A STUDY ON HEART DISEASE PREDICTION USING DIFFERENT CLASSIFICATION MODELS BASED ON CROSS VALIDATION METHOD

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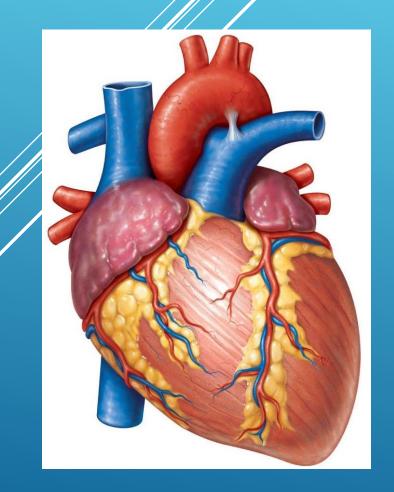
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INTRODUCTION

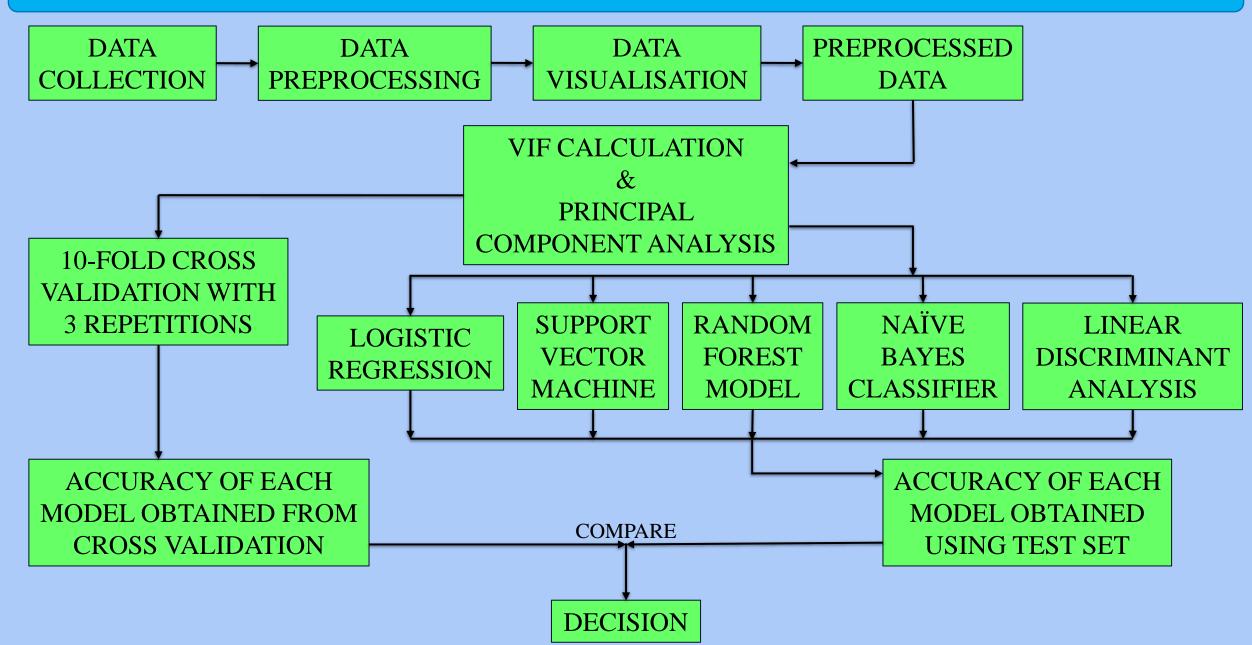
According to World Health Organization (WHO), heart disease is the no. 1 cause of death in world. It is responsible for 16% of total deaths in world [1]. Since 2000, the largest increase in deaths has been for heart disease, rising by more than 2 million to 8.9 million deaths in 2019 [1]. Also in India, heart disease is the leading cause of death. According to Global Burden of Disease, 24.8% of all deaths in India is due to heart disease [2]. Heart disease may happen for various reasons. Most common heart disease is coronary artery disease, which happens due to building up of fatty plaques in arteries (atherosclerosis). Heart disease can show various symptoms like chest pain, suffocation, weakness and many more according to the type of heart disease. It can be prevented by maintaining proper diet, following healthy lifestyle, doing regular exercise etc. Though a great amount of statistical and scientific researches is being done, heart disease continues to be the largest killer of world. By early detection of heart disease and proper treatment, chance of survival of a heart disease patient can be increased.

OBJECTIVES

- > To compare five classification models and find the best model, out of these, in terms of accuracy.
- > To find the importance of variables for classification.

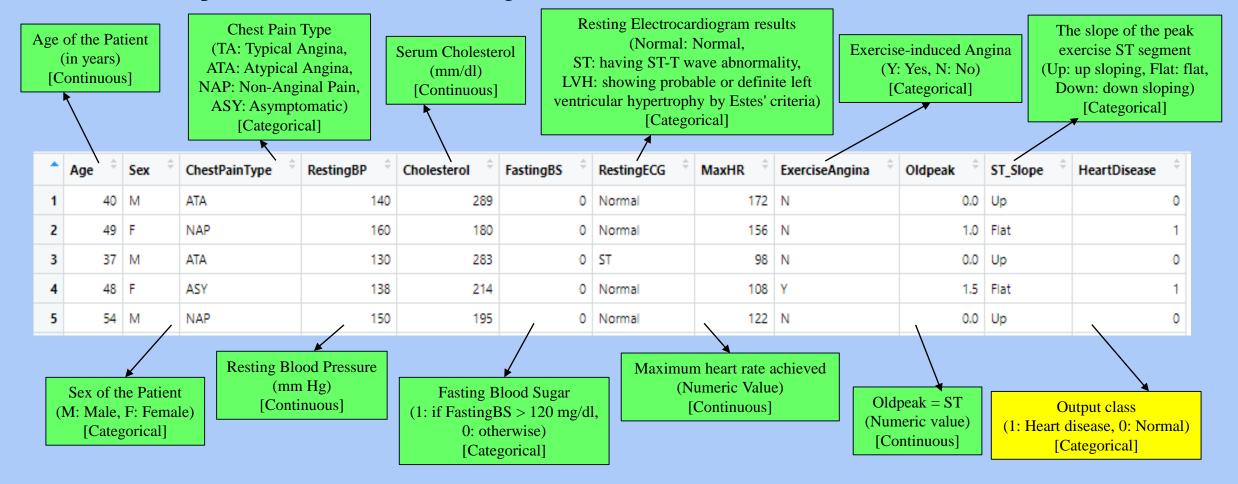
Keywords: - Heart Disease, Logistic Regression model, Support Vector Machine, Random Forest Model, Naïve Bayes classifier, Linear Discriminant Analysis, Cross validation

METHODOLOGY



DATA DESCRIPTION

- ➤ The dataset is downloaded from Kaggle website [3]. This is an open-access dataset.
- The dataset is of 918 observations and contains 11 independent variables and a categorical variable, whether there exists heart disease or not, as target variable.
- Out of 11 independent variables, 6 are categorical variables and rest 5 are continuous variables.

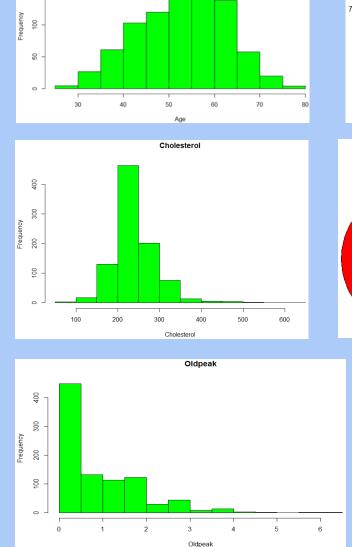


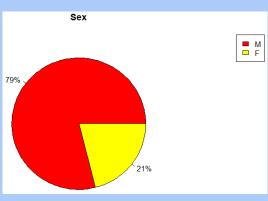
DATA PREPROCESSING

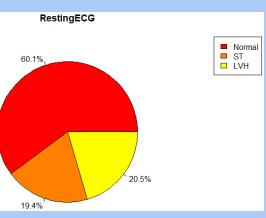
- ➤ There is no missing value in our dataset.
- There are some 0 values in the columns RestingBP and Cholesterol. But Resting Blood Pressure and Serum Cholesterol of a person can never be 0. So, these are bad values. These zeros are replaced with median values of the corresponding columns.
- > There are some negative values in the column Oldpeak. These negative values are converted to positive.
- ➤ It is found that about 77% values of the column FastingBS are 0. So, this column will not impact greatly on classification. So FastingBS column is dropped.
- ➤ Values of some columns are categorical variables. So, we code them into numbers. The changes are shown in the form of a table.

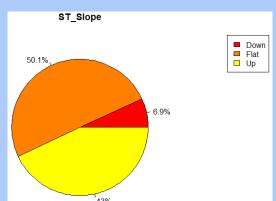
Column Name	Value	Coded Value
Sex	'M'	1
	'F'	2
ChestPainType	'ATA'	1
	'NAP'	2
	'ASY'	3
	'TA'	4
RestingECG	'Normal'	1
	'ST'	2
	'L∨H'	3
ExerciseAngina	'Υ'	1
	'N'	0
ST_Slope	'Down'	-1
	'Flat'	0
	'Up'	1

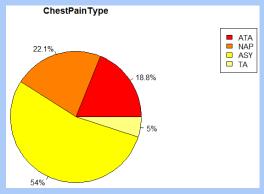
DATA VISUALISATION

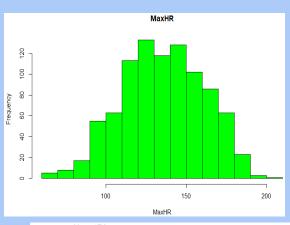


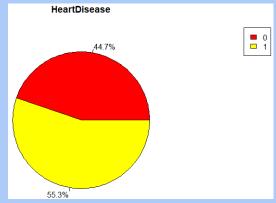


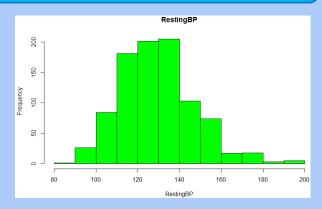


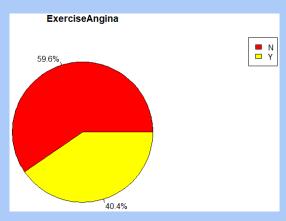












A good interpretation of the dataset can be made using these information and visualisations.

EXPERIMENTAL RESULTS

VIF CALCULATION

Variables	VIF
Age	1.361663
Sex	1.092017
ChestPainType	1.258605
RestingBP	1.100360
Cholesterol	1.038561
RestingECG	1.090604
MaxHR	1.428407
ExerciseAngina	1.455541
Oldpeak	1.539348
ST_Slope	1.622914

We can see that the VIFs are very close to 1. So, there is no significant multicollinearity in the data.

PRINCIPAL COMPONENT ANALYSIS

	Standard Deviation	Proportion of Variance	Cumulative Proportion
PC1	1.829433	0.304260	0.304260
PC2	1.14266	0.11870	0.42295
PC3	1.023245	0.095180	0.518140
PC4	0.9901791	0.0891300	0.6072700
PC5	0.9213231	0.0771700	0.6844400
PC6	0.908279	0.075000	0.759440
PC7	0.8358615	0.0635100	0.8229500
PC8	0.7840512	0.0558900	0.8788400
PC9	0.7166974	0.0467000	0.9255300
PC10	0.6671866	0.0404700	0.9660000
PC11	0.6115684	0.0340000	1.0000000

To explain 95% variance, 10 out of 11 principal components is required. So, no significant dimension reduction is possible. This also indicates non-existence of multicollinearity, which supports the information obtained from the VIF values.

So we are good to fit various classification models.

EXPERIMENTAL RESULTS

We have fitted various model to the train set and predicted the test set. Then 10-fold cross validation is done with 3 repetitions. Then following accuracies are obtained.

ACCURACY OBTAINED USING TEST SET

Model	Accuracy	
Logistic Regression	0.832	
Support Vector Machine	0.832	
Random Forest Model	0.864	
Naïve Bayes Classifier	0.832	
Linear Discriminant Analysis	0.832	

ACCURACY OBTAINED FROM CROSS VALIDATION

	Minimum	1 st Quartile	Median	Mean	3 rd Quartile	Maximum
LR	0.783	0.818	0.847	0.848	0.877	0.924
SVM	0.793	0.835	0.848	0.852	0.877	0.913
RF	0.772	0.848	0.869	0.869	0.891	0.924
NB	0.761	0.826	0.848	0.842	0.859	0.902
LDA	0.783	0.817	0.852	0.849	0.870	0.935

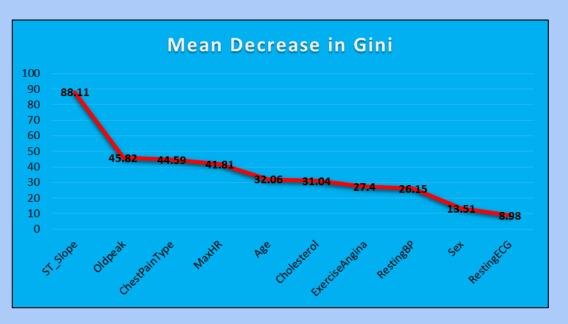
Random Forest model has the highest accuracy of 86.4% when observations of test set is predicted using the fitted models. Resampling also proposes Random Forest as best model with a mean accuracy of 86.9%, which is highest among all models.

EXPERIMENTAL RESULTS

From Random Forest model, variable importance is obtained by of Mean Decrease in Gini coefficient. It is produced simultaneously during training of the model. Mean Decrease in Gini coefficients for each predictor variable obtained from the Random Forest model fitted to the training data is shown below.

VARIABLE IMPORTANCE

Variables	Mean Decrease in Gini
ST_Slope	88.109127
Oldpeak	45.819191
ChestPainType	44.594254
MaxHR	41.805309
Age	32.061294
Cholesterol	31.044686
Exercise Angina	27.399798
RestingBP	26.154423
Sex	13.510075
RestingECG	8.977059



- > ST_Slope has a Mean Decrease in Gini coefficient value of 88.11, which is highest among all predictor variables.
- ➤ Oldpeak, ChestPainType and MaxHR also have close values, 45.82, 44.59 and 41.81 respectively.
- RestingECG has the lowest value of 8.98.

SIGNIFICANCE & CONCLUSION

- So, it can be concluded that the Random Forest model gives the best predictions of existence of heart disease. From the results of 10- fold cross-validation, it can be said that resampling also supports the fact of the Random Forest being the best model out of our experimented models. Rest of the models give more or less similar performance in terms of correctness of prediction.
- ➤ There are some advantages of using Random Forest model for classification. Decision tree is highly biased and it has greater variance. But as Random Forest is collection of multiple decision trees, it has less bias and less variance. Also Random Forest solves the problem of overfitting of Decision tree.
- The slope of the peak exercise ST segment is the most important factor for classification of heart disease. Also, old peak, type of chest pain and maximum heart rate should also be seriously considered. Resting electrocardiogram result and sex contributes least for classification.

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I would like to thank my parents, family members and friends for support in various fields of this project.

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THANK YOU!!