \sum_{m=1}^N D_{t+1}(m)$

8. Step 6: Repeat steps 2-5 for  $T$  iterations.

9. Step 7: Combine weak classifiers to create the boosted hypothesis o      For a new input sample  $x$ :

Calculate the boosted hypothesis  $H(x)$  as:  $H(x) = \text{sign}(\sum_{t=1}^T \alpha_t h_t(x))$

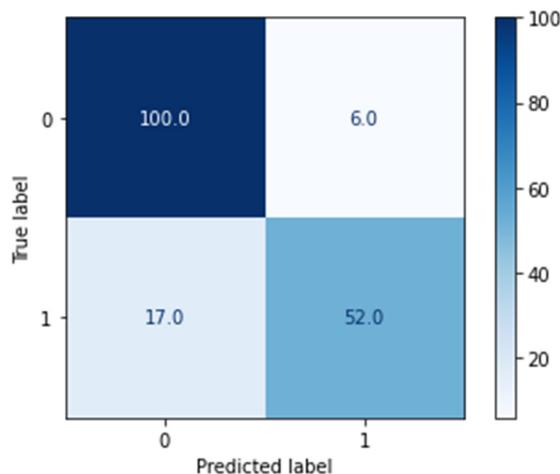
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## Appendix 2 (Validation)

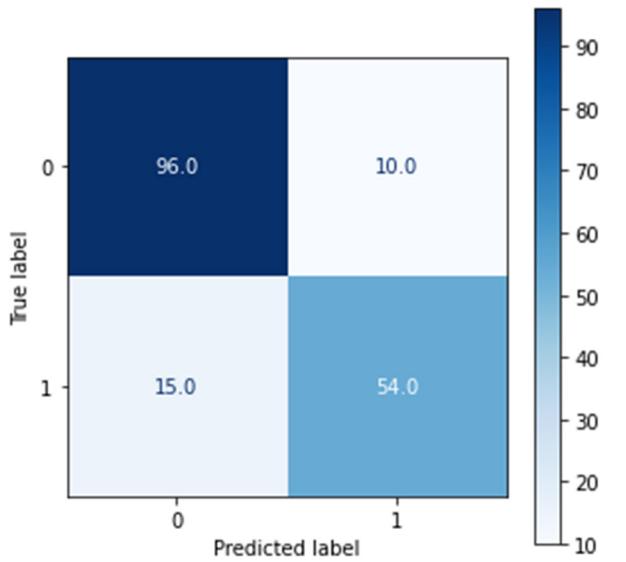
### Random Forest Results

Accuracy [ with RF]: 86.85714285714286 %  
Recall [ with RF]: 75.36231884057972 %  
precision [ with RF]: 89.65517241379311 %  
MCC [ with RF]: 0.723628103990507

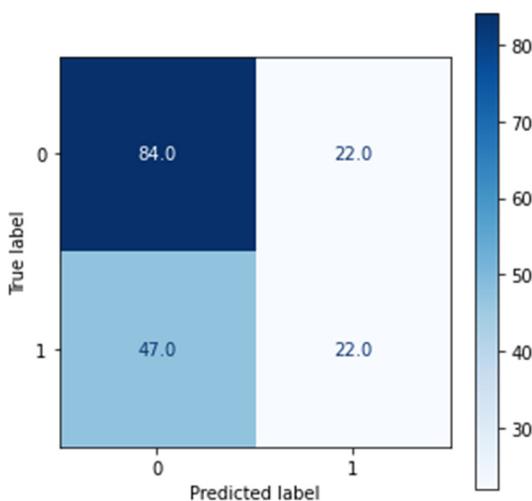


### AdaBoost Random Forest Results

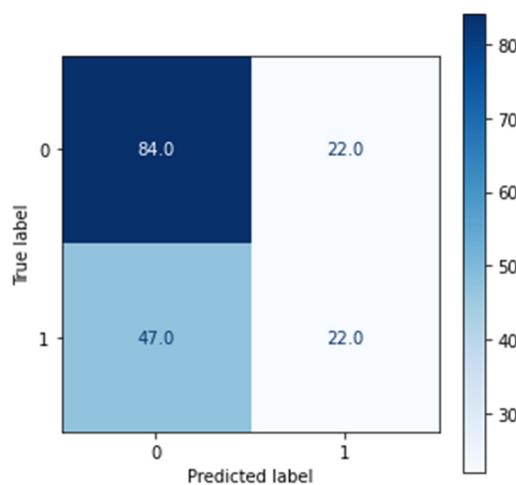
Accuracy [AdaBoost with RF]: 85.71428571428571 %  
Recall [AdaBoost with RF]: 78.26086956521739 %  
precision [AdaBoost with RF]: 84.375 %  
MCC [ AdaBoost with RF]: 0.6983678802480235



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Logistic Regression Results
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Accuracy [ with LR]: 60.57142857142858 %
Recall [ with LR]: 31.88405797101449 %
precision [ with LR]: 50.0 %
MCC [ with LR]: 0.1253674928685844
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AdaBoost LR Results
=====
Accuracy [AdaBoost with LogisticRegression]: 60.57142857142858 %
Recall [AdaBoost with LogisticRegression]: 31.88405797101449 %
precision [AdaBoost with LogisticRegression]: 50.0 %
=====
MCC [ AdaBoost with LR]: 0.1253674928685844
```

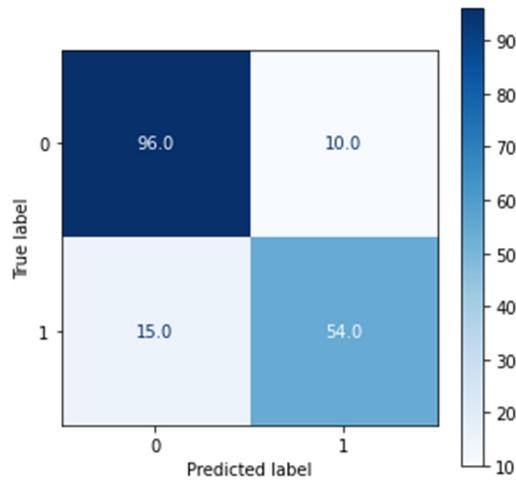


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**Extra Tree Results**

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Accuracy [ with ET]: 85.71428571428571 %  
 Recall [ with ET]: 78.26086956521739 %  
 precision [ with ET]: 84.375 %  
 MCC [ with ET]: 0.6983678802480235

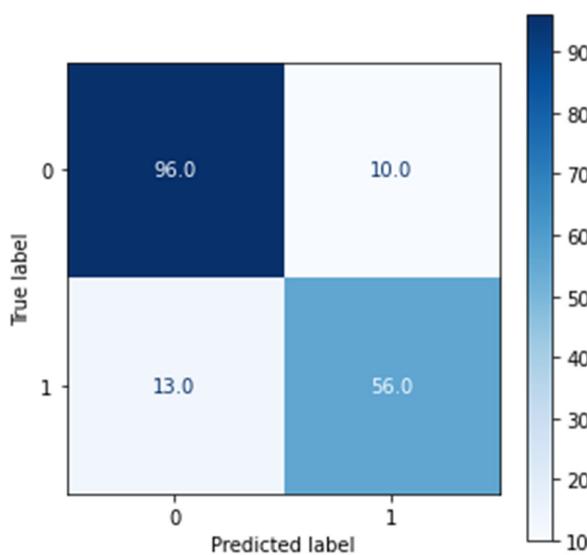


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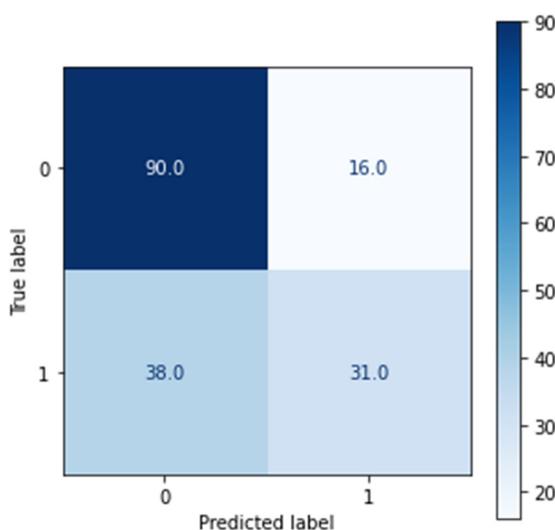
**Extra TreeForest Results**

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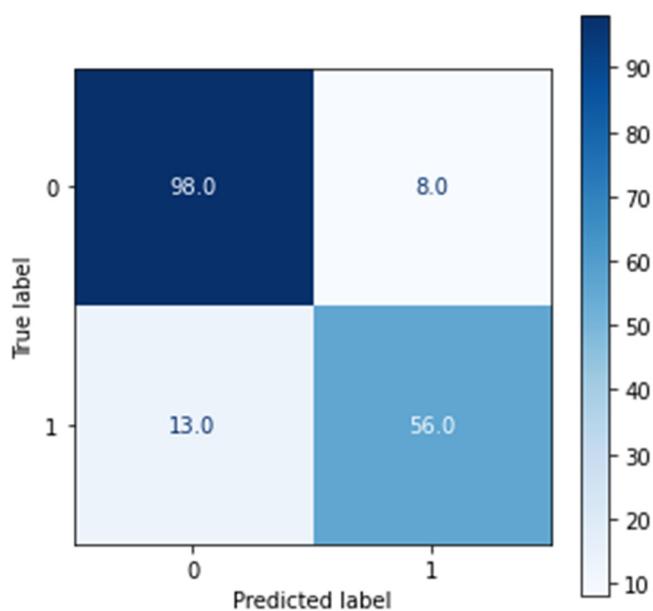
Accuracy [AdaBoost with ET]: 86.85714285714286 %  
 Recall [AdaBoost with ET]: 81.15942028985508 %  
 precision [AdaBoost with ET]: 84.84848484848484 %  
 MCC [ AdaBoost with ET]: 0.7232119465299363



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XGB Results
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Accuracy [ with ET]: 69.14285714285714 %
Recall [ with ET]: 44.927536231884055 %
precision [ with ET]: 65.95744680851064 %
MCC [ with ET]: 0.328945049232401
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XGB TreeForest Results
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Accuracy [AdaBoost with XGB]: 88.0 %
Recall [AdaBoost with XGB]: 81.15942028985508 %
precision [AdaBoost with XGB]: 87.5 %
MCC [ AdaBoost with XGB]: 0.7469234539641157
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