

Cardiovascular Disease Prediction using Supervised Machine Learning



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Abstract

Cardiovascular disease (CVD) or heart disease accounts for the leading cause of death worldwide. It is a matter for us to be concerned about today's highly chaotic lifestyle that leads to various diseases. Early prediction and identification of heart-related diseases can bring down the mortality rate. Consequently, many studies explore the realm of classification algorithms to find the best models. This paper follows the CRISP-DM Methodology to solve the problem of heart disease prediction. A relatively large dataset is selected from Kaggle, and it is cleaned by various preprocessing methods like imputation, normalization, encoding, transformation, and resampling. Although many ML models can be used, this paper focuses on Logistic Regression and Random Forest. In order to obtain a comprehensive result, the dataset is prepared using five different approaches where the combination of preprocessing methods has been changed. Since the dataset used in this study is highly imbalanced, a combination of various evaluation metrics like Confusion Matrix, Precision, Recall, and ROC curve have been used to analyze the performance of the two ML models. Finally, a detailed analysis of the result has been presented to explain how Random Forest is more appropriate for predicting heart diseases.

Data Preparation

Data Cleaning

Missing Values

Two terms to call missing of data values in this dataset. Given missing values that can detected by function, like NaN. Another one is values that have similar meaning with missing values, such as refused, not sure, & never checked.

Handle: Listwise deletion **Statistical Imputation**

Outliers exist in all numerical features. In feature BMI, outliers are under the lower bound, while the other two features are on the top of upper boud. Handle:

> Remove with IQR **Capping outliers**

Data Pre-Processing

Scaling / Normalization

The data is scaled using the minimum-maximum valuebased method. It scales the the range of 0 to 1. method is suitable for feature BMI, MentHlth, and PhysHlth, since they have skewed distribution.

Log Transformation

assume the data follow a normal distribution. A log value of the features within transform is helpful when dealing with numbers with a heavy-tailed distribution. All numerical features have positively skewed distributions.

Categorical Encoding

There are 2 methods applied to encode the data. Label Encoding

The target variable and all categorical

features, except Sex, are encoded

using this technique. **One-hot Encoding**

Feature Sex was applied to this method. It consists of two unique values: male and female.

Problem Understanding

Background of Study

The number of people suffering from cardiovascular Manual diagnosis and complex treatment. disease (CVD) or so-called heart disease is The study of the patient's medical history significantly rising every year. People do less manually is still used, reports of physical physical work because everything is automated. **The** examinations performed by medical medical equipment that exists today still takes time, professionals are less accurate and take a less accurate, and costly. Machine learning can be long time. personalize medical treatments for the people who have the probability to have CVD earlier.

Problem Statement

used to predict CVD. It can help the people take Inaccurate detecting technique. It is critical to the precaution steps and the medical officers to recognize the presence of this condition as soon as

> Lack of data. Accuracy could not improve due to a lack of large amounts of data or the nature of the dataset.

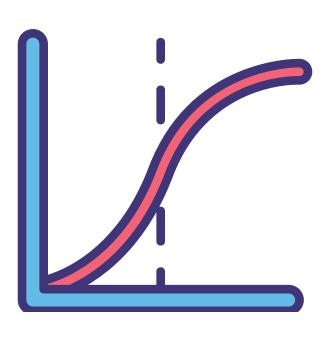
Objective Motivation

The main objective of this study is to predict the probability of patients to have cardiovascular disease (CVD) in the future by using machine learning. Random forest and logistic regression algorithms are being employed to predict CVD outcomes. The focus of this project is to maximize recall score by at least 95%, for it is risky to misdiagnose patients and could lead to death.

Algorithms

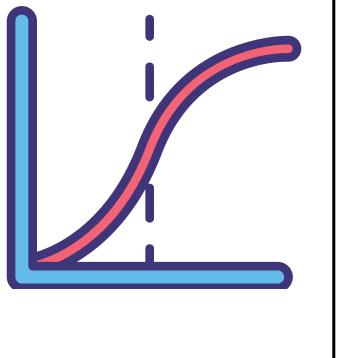
Logistic Regression

Logistic regression can solve classification problems through fits an S-shaped curve logistic function using the maximum likelihood In this case, the Logistic Regression model can show the relationship between the target variable and features, which features contribute the most to predict whether the people have a high probability to get heart disease or

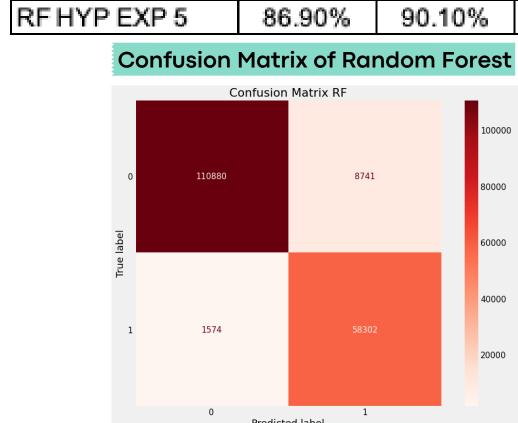


Random Forest

Random forest is one of the supervised learning algorithms which can handle both linear and non-linear data. Random forest predicts the new instance made by the trees that have low correlations. It is faster to fit the train data and predict the new instance with a large dataset. Furthermore, the combination of Random Forest and the resampling method can overcome the imbalanced class problem. Hence this is one of the proposed algorithms used in this project.



LR EXP 5 LR HYP EXP 5



Resampling methods

The imbalanced dataset

is commonly found in

medical diagnosis data.

In order to balance the

dataset, 3 resampling

Random Undersampling

Random Oversampling

Comparison of Model Performance

55.40%

62.60%

Results & Analysis

12.50%

9.10%

14.50%

5.60%

12.30%

9.80%

14.50%

5.30%

61.50%

61.00%

62.10%

62.00%

61.00%

61.00%

63.80%

81.30%

81.60%

92.10%

SMOTE

Model Experiment Precision

LR EXP 1

RF EXP 1

LR EXP 2

RF EXP 2

LR EXP 3

RF EXP 3

LR EXP 4

RF EXP 4

RF EXP 5

LR HYP EXP 1

RF HYP EXP

LR HYP EXP 2

RF HYP EXP 2

LR HYP EXP 3

RF HYP EXP 3

LR HYP EXP 4

RF HYP EXP 4

methods were applied:

72.50%

Figure above shows the importance of each features in both LR and RF respectively. These graphs shows how the two algorithms uses very different approach to solve the same problem. For LR, the gender of the patients played the most important roles in the classification process. But for RF, BMI and Age has the most influence in the model prediction.

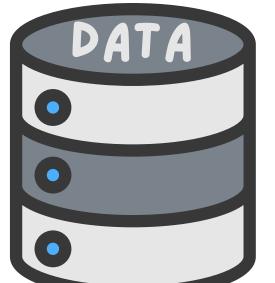
Feature Importances

Logistic regression (LR) has significantly higher precision than Random Forest (RF) in the first two experiments, where the imbalanced data is not handled. However, both Recall and AUC score was slighting higher for RF in these two experiments. Moving onto experiment 3 where random undersampling is applied, all the scores are significantly increased from the first and second experiments. The results from experiments 4 and 5 show that RF clearly outperforms LR in all Precision, Recall and AUC when the imbalanced dataset is handled with **oversampling.** Moreover, we can see that the random oversampling method give the best best for all the metrics in RF, whereas SMOTE-overasampling gives the best result for LR.

Conclusion

From 5 experiments were done. It can be concluded that the combination Random Forest algorithm and random oversampling (ROS) method achieve the best score for all evaluation metrics, with the precision of 86.9%, recall of 97.3%, and AUC score of 94.9%. That means a combination of Random Forest and ROS can do the classification properly. In comparison, Logistic Regression achieved the highest score with SMOTE technique. However, it is not as good as the Random Forest score. For future works, it is suggested to use TensorFlow to handle large-scale data in machine learning so that the model can be run faster. It also might be possible to improve the score using another algorithm and combination of resampling method.

Data Understanding



- Categorical 15 Attributes

- HighBP
- Veggies Fruits HighChol
- CholCheck Smoker
- AnyHealthcare GenHlth
- Stroke
- DiffWalk Diabetes Sex
- PhysActivity Age group HvyAlcoholConsump

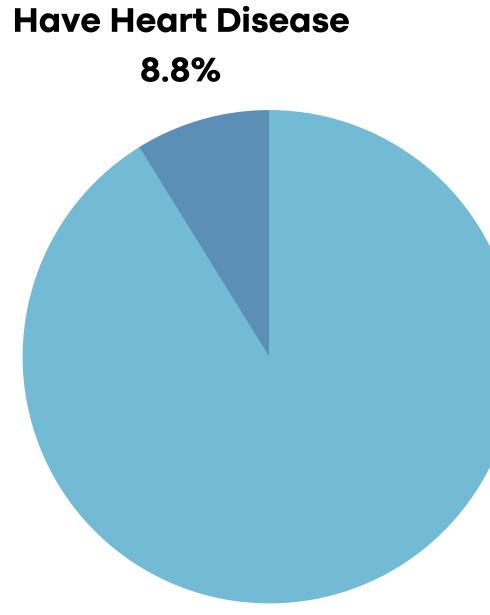
Numerical Attributes

- BMI
- MentHlth

PhysHlth Kaggle Dataset source : RFSS 2015 : https://rb.gy/unj6fv Alex Teboul : https://rb.gy/8thljp

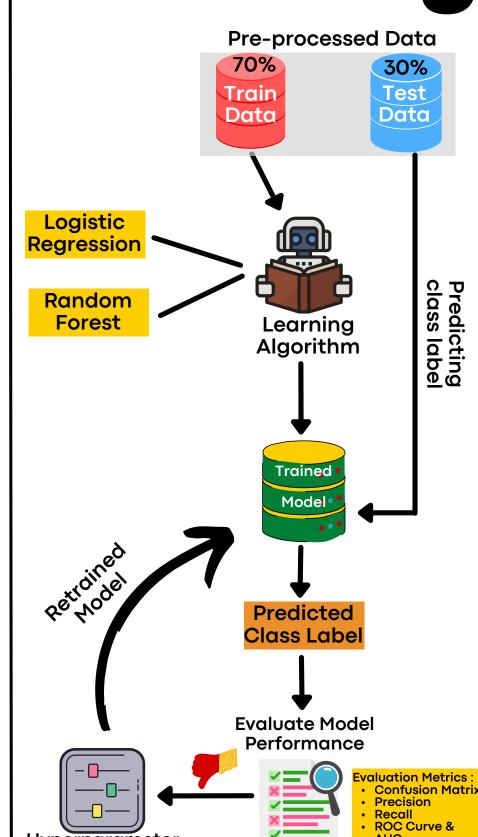
Target

? Class Labels





Modeling



Random Forest

Tools

- Python 3.9
- Pandas Scikit Learn
- Seaborn

Jupyter Lab

- Matplotlib

Experiment 1

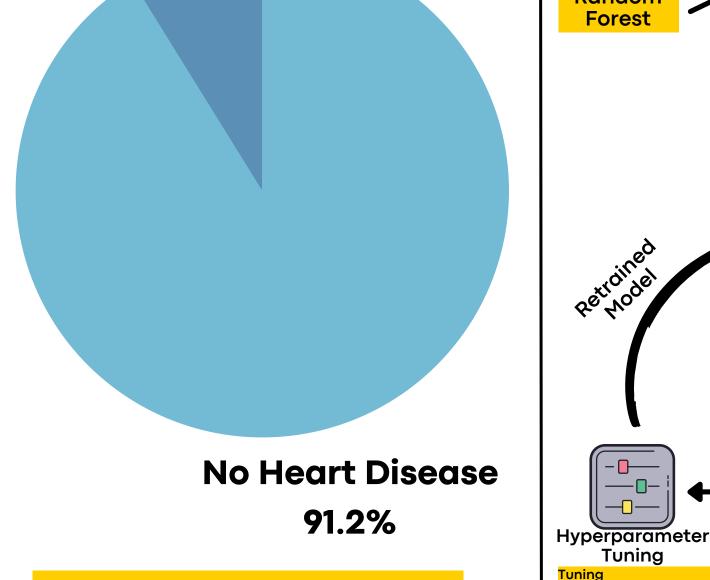
- Remove all missing values
- Remove outliers-IQR
- Normalization Log Transform
- Encoding

Experiment 3

Experiment 3 is the same with experiment 2, but Random Undersampling (RUS) is applied to **handle** imbalanced class.

Experiment 4 is the same with experiment 2, but Random is applied to **handle** imbalanced class.

same with experiment 2, but **SMOTE** is applied to handle imbalanced class.



Experiment Setup

- **Statistical** imputation for all
 - missing values Capping
 - outliers-IQR Normalization Log Transform

Encoding

Experiment 4

Experiment 2

Oversampling (ROS)

Experiment 5

Experiment 5 is the