**Cardiovascular Risk Prediction**

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**1.Introduction:**

Heart disease is the major cause of morbidity and mortality globally: it accounts for more deaths annually than any other cause. According to the WHO, an estimated 17.9 million people died from heart disease in 2016, representing 31% of all global deaths. Over three quarters of these deaths took place in low- and middle-income countries.

Of all heart diseases, coronary heart disease (aka heart attack) is by far the most common and the most fatal. In the United States, for example, it is estimated that someone has a heart attack every 40 seconds and about 805,000 Americans have a heart attack every year (CDC 2019).

The silver lining is that heart attacks are highly preventable and simple lifestyle modifications(such as reducing alcohol and tobacco use; eating healthily and exercising) coupled with early treatment greatly improves its prognosis. It is, however, difficult to identify high risk patients because of the multi-factorial nature of several contributory risk factors such as diabetes, high blood pressure, high cholesterol, et cetera. This is where machine learning and data mining come to the rescue.

Doctors and scientists alike have turned to machine learning (ML) techniques to develop screening tools and this is because of their superiority in pattern recognition and classification as compared to other traditional statistical approaches.

In this article, I will be giving you a walk through on the development of a screening tool for predicting whether a patient has 10-year risk of developing coronary heart disease(CHD) using different Machine Learning techniques on the Framingham dataset .

**2.Data and Problem Statement:**

The dataset is from an ongoing cardiovascular study on residents of the town of Framingham, Massachusetts. The classification goal is to predict whether the patient has a 10-year risk of future coronary heart disease (CHD). The dataset provides the patients’ information. It includes over 3,000 records and 15 attributes.

Variables

Each attribute is a potential risk factor. There are both demographic, behavioral, and medical risk factors.

Data Description

Demographic:

• Sex: male or female("M" or "F")

• Age: Age of the patient;(Continuous - Although the recorded ages have been truncated to whole numbers, the concept of age is continuous)

Behavioral

• is\_smoking: whether or not the patient is a current smoker ("YES" or "NO")

• Cigs Per Day: the number of cigarettes that the person smoked on average in one day.(can be considered continuous as one can have any number of cigarettes, even half a cigarette.)

Medical( history)

• BP Meds: whether or not the patient was on blood pressure medication (Nominal) • Prevalent Stroke: whether or not the patient had previously had a stroke (Nominal) • Prevalent Hyp: whether or not the patient was hypertensive (Nominal)

• Diabetes: whether or not the patient had diabetes (Nominal)

Medical(current)

• Tot Chol: total cholesterol level (Continuous)

• Sys BP: systolic blood pressure (Continuous)

• Dia BP: diastolic blood pressure (Continuous)

• BMI: Body Mass Index (Continuous)

• Heart Rate: heart rate (Continuous - In medical research, variables such as heart rate though in fact discrete, yet are considered continuous because of large number of possible values.)

• Glucose: glucose level (Continuous)

Predict variable (desired target)

• 10-year risk of coronary heart disease CHD(binary: “1”, means “Yes”, “0” means “No”) - DV

**3. Steps involved:**

* **Missing values & Outliers Treatment**

Our dataset contain less number of outliers which might tend to disturb our accuracy hence we dropped them at the beginning of our project in order to get a better result.and also treated missing values

* **Exploratory Data Analysis**

After loading the dataset we performed this method by comparing our target variable that is ‘TenYearCHD’ with other independent variables. This process helped us figuring out various aspects and relationships among the target and the independent variables. It gave us a better idea of which feature behaves in which manner compared to the target variable.

* **Feature Selection**

To idenfify the features that have larger contribution towards the outcome

variable,TenYearCHD and selected 8 best features apart fromTenYearCHD

* **Fitting different models**

For modelling we tried various classification algorithms like:

* **Logistic Regressor**
* **K-Nearest Neighbour Classifier**
* **Random Forest Classifier**
* **Decision Tree Classifier**
* **Gradient Boosting Classifier**

**4. Algorithms:**

* **Logistic Regression:**

Logistic regression is basically a supervised classification algorithm. In a classification problem, the target variable(or output), y, can take only discrete values for a given set of features(or inputs), X.

Contrary to popular belief, logistic regression IS a regression model. The model builds a regression model to predict the probability that a given data entry belongs to the category numbered as “1”. Just like Linear regression assumes that the data follows a linear function, Logistic regression models the data using the sigmoid function.

g(z) = \frac{1}{1 + e^-^z}\



Logistic regression becomes a classification technique only when a decision threshold is brought into the picture. The setting of the threshold value is a very important aspect of Logistic regression and is dependent on the classification problem itself.

The decision for the value of the threshold value is majorly affected by the values of precision and recall. Ideally, we want both precision and recall to be 1, but this seldom is the case. In the case of a Precision-Recall tradeoff, we use the following arguments to decide upon the threshold:-

1. Low Precision/High Recall: In applications where we want to reduce the number of false negatives without necessarily reducing the number of false positives, we choose a decision value that has a low value of Precision or a high value of Recall. For example, in a cancer diagnosis application, we do not want any affected patient to be classified as not affected without giving much heed to if the patient is being wrongfully diagnosed with cancer. This is because the absence of cancer can be detected by further medical diseases but the presence of the disease cannot be detected in an already rejected candidate.

2. High Precision/Low Recall: In applications where we want to reduce the number of false positives without necessarily reducing the number of false negatives, we choose a decision value that has a high value of Precision or a low value of Recall. For example, if we are classifying customers whether they will react positively or negatively to a personalized advertisement, we want to be absolutely sure that the customer will react positively to the advertisement because otherwise, a negative reaction can cause a loss of potential sales from the customer

.**4.2 K-Nearest Neighbour Classifier:**

K-Nearest Neighbour is one of the simplest Machine Learning algorithms based on Supervised Learning technique.

K-NN algorithm assumes the similarity between the new case/data and available cases and put the new case into the category that is most similar to the available categories.

K-NN algorithm stores all the available data and classifies a new data point based on the similarity. This means when new data appears then it can be easily classified into a well suite category by using K- NN algorithm.

K-NN algorithm can be used for Regression as well as for Classification but mostly it is used for the Classification problems.

K-NN is a non-parametric algorithm, which means it does not make any assumption on underlying data.

It is also called a lazy learner algorithm because it does not learn from the training set immediately instead it stores the dataset and at the time of classification, it performs an action on the dataset.

KNN algorithm at the training phase just stores the dataset and when it gets new data, then it classifies that data into a category that is much similar to the new data.

Example: Suppose, we have an image of a creature that looks similar to cat and dog, but we want to know either it is a cat or dog. So for this identification, we can use the KNN algorithm, as it works on a similarity measure. Our KNN model will find the similar features of the new data set to the cats and dogs images and based on the most similar features it will put it in either cat or dog category.



K-Nearest Neighbor(KNN) Algorithm for Machine Learning

Why do we need a K-NN Algorithm?

Suppose there are two categories, i.e., Category A and Category B, and we have a new data point x1, so this data point will lie in which of these categories. To solve this type of problem, we need a K-NN algorithm. With the help of K-NN, we can easily identify the category or class of a particular dataset. Consider the below diagram:



* **Random Forest Classifier:**

Random forest, like its name implies, consists of a large number of individual decision trees that operate as an ensemble. Each individual tree in the random forest spits out a class prediction and the class with the most votes becomes our model’s prediction (see figure below).



Visualization of a Random Forest Model Making a Prediction

The fundamental concept behind random forest is a simple but powerful one — the wisdom of crowds. In data science speak, the reason that the random forest model works so well is:

A large number of relatively uncorrelated models (trees) operating as a committee will outperform any of the individual constituent models.

The low correlation between models is the key. Just like how investments with low correlations (like stocks and bonds) come together to form a portfolio that is greater than the sum of its parts, uncorrelated models can produce ensemble predictions that are more accurate than any of the individual predictions. The reason for this wonderful effect is that the trees protect each other from their individual errors (as long as they don’t constantly all err in the same direction). While some trees may be wrong, many other trees will be right, so as a group the trees are able to move in the correct direction. So the prerequisites for random forest to perform well are:

There needs to be some actual signal in our features so that models built using those features do better than random guessing.

The predictions (and therefore the errors) made by the individual trees need to have low correlations with each other.

* **Decision Tree Classifier:**

Decision Tree is a Supervised learning technique that can be used for both classification and Regression problems, but mostly it is preferred for solving Classification problems. It is a tree-structured classifier, where internal nodes represent the features of a dataset, branches represent the decision rules and each leaf node represents the outcome.

In a Decision tree, there are two nodes, which are the Decision Node and Leaf Node. Decision nodes are used to make any decision and have multiple branches, whereas Leaf nodes are the output of those decisions and do not contain any further branches.

The decisions or the test are performed on the basis of features of the given dataset.

It is a graphical representation for getting all the possible solutions to a problem/decision based on given conditions.

It is called a decision tree because, similar to a tree, it starts with the root node, which expands on further branches and constructs a tree-like structure.

In order to build a tree, we use the CART algorithm, which stands for Classification and Regression Tree algorithm.

A decision tree simply asks a question, and based on the answer (Yes/No), it further split the tree into subtrees.

Below diagram explains the general structure of a decision tree:



**Gradient Boosting Classifier:**

Gradient boosting classifiers are a group of machine learning algorithms that combine many weak learning models together to create a strong predictive model. Decision trees are usually used when doing gradient boosting. Gradient boosting models are becoming popular because of their effectiveness at classifying complex datasets

Classification refers to the task of giving a machine learning algorithm features, and having the algorithm put the instances/data points into one of many discrete classes. Classes are categorical in nature, it isn't possible for an instance to be classified as partially one class and partially another. A classic example of a classification task is classifying emails as either "spam" or "not spam" - there's no "a bit spammy" email

Gradient boosting classifiers are specific types of algorithms that are used for classification tasks, as the name suggests.

The idea behind "gradient boosting" is to take a weak hypothesis or weak learning algorithm and make a series of tweaks to it that will improve the strength of the hypothesis/learner. This type of Hypothesis Boosting is based on the idea of Probability Approximately Correct Learning (PAC).

This PAC learning method investigates machine learning problems to interpret how complex they are, and a similar method is applied to Hypothesis Boosting.

In hypothesis boosting, you look at all the observations that the machine learning algorithm is trained on, and you leave only the observations that the machine learning method successfully classified behind, stripping out the other observations. A new weak learner is created and tested on the set of data that was poorly classified, and then just the examples that were successfully classified are kept.

**5.Hyperparameter Tuning for best Classifier:**

Hyperparameters are sets of information that are used to control the way of learning an algorithm. Their definitions impact parameters of the models, seen as a way of learning, change from the new hyperparameters. This set of values affects performance, stability and interpretation of a model. Each algorithm requires a specific hyperparameters grid that can be adjusted according to the business problem. Hyperparameters alter the way a model learns to trigger this training algorithm after parameters to generate outputs

We used Random Search CV for hyperparameter tuning. This also results in cross validation and in our case we divided the dataset into different folds.

**6. Conclusion:**

⦁ Read the file and displayed its columns.​

⦁ Handled missing values, Outliers and Duplicate Data.​

⦁ Calculated basic statistics of the data (count, mean, std, etc), did exploratory analysis and described my observations.​

⦁ Selected columns that will probably be important to predict heart disease.​

⦁ Created training and testing sets (using 60% of the data for the training and reminder for testing) and scaled the data using MinMaxScaler.​

⦁ Built 5 different machine learning models to predict TenYearCHD:​

⦁ Logistic Regression - 65% Accuracy​

⦁ kNN Classification - 93% Accuracy​

⦁ Random Forest Classification - 94% Accuracy​

⦁ Decision Tree Classification - 91% Accuracy​

⦁ Gradient Boosting Classification - 72% Accuracy​

⦁ Hyperparameter tuned the RandomForestClassification – 98%(Recall)​

⦁ Evaluated each model (f1 score, Accuracy, Precision ,Recall and Confusion Matrix) and plotted a graph for the false positive rate and true positive rate for each model.​

⦁ Ensembled the four best models using Stacking technique to further increase the accuracy of the model and achieved an Recall score of 98%​

⦁ Concluded that Ensembling all the four most important models, with Random Forest Classification leading the way, has resulted in a very high accuracy and Recall

Business Use case:

We took Recall as a right parameter because if a doctor gives tablets to patients which are might be harmful to the patients which are not having cardiovascular disease then it might causes some irrelevance that is why we took Recall which enabled not to miss out any patients with coronary heart disease