#### CASE STUDY REPORT

#### **Coronary Heart Disease Classification Problem**

Group No.: 11

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### **Executive Summary:**

Cardiovascular disease is one of the most significant reasons for the number of deaths among all people around the world. The early forecast of cardiovascular ailments can help in compelling high-risk patients to change their risky habits and follow a healthy routine such that any unfavorable events can be prevented.

The goal of this study was to forecast if a patient would get coronary heart disease in a 10-year span or not according to the traits he possesses.

The dataset is publicly accessible on the Kaggle website, and it is from an in progress cardiovascular examination, on the inhabitants of the town of Framingham, Massachusetts. The dataset has over 4000 records and 15 features or attributes. Each attribute is a potential risk factor. There are both demographic, behavioral, and medical risk factors.

While exploring our data, we found that the total number of rows with missing values were 582 which is only 13 percent of the entire dataset, so we excluded them.

As we have a classification problem, we have applied kNN, Naive Bayes, Random Forest, Logistic Regression, Neural Net, Linear Discriminant Analysis and Support Vector Machine and then evaluated the performances of each model.

We formulated 7 different supervised learning models and created confusion matrices, Lift Charts/ ROC curves and found out accuracies for all of them. Men appear to be more vulnerable to coronary illness than Women. An expansion in age, number of cigarettes smoked every day and systolic Blood Pressure additionally show expanded chances of having coronary illness.

Out of all the 7 supervised learning models, our KNN model is most accurate with an accuracy of 86%.

### I. Background and Introduction

Cardiovascular diseases are probably the most significant reason for the number of deaths among all people around the world. The prediction of cardiovascular diseases is viewed as one of the most significant subjects in the field of data analytics under the healthcare domain. World Health Organization has evaluated 12 million demises happening around the world, consistently because of heart related problems. Around 610,000 individuals fall at the hands of coronary illness in the United States each year—that is 1 in every four demises; that is, one person dies every 37 seconds due to it.

Coronary illness portrays a scope of conditions that influences the heart. Sicknesses under the coronary illness umbrella comprises of blood vessels ailments, heart rhythm issues (arrhythmias); and heart absconds you're brought into the world with (congenital heart defects), etc.

The expression "coronary illness" is frequently utilized conversely with the expression "cardiovascular diseases." Cardiovascular sickness by and large alludes to conditions that include restricted or blocked vessels that can prompt a respiratory failure, chest torment (angina) or stroke. Other heart conditions, for example, those that influence your heart's muscle, valves or rhythm, likewise are viewed as types of coronary illness. Numerous types of coronary illness can be prevented or treated by following and implementing solid & healthy lifestyle decisions. The early forecast of cardiovascular ailments can help in compelling high-risk patients to change their risky habits and follow a healthy routine such that any unfavorable events can be prevented.

Our study means to pinpoint the most pertinent/hazardous elements involved in the cardiovascular diseases and to anticipate the general hazard. The classification goal is to forecast if a patient would get coronary heart disease in a 10-year span.

## II. Data Exploration and Visualization

We utilized different visualization techniques to understand data. We checked the multi-collinearity for linear models.

In the end, the model is intended to pinpoint the most relevant/risk factors of heart disease as well as predict the overall risk. It's a binary classification problem.

Predictors are Age, Glucose level, Gender, Education, Current Smoker, Cigarette per day, Blood Pressure Medications, Prevalent Stroke, Prevalent Hypertension, Diabetes, Total Cholesterol, Systolic Blood Pressure, Diastolic Blood Pressure, Body Mass Index & Heart rate. The Outcome variable is a binary variable, the 10-year risk of coronary heart disease CHD with the class either Yes or No.

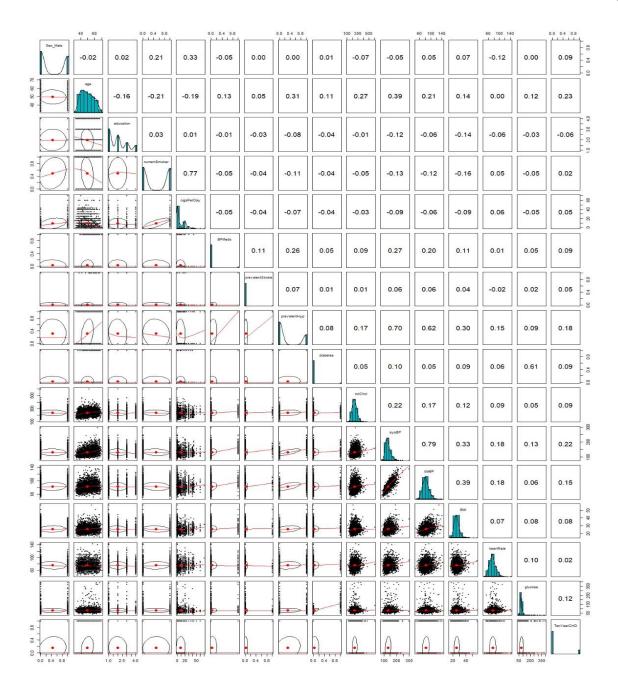


Fig.1 Correlation Plot

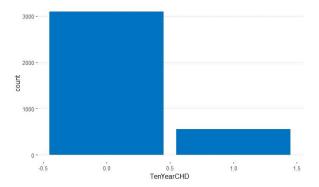


Fig.2 Count plot

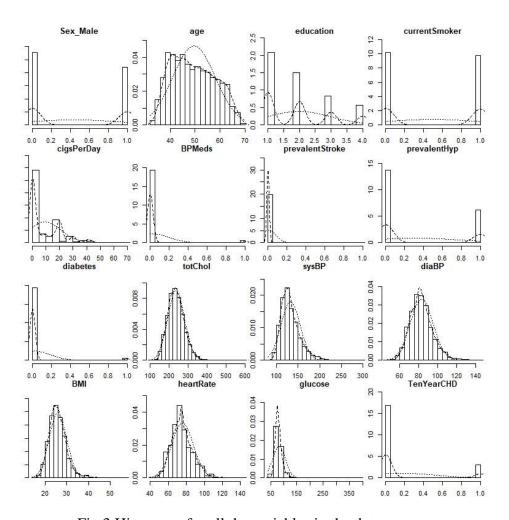


Fig.3 Histogram for all the variables in the data set.

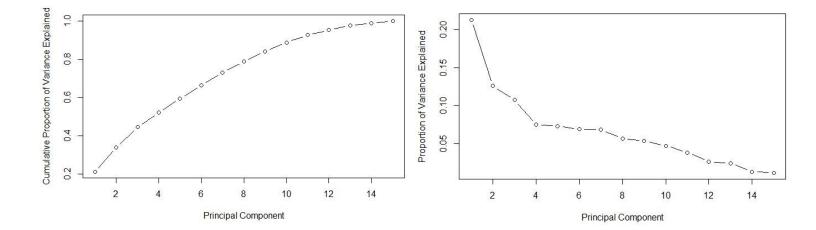
Fig.2 shows that our data has \_ patients who will not have heart disease while \_ patients will have heart disease in the next 10 years.

Fig.3 shows the distribution of individual features in our dataset.

# III. Data Preparation and Preprocessing

We had checked for missing or NA values in the dataset & descriptive statistic of each variable to see if there were any outliers. We eliminated the rows which had missing values because if we had imputed the values here, I would create a bias in our features Total number of rows with missing values was 582. Since it was only 13.72 percent of the entire dataset the rows with missing values were excluded.

```
cigsPerDay
                                                                                        RPMeds
   Sex Male
                                   education
                                                 currentSmoker
Min.
       :0.0000
                 Min.
                        :32.00
                                 Min.
                                        :1.00
                                                 Min.
                                                        :0.0000
                                                                  Min.
                                                                         : 0.000
                                                                                    Min.
                                                                                           :0.00000
1st Qu.:0.0000
                 1st Qu.:42.00
                                                 1st Qu.:0.0000
                                                                  1st Qu.: 0.000
                                                                                    1st Qu.:0.00000
                                 1st Qu.:1.00
Median :0.0000
                 Median :49.00
                                                 Median :0.0000
                                                                  Median : 0.000
                                                                                    Median :0.00000
                                 Median :2.00
      :0.4437
                        :49.55
                                        :1.98
                                                 Mean
                                                        :0.4891
                                                                         : 9.025
                                                                                           :0.03034
                 Mean
                                 Mean
                                                                  Mean
                                                                                    Mean
Mean
3rd Qu.:1.0000
                 3rd Qu.:56.00
                                 3rd Qu.:3.00
                                                 3rd Qu.:1.0000
                                                                  3rd Qu.: 20.000
                                                                                    3rd Qu.:0.00000
Max.
       :1.0000
                 Max.
                       :70.00
                                 Max.
                                         :4.00
                                                 Max.
                                                        :1.0000
                                                                  Max.
                                                                         :70.000
                                                                                    Max.
                                                                                           :1.00000
prevalentStroke
                    prevalentHyp
                                        diabetes
                                                          totChol
                                                                            sysBP
                                                                                            diaBP
Min.
       :0.000000
                   Min.
                         :0.0000
                                    Min.
                                           :0.00000
                                                       Min.
                                                              :113.0
                                                                       Min.
                                                                              : 83.5
                                                                                        Min.
                                                                                              : 48.00
                   1st Qu.:0.0000
                                    1st Qu.:0.00000
1st Ou.:0.000000
                                                       1st Ou.: 206.0
                                                                       1st Qu.:117.0
                                                                                       1st Qu.: 75.00
Median :0.000000
                                                       Median :234.0
                   Median :0.0000
                                    Median :0.00000
                                                                       Median :128.0
                                                                                        Median : 82.00
      :0.005741
Mean
                   Mean
                         :0.3116
                                    Mean
                                           :0.02706
                                                       Mean
                                                             :236.8
                                                                       Mean
                                                                              :132.4
                                                                                        Mean
                                                                                              : 82.92
3rd Qu.:0.000000
                   3rd Qu.:1.0000
                                    3rd Qu.:0.00000
                                                       3rd Qu.: 263.0
                                                                       3rd Qu.:143.9
                                                                                        3rd Qu.: 90.00
      :1.000000
                   Max. :1.0000
                                    Max.
                                           :1.00000
                                                       Max.
                                                              :600.0
                                                                       Max.
                                                                              :295.0
                                                                                       Max.
                                                                                              :142.50
Max.
    RMT
                  heartRate
                                                     TenYearCHD
                                    glucose
                                                         :0.0000
Min.
      :15.54
                Min.
                      : 44.00
                                 Min.
                                        : 40.00
                                                   Min.
1st Qu.:23.08
                1st Qu.: 68.00
                                 1st Qu.: 71.00
                                                   1st Qu.:0.0000
Median :25.38
                Median : 75.00
                                 Median : 78.00
                                                   Median :0.0000
                                 Mean : 81.85
      :25.78
                      : 75.73
                                                         :0.1523
Mean
                Mean
                                                   Mean
3rd Qu.:28.04
                3rd Qu.: 82.00
                                 3rd Qu.: 87.00
                                                   3rd Qu.:0.0000
Max.
      :56.80
                Max.
                       :143.00
                                 Max.
                                        :394.00
                                                   Max.
                                                          :1.0000
```



We performed Principal Component Analysis. PCA looks for properties that show as much variation across classes as possible to build the principal component space. The algorithm uses the concepts of variance matrix, covariance matrix, eigenvector and eigenvalues pairs to perform PCA, providing a set of eigenvectors and its respectively eigenvalues as a result. It is very simple; the eigenvectors represent the new set of axes of the principal component space and the eigenvalues carry the information of quantity of variance that each eigenvector have. So, in order to reduce the dimension of the dataset we are going to choose those eigenvectors that have more variance and discard those with less variance.

#### IV. Data Mining Techniques and Implementation

As we have a classification problem, we have applied kNN, Naive Bayes, Random Forest, Logistic Regression, Neural Net, Linear Discriminant Analysis and Support Vector Machine. We will compare the models to find the best fit model for our problem.

#### 1. K-Nearest Neighbors

RMSE was used to select the optimal model using the smallest value. The final value used for the model was k = 43. The model is classified as level 0 thus the patient will not have coronary heart disease in 10 years.

```
[1] "2205" "4036" "1554"
[1] 0
attr(,"nn.index")
      [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1] [,1]
[1,] 3157 3260 2654 228 3402 1886 3473 1348 3220 2690 2125
                                                               667 2117 2308
                                                                                 409
                                                                                       564
     [,20] [,21] [,22] [,23] [,24] [,25] [,26] [,27] [,28] [,29] [,30] [,31] [,32] [,33] [,34] [,35] [,36] [,37]
[1,] 1749 3102 2967
                        406 1604 2295
                                          272 1636 2746 3276 2369 1127 3467 1869
[,38] [,39] [,40] [,41] [,42] [,43]
[1,] 1345 3048 2451 1012 2205 3627
attr(,"nn.dist")
         [,1]
                  [,2]
                           [,3]
                                    [,4]
                                             [,5]
                                                     [,6]
                                                              [,7]
                                                                       [,8]
                                                                               [,9]
                                                                                      [,10]
                                                                                                [,11]
                                                                                                         [,12]
[1,] 6.977703 6.979081 7.248525 7.687337 7.827544 8.53705 8.629625 8.782091 8.898135 8.91105 8.911886 8.920804
                                                    [,18]
                                                                                      [,22]
                 [,14]
                          [,15]
                                  [,16]
                                           [,17]
                                                            [,19]
                                                                      [,20]
                                                                              [,21]
                                                                                                [,23]
        [,13]
[1,] 8.930859 9.203902 9.204123 9.42979 9.457254 9.562825 9.691679 9.932137 10.10613 15.3833 15.51666 15.76507
        [,25]
                 [,26]
                          [,27]
                                   [,28]
                                           [,29]
                                                    [,30]
                                                             [,31]
                                                                     [,32]
                                                                              [,33]
                                                                                       [,34]
                                                                                                [,35]
[1,] 15.80075 16.12166 16.12229 16.15352 16.16135 16.19502 16.19532 16.2192 16.22703 16.23609 16.25262 16.27233
               [,38]
                        [,39]
                                 [,40]
                                                   [,42]
                                                            [,43]
        [,37]
                                          [,41]
[1,] 16.27513 16.2839 16.30901 16.31181 16.31598 16.33695 16.33944
Levels: 0
```

### 2. Naïve Bayes:-

```
Naive Bayes Classifier for Discrete Predictors
     naiveBayes.default(x = X, y = Y, laplace = laplace)
     A-priori probabilities:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prevalentStroke
0.8463993 0.1536007
Conditional probabilities:
               [,1] [,2]
0 0.4221863 0.4940410
1 0.5489614 0.4983369
               33 34 35 36 37 38 39 40 41 0 0.001077006 0.005385030 0.009593053 0.028540657 0.022078621 0.034464159 0.046849758 0.055542811 0.044157243 1 0.000000000 0.0000000000 0.005934718 0.00267539 0.008902077 0.017804154 0.008902077 0.023738872 0.020771513
                 ege 4 43 44 45 46 47 48 49 0.04457472752 0.042541734 0.036612201 0.048465267 0.032310178 0.037165704 0.020549657 0.035310178 0.037165704 0.020549657 0.035310178 0.037165704 0.020549657 0.0353541195 1 0.0229573591 0.041836795 0.020771513 0.023738872 0.029673591 0.023738872 0.04183680 0.041543627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          159 160 151 152 163 164 155 166 167 3 0.002154012 0.002352131 0.00225154012 0.004308024 0.002154012 0.002508000 0.000200000 0.000200000 0.002267359 0.00200000 0.000200000 0.002267359 0.002267359 0.002000000 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267359 0.002267250 0.002267359 0.002267359 0.002267250 0.002267359 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.002267250 0.0
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            69 76
0 0.001615509 0.000538503
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               1 2 3 4
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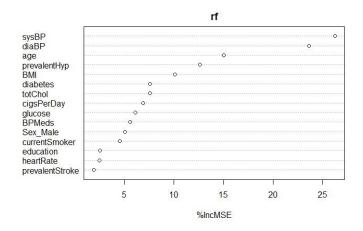
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 180 1 180.5 180.5 181.5 182.5 182.5 182.5 183 184 185 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 185.5 
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              giucose 67 68 970 71 72 73 74 0 0.01386107 0.025340657 0.025340645 0.019924610 0.043080273 0.021540118 0.02530639 0.0408387722 0.036618201 0 0.043080273 0.02534059 0.02530639 0.040838775 0.025340575 0.014836795 0.011869436 0.035608309 0.011869436 0.038575668 0.017804742 0.025073591
              glucose 7 76 77 78 79 80 81 82 83 83 1 0.05367272752 0.032848681 0.04516740 0.032673591 0.027463651 0.038772273 0.032848681 0.04517360 0.038373710 0.027463651 0.038772213 0.016155089 0.025309639 0.037156704 1 0.065379822 0.041543027 0.044510386 0.038575668 0.022738872 0.020771513 0.011869436 0.017804154 0.0594451040 glucose 84
              85 86 87 88 89 90 91 92 0 0.031771675 0.030156166 0.021081616 0.02261763 0.016155899 0.00592333 0.020463113 0.0080877544 0.007000532 0 0.02670231 0.025967391 0.014838795 0.038375668 0.026771513 0.005934718 0.0273738872 0.0025967399 0.0080922877
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206
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.005934718
394
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.005934718

#### 3. Random Forest:-

Percentage of predicted classifications correct 84.6994% The model is classified as level 0 thus the patient will not have coronary heart disease in 10 years.



```
Confusion Matrix and Statistics
          Reference
             0
Prediction
        0 1217
                212
           18
              Accuracy : 0.8429
                 95% CI: (0.8232, 0.8612)
   No Information Rate: 0.8436
   P-Value [Acc > NIR] : 0.5462
                  Kappa : 0.0911
Mcnemar's Test P-Value : <2e-16
            Sensitivity: 0.98543
            Specificity: 0.07424
         Pos Pred Value : 0.85164
        Neg Pred Value : 0.48571
             Prevalence: 0.84358
        Detection Rate : 0.83128
   Detection Prevalence : 0.97609
      Balanced Accuracy: 0.52983
       'Positive' Class: 0
```

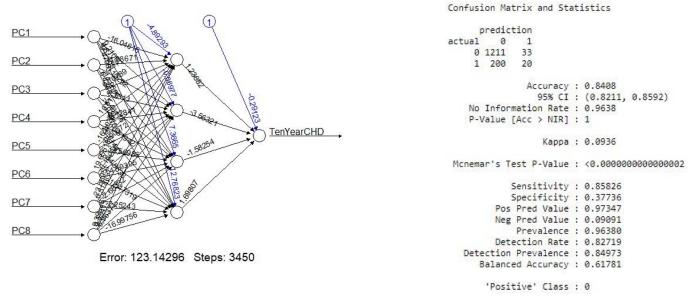
### 4. Logistic Regression:-

For logistic regression algorithm, we found the accuracy to be 83.87%.

```
Call:
glm(formula = as.factor(TenYearCHD) ~ ., family = "binomial",
    data = train.df)
Deviance Residuals:
Min 1Q Median 3Q
-1.6873 -0.5937 -0.4265 -0.2882
                                        2.7505
Coefficients:
                  Estimate Std. Error z value
                                                              Pr(>|z|)
(Intercept)
                 -8.401462
                              0.930109
                                         -9.033 < 0.00000000000000000 ***
Sex_Male
                  0.436716
                              0.139550
                                          3,129
                                                              0.001751 **
                  0.057447
                              0.008656
                                                       0.00000000000321
age
                                          6.636
education
                 -0.141695
                              0.065933
                                          -2.149
                                                              0.031629
                  0.168800
                              0.198921
                                          0.849
currentSmoker
                                                              0.396117
cigsPerDay
                              0.007818
                  0.015696
                                          2.008
BPMeds
                 -0.017661
                              0.301117
                                         -0.059
                                                              0.953231
prevalentStroke
                 1.675153
                              0.815762
                                                              0.040026
                                          2.053
prevalentHyp
                  0.016324
                              0.174708
                                          0.093
                                                              0.925557
diabetes
                  0.230871
                              0.377203
                                          0.612
                                                              0.540498
                  0.003407
                              0.001408
                                          2.420
totChol
                                                              0.015535
                              0.004893
0.008306
sysBP
                  0.018204
                                          3.720
                                                              0.000199 ***
diaBP
                  0.002043
                                          0.246
                                                              0.805748
BMI
                  0.001437
                              0.016413
                                          0.088
                                                              0.930215
heartRate
                 -0.007146
                              0.005553
                                         -1.287
                                                              0.198117
                  0.006026
                              0.002818
                                                              0.032472 *
                                          2.139
glucose
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 1875.2 on 2193 degrees of freedom
Residual deviance: 1647.4 on 2178 degrees of freedom AIC: 1679.4
Number of Fisher Scoring iterations: 5
Min. 1st Qu. Median Mean 3rd Qu. Max.
0.01372 0.06422 0.11638 0.15269 0.19968 0.90813
0.1348806 0.2515136
    FALSE TRUE
  0 1840
            19
      301
```

#### 5. Neural Net:-

The model is classified as level 0 thus the patient will not have coronary heart disease in 10 years.



### 6. Linear Discriminant Analysis:-

```
lda(TenYearCHD ~ ., data = train.df, na.action = "na.omit")
Prior probabilities of groups:
0.8473108 0.1526892
Group means:
Sex_Male age education currentSmoker cigsPerDay BPMeds prevalentStroke prevalentHyp diabetes 0 0.4410974 48.79828 1.997848 0.4862829 8.844002 0.02689618 0.001613771 0.2974718 0.02313072 1 0.5313433 54.32537 1.740299 0.5164179 10.447761 0.06865672 0.011940299 0.5223881 0.07761194
totChol sysBP diaBP BMI heartRate glucose 0 235.9252 130.752 82.38811 25.78779 75.41151 80.64497 1 249.3642 145.806 88.06119 26.65618 76.03582 89.57313
Coefficients of linear discriminants:
Sex_Male
                      0.385006951
                      0.056147895
age
education
                      -0.141663379
currentSmoker 0.173434582
cigsPerDay
                       0.017049104
BPMeds
                       0.162248582
prevalentStroke 3.024922180
prevalentHyp -0.027685808
                      0.482156884
diabetes
                      0.002575647
totChol
sysBP
                      0.026878585
diaBP
                      -0.003697390
BMI
                      -0.007378632
heartRate
                     -0.007452305
glucose
0 1
1420 44
                      0.009190746
```

## 7. Support Vector Machine:-

The model is classified as level 0 thus the patient will not have coronary heart disease in 10 years.

```
Support Vector Machines with Linear Kernel

2561 samples
15 predictor
2 classes: '0', '1'

Pre-processing: centered (15), scaled (15)
Resampling: Cross-Validated (10 fold, repeated 3 times)
Summary of sample sizes: 2305, 2304, 2306, 2305, 2305, 2305, ...
Resampling results:

Accuracy Kappa
0.8508412 0

Tuning parameter 'C' was held constant at a value of 1
```

#### V. Performance Evaluation

Evaluating our machine learning algorithm is an essential part of any project. Most of the times we use classification accuracy to measure the performance of our model, however it is not enough to truly judge our model. In this project, we have made use of confusion matrices, ROC curves & Lift charts to evaluate different models.

Classification Accuracy is what we usually mean, when we use the term accuracy. It is the ratio of number of correct predictions to the total number of input samples.

Confusion Matrix as the name suggests gives us a matrix as output and describes the complete performance of the model.

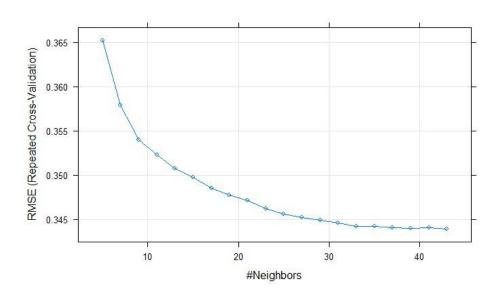
## 1. K-Nearest Neighbors: -

The confusion matrix is created: -

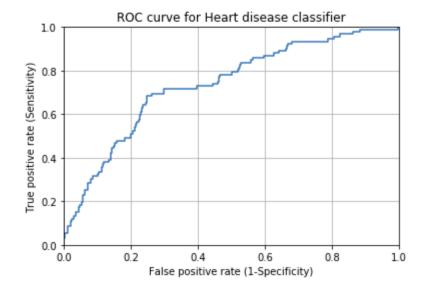
knn.pred	0	1
0	1257	204
1	1	2
Accuracy		0.86
Precision	n:	0.999
Recall:		0.86
F-measure	2:	0.925

k	RMSE	Rsquared	MAE
5	0.3651653	0.03613560	0.2229062
7	0.3579119	0.04043436	0.2237255
9	0.3539518	0.04243793	0.2242940
11	0.3523180	0.04144572	0.2247124
13	0.3507094	0.04266692	0.2252169
15	0.3497306	0.04248450	0.2257110
17	0.3485067	0.04454184	0.2258991
19	0.3477617	0.04582425	0.2260211
21	0.3471409	0.04663127	0.2261944
23	0.3462341	0.04932322	0.2263275
25	0.3456145	0.05075775	0.2261673
27	0.3452036	0.05170426	0.2262544
29	0.3448638	0.05272717	0.2262164
31	0.3445690	0.05398300	0.2262074
33	0.3441698	0.05567272	0.2262103
35	0.3441968	0.05547592	0.2262222
37	0.3440556	0.05621621	0.2262110
39	0.3439878	0.05617782	0.2261798
41	0.3440141	0.05575533	0.2263301
43	0.3439350	0.05621896	0.2263700

RMSE was used to select the optimal model using the smallest value. The final value used for the model was  $k\,=\,43$ .

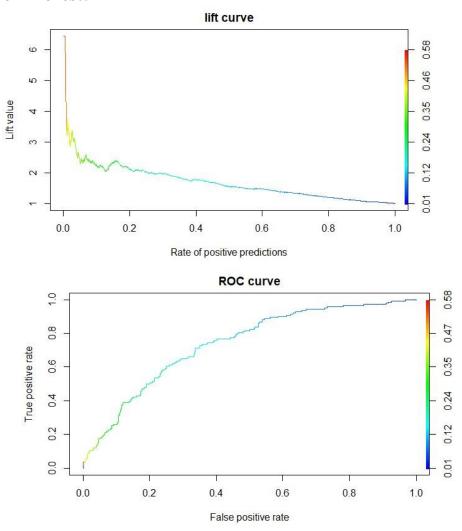


# 2. Naïve Bayes:-

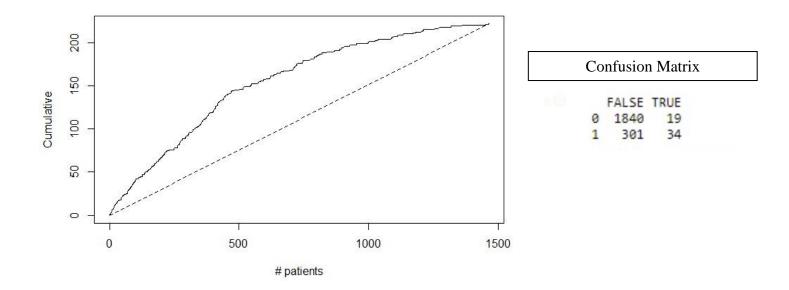


Confusion Matrix & Statistics prediction actual 0 1 0 658 1 1 88 4

# 3. Random Forest:-



# 4. Logistic Regression:-



#### 5. Neural Net:-

Confusion Matrix and Statistics

prediction 1 actual 0 33 0 1211 1 200 20

Accuracy: 0.8408

95% CI: (0.8211, 0.8592)

No Information Rate: 0.9638 P-Value [Acc > NIR] : 1

Kappa: 0.0936

Mcnemar's Test P-Value : <0.00000000000000002

Sensitivity: 0.85826 Specificity: 0.37736 Pos Pred Value : 0.97347 Neg Pred Value : 0.09091 Prevalence: 0.96380

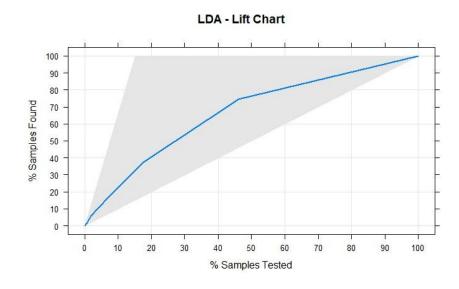
Detection Rate: 0.82719 Detection Prevalence: 0.84973 Balanced Accuracy: 0.61781

'Positive' Class: 0

# 6. Linear Discriminant Analysis:-

Confusion Matrix and Statistics

0 1 0 1217 203 25 19 Accuracy: 0.8443 95% CI: (0.8247, 0.8625) No Information Rate : 0.8484 P-Value [Acc > NIR] : 0.6844 Kappa : 0.0976 Mcnemar's Test P-Value : <0.00000000000000000 Sensitivity: 0.08559 Specificity: 0.97987 Pos Pred Value : 0.43182 Neg Pred Value: 0.85704 Prevalence: 0.15164 Detection Rate: 0.01298 Detection Prevalence : 0.03005



'Positive' Class : 1

Balanced Accuracy: 0.53273

## 7. Support Vector Machine:-

Confusion Matrix and Statistics

test\_pred 0 1 0 922 175 1 0 0

Accuracy: 0.8405

95% CI : (0.8174, 0.8617)

No Information Rate: 0.8405

P-Value [Acc > NIR] : 0.5202

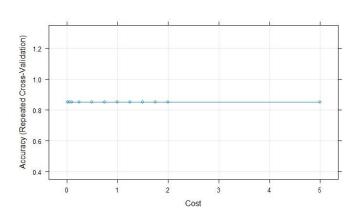
Kappa: 0

Mcnemar's Test P-Value : <0.000000000000000002

Sensitivity: 1.0000
Specificity: 0.0000
Pos Pred Value: 0.8405
Neg Pred Value: NaN
Prevalence: 0.8405
Detection Rate: 0.8405
Detection Prevalence: 1.0000

Balanced Accuracy : 0.5000

'Positive' Class : 0



#### VI. Discussion and Recommendation

We formulated 7 different supervised learning models and created confusion matrices, Lift Charts/ ROC curves and found out accuracies for all of them. Out of all the 7 supervised learning models, our KNN model is most accurate with an accuracy of 86%.

- Men appear to be more vulnerable to coronary illness than Women. An expansion in age, number of cigarettes smoked every day and systolic Blood Pressure additionally show expanded chances of having coronary illness.
- Total cholesterol shows no huge change in the chances of getting coronary heart disease. This could be because of the presence of 'good cholesterol(HDL) while the total cholesterol was calculated. Glucose also causes a truly irrelevant change in the chances of getting coronary heart disease.
- Overall model could be improved if more amount of data is available.

### VII. Summary

The goal of our study was to forecast if a patient would get coronary heart disease in a 10-year span or not according to the traits he possesses. The early forecast of cardiovascular ailments can help in compelling high-risk patients to change their risky habits and follow a healthy routine such that any unfavorable events can be prevented. For our classification problem, we have applied kNN, Naive Bayes, Random Forest, Logistic Regression, Neural Net, Linear Discriminant Analysis and Support Vector Machine and then evaluated the performances of each model.

Out of all the 7 supervised learning models, our KNN model is most accurate with an accuracy of 86%. Changes in features like age, number of cigarettes smoked every day and systolic Blood Pressure have a directly proportional relationship with the 10 year coronary heart disease. Changes in the level total cholesterol and glucose level do not cause much changes in the chances of getting coronary heart disease.

### Appendix: R Code for use case study

```
library(stats)
heart <- read.csv("D:/Data Mining/Data Mining - Group 11/Project/framingham.csv", na.strings = "",
stringsAsFactors = FALSE)
dim(heart)
str(heart)
heart[heart == "NA"] <- NA
heart update <- na.omit(heart)
i < c(3,5,6,10,13,14,15)
                                            # Specify columns you want to change
#We can now use the apply function to change columns 2, 3, 5, 6, 10, 13, 14, and 15 to numeric:
heart_update[, i] <- apply(heart_update[, i], 2,
                                                     # Specify own function within apply
            function(x) as.numeric(as.character(x)))
#Let's check the classes of the variables of our data frame:
sapply(heart_update, class)
                                          # Get classes of all columns
names(heart_update)[1] <- "Sex_Male"
summary(heart_update)
library(plyr)
library(psych)
multi.hist(heart_update[,sapply(heart_update, is.numeric)])
library(ggplot2)
library(ggpubr)
theme_set(theme_pubr())
ggplot(heart_update, aes(TenYearCHD)) +
 geom_bar(fill = "#0073C2FF") +
 theme_pubclean()
library(psych)
pairs.panels(heart update,
       method = "pearson", # correlation method
       hist.col = "#00AFBB",
        density = TRUE, # show density plots
        ellipses = TRUE # show correlation ellipses
        )
#PCA
set.seed(112)
index = sample(1:nrow(heart update), nrow(heart update) * 0.6, replace = FALSE)
trainset = heart_update[index, ]
test = heart update[-index, ]
testset = test[, 1:15]
pca_trainset = trainset[, 1:15]
pca testset = testset
pca = prcomp(pca trainset, scale = T)
# variance
pr var = (pca\$sdev)^2
# % of variance
prop_varex = pr_var / sum( pr_var )
# Plot
plot( prop_varex, xlab = "Principal Component",
          ylab = "Proportion of Variance Explained", type = "b")
plot(cumsum( prop_varex ), xlab = "Principal Component",
                 ylab = "Cumulative Proportion of Variance Explained", type = "b")
```

```
# Creating a new dataset
train = data.frame(TenYearCHD = trainset$TenYearCHD, pca$x)
t = as.data.frame(predict(pca, newdata = pca testset))
new_trainset = train[, 1:9]
new testset = t[.1:8]
# Build the neural network (NN)
library( neuralnet )
n = names( new trainset )
f = as.formula( paste( "TenYearCHD ~", paste( n[!n %in% "TenYearCHD" ], collapse = "+" ) ) )
nn = neuralnet(f, new_trainset, hidden = 4, linear.output = FALSE, threshold=0.01)
# Plot the NN
plot(nn, rep = "best")
# Test the resulting output
nn.results = compute(nn, new testset)
# Results
results = data.frame( actual = test$TenYearCHD,
             prediction = round(nn.results$net.result))
# Confusion Matrix
library(caret)
t = table(results)
print(confusionMatrix(t))
Predict=compute(nn,new_testset)
prob.result <- Predict$net.result</pre>
prob <- Predict$net.result</pre>
# Model 1: kNN
set.seed(743)
train.index <- sample(row.names(heart update), 0.6 * dim(heart update))
valid.index <- setdiff(row.names(heart_update), train.index)</pre>
train.df <- heart_update[train.index, ]</pre>
valid.df <- heart_update[valid.index, ]</pre>
# new patient
new.df <- data.frame(Sex Male = 1, age = 52, education = 2, currentSmoker = 1, cigsPerDay = 20,
BPMeds = 1, prevalentStroke = 1, prevalentHyp = 1, diabetes = 1, totChol = 200, sysBP = 145, diaBP =
100, BMI = 28, heartRate = 80, glucose = 70)
# initialize normalized training, validation data, complete data frames to originals
train.norm.df <- train.df
valid.norm.df <- valid.df
heart.norm.df <- heart_update
# use preProcess() from the caret package to normalize variables.
library(caret)
norm.values <- preProcess(train.df[, 1:15], method=c("center", "scale"))
train.norm.df[, 1:15] <- predict(norm.values, train.df[, 1:15])
valid.norm.df[, 1:15] <- predict(norm.values, valid.df[, 1:15])
heart.norm.df[, 1:15] <- predict(norm.values, heart_update[, 1:15])
new.norm.df <- predict(norm.values, new.df)</pre>
# use knn() to compute knn.
# knn() is available in library FNN (provides a list of the nearest neighbors)
# and library class (allows a numerical output variable).
library(FNN)
nn < knn(train = train.norm.df[, 1:15], test = new.norm.df, cl = train.norm.df[, 16], k = 3)
```

```
row.names(train.df)[attr(nn, "nn.index")]
nn
library(caret)
# initialize a data frame with two columns: k, and accuracy.
accuracy.df \leftarrow data.frame(k = seq(1, 45, 1), accuracy = rep(0, 45))
# compute knn for different k on validation.
valid.norm.df$TenYearCHD <- as.factor(valid.norm.df$TenYearCHD)
for(i in 1:45) {
 knn.pred <- knn(train.norm.df[, 1:15], valid.norm.df[, 1:15],
           cl = train.norm.df[, 16], k = i)
accuracy.df[i, 2] <- confusionMatrix(knn.pred, valid.norm.df[, 16])$overall[1]
accuracy.df
knn.pred.new <- knn(heart.norm.df[, 1:15], new.norm.df,
cl = heart.norm.df[, 16], k = 43)
row.names(train.df)[attr(nn, "nn.index")]
knn.pred.new
xtab = table(knn.pred, valid.norm.df[, 16])
print(xtab)
accuracy = sum(knn.pred == valid.norm.df[, 16])/length(valid.norm.df[, 16])
precision = xtab[1,1]/sum(xtab[,1])
recall = xtab[1,1]/sum(xtab[1,])
f = 2 * (precision * recall) / (precision + recall)
cat(paste("Accuracy:\t", format(accuracy, digits = 3), "\n",sep=" "))
cat(paste("Precision:\t", format(precision, digits = 3), "\n",sep=" "))
cat(paste("Recall:\t\t", format(recall, digits = 3), "\n", sep=" "))
cat(paste("F-measure:\t", format(f, digits = 3), "\n",sep=" "))
library(ISLR)
library(caret)
set.seed(300)
#Spliting data as training and test set. Using createDataPartition() function from caret
indxTrain <- createDataPartition(y = heart update$TenYearCHD, p = 0.75,list = FALSE)
training <- heart update[indxTrain.]
testing <- heart_update[-indxTrain,]
#Checking distibution in origanl data and partitioned data
prop.table(table(training$TenYearCHD)) * 100
prop.table(table(testing$TenYearCHD)) * 100
prop.table(table(heart update$TenYearCHD)) * 100
trainX <- training[,names(training) != "TenYearCHD"]</pre>
preProcValues <- preProcess(x = trainX,method = c("center", "scale"))</pre>
preProcValues
set.seed(400)
ctrl <- trainControl(method="repeatedcv",repeats = 3) #,classProbs=TRUE,summaryFunction =
twoClassSummary)
knnFit <- train(TenYearCHD ~ ., data = training, method = "knn", trControl = ctrl, preProcess =
c("center", "scale"), tuneLength = 20)
#Output of kNN fit
knnFit
#Plotting yields Number of Neighbours Vs accuracy (based on repeated cross validation)
plot(knnFit)
knnPredict <- predict(knnFit,newdata = testing)
#Get the confusion matrix to see accuracy value and other parameter values
```

```
#confusionMatrix(knnPredict, testing$TenYearCHD)
mean(knnPredict == testing$TenYearCHD)
#Now verifying 2 class summary function
     <- trainControl(method="repeatedcv",repeats</p>
                                                                3,classProbs=TRUE,summaryFunction
twoClassSummary)
#knnFit <- train(TenYearCHD ~ ., data = training, method = "knn", trControl = ctrl, preProcess =
c("center", "scale"), tuneLength = 20)
#Output of kNN fit
knnFit
#Plotting yields Number of Neighbours Vs accuracy (based on repeated cross validation)
plot(knnFit, print.thres = 0.5, type="S")
knnPredict <- predict(knnFit,newdata = testing)
#Get the confusion matrix to see accuracy value and other parameter values
#confusionMatrix(knnPredict, testing$TenYearCHD)
mean(knnPredict == testing$TenYearCHD)
library(pROC)
## Naive bayes
library(e1071)
train.index <- sample(c(1:dim(heart update)[1]), dim(heart update)[1]*0.6)
train.df <- heart update[train.index, ]
valid.df <- heart update[-train.index, ]
train.df$age <- as.factor(train.df$age)</pre>
train.df$education <- as.factor(train.df$education)</pre>
train.df\( cigsPerDay <- as.factor(train.df\( cigsPerDay \)
train.df$totChol <- as.factor(train.df$totChol)</pre>
train.df$sysBP <- as.factor(train.df$sysBP)
train.df$diaBP <- as.factor(train.df$diaBP)</pre>
train.df$heartRate <- as.factor(train.df$heartRate)</pre>
train.df$glucose <- as.factor(train.df$glucose)</pre>
disease.nb <- naiveBayes(TenYearCHD ~ ., data = train.df)
disease.nb
library(rpart)
library(rpart.plot)
# partition
set.seed(1)
train.index <- sample(c(1:dim(heart update)[1]), dim(heart update)[1]*0.6)
train.df <- heart_update[train.index, ]</pre>
valid.df <- heart_update[-train.index, ]</pre>
# classification tree
default.ct <- rpart(TenYearCHD ~ ., data = train.df, method = "class")
# plot tree
prp(default.ct, type = 1, extra = 1, under = TRUE, split.font = 1, varlen = -10)
# set argument type = "class" in predict() to generate predicted class membership.
default.ct.point.pred.train <- predict(default.ct, train.df, type = "class")
# generate confusion matrix for training data
train.df$TenYearCHD <- as.factor(train.df$TenYearCHD)</pre>
confusionMatrix(default.ct.point.pred.train, train.df$TenYearCHD)
# repeat the code for the validation set
valid.df$TenYearCHD <- as.factor(valid.df$TenYearCHD)
default.ct.point.pred.valid <- predict(default.ct, valid.df, type = "class")</pre>
confusionMatrix(default.ct.point.pred.valid, valid.df$TenYearCHD)
```

```
# argument xval refers to the number of folds to use in rpart's built-in
# cross-validation procedure
# argument cp sets the smallest value for the complexity parameter.
cv.ct <- rpart(TenYearCHD ~ ., data = train.df, method = "class", cp = 0.00001, minsplit = 5, xval = 5)
# use printcp() to print the table.
printcp(cv.ct)
## random forest
library(randomForest)
## random forest
rf <- randomForest(TenYearCHD ~ .. data = train.df, ntree = 500, mtry = 4, nodesize = 5, importance =
TRUE)
## variable importance plot
varImpPlot(rf, type = 1)
## confusion matrix
rf.pred <- predict(rf, valid.df)
confusionMatrix(rf.pred, valid.df$TenYearCHD)
require(rpart)
# Split randomly
x <- heart_update[sample(1:nrow(heart_update), nrow(heart_update), replace = F),]
x.train <- heart update[1:floor(nrow(x)*.75), ]
x.evaluate <- heart update[(floor(nrow(x)*.75)+1):nrow(x), ]
# Create a model using "random forest and bagging ensemble algorithms
# utilizing conditional inference trees."
require(party)
x.model <- cforest(as.factor(TenYearCHD) ~ ., data = x.train, control = cforest_unbiased(mtry = 3))
# Alternatively, use "recursive partitioning [...] in a conditional
# inference framework."
# ctree plots nicely (but cforest doesn"t plot)
# plot (x.model)
# Use the model to predict the evaluation.
x.evaluate$prediction <- predict(x.model, newdata=x.evaluate)
# Calculate the overall accuracy.
x.evaluate$correct <- x.evaluate$prediction == x.evaluate$TenYearCHD
print(paste("% of predicted classifications correct", mean(x.evaluate$correct) * 100))
# Extract the class probabilities.
x.evaluate$probabilities
                                                  unlist(treeresponse(x.model,
                                                                                      newdata=x.evaluate),
use.names=F)[seq(1,nrow(x.evaluate)*2,2)]
# Plot the performance of the model applied to the evaluation set as
# an ROC curve.
require(ROCR)
pred <- prediction(x.evaluate$probabilities, x.evaluate$TenYearCHD)</pre>
perf <- performance(pred,"tpr","fpr")</pre>
plot(perf, main="ROC curve", colorize=T)
# And then a lift chart
perf <- performance(pred,"lift","rpp")</pre>
plot(perf, main="lift curve", colorize=T)
# run logistic regression
# partition data
set.seed(2)
train.index <- sample(c(1:dim(heart update)[1]), dim(heart update)[1]*0.6)
```

```
train.df <- heart_update[train.index, ]
valid.df <- heart_update[-train.index.]
# use glm() (general linear model) with family = "binomial" to fit a logistic
# regression.
logit.reg <- glm(as.factor(TenYearCHD) ~ ., data = train.df, family = "binomial")
options(scipen=999)
summary(logit.reg)
predictTrain = predict(logit.reg, type = "response")
summary(predictTrain)
tapply(predictTrain, train.df$TenYearCHD, mean)
table(train.df$TenYearCHD, predictTrain > 0.5)
Sensitivity <- 34/335
\#Sensitivity = 0.1014925
Specificity <- 1840/ 1859
#Specificity = 0.9897795
# use predict() with type = "response" to compute predicted probabilities.
logit.reg.pred <- predict(logit.reg, valid.df, type = "response")</pre>
# first 5 actual and predicted records
data.frame(actual = valid.df$TenYearCHD[1:5], predicted = logit.reg.pred[1:5])
library(gains)
gain <- gains(valid.df$TenYearCHD, logit.reg.pred, groups = length(logit.reg.pred))
# plot lift chart
plot(c(0, gain\$cume.pct.of.total * sum(valid.df\$TenYearCHD)) \sim c(0, gain\$cume.obs), xlab = "# patients",
ylab = "Cumulative", main = "", type = "l")
lines(c(0, sum(valid.df\TenYearCHD)) \sim c(0, dim(valid.df)[1]), lty = 2)
#NeuralNet
library(neuralnet)
library(nnet)
library(caret)
# partition the data
set.seed(2)
train.index <- sample(c(1:dim(heart update)[1]), dim(heart update)[1]*0.6)
train.df <- heart_update[train.index.]
valid.df <- heart_update[-train.index, ]</pre>
valid.index=setdiff(row.names(heart_update), train.index)
nn < -neuralnet(TenYearCHD \sim ., data = train.df, hidden = 2)
training.prediction = compute(nn, train.df)
training.class = apply(training.prediction$net.result, 1, which.max) - 1
training.class = as.factor(training.class)
confusionMatrix(training.class, as.factor(heart_update[train.index,]$TenYearCHD))
validation.prediction = compute(nn, valid.df)
validation.class = apply(validation.prediction$net.result,1,which.max) - 1
validation.class = as.factor(validation.class)
#confusionMatrix(validation.class, as.factor(heart_update[valid.index,]$TenYearCHD))
prob = compute(nn, valid.df[, -ncol(valid.df)] )
prob.result <- prob$net.result
detach(package:neuralnet,unload = T)
library(ROCR)
nn.pred = prediction(prob.result, valid.df$TenYearCHD)
pref <- performance(nn.pred, "tpr", "fpr")</pre>
plot(pref)
library(DiscriMiner)
```

```
da.reg <- linDA(heart_update[,1:15], heart_update[,16])
da.reg$functions
da.reg <- linDA(heart update[, 1:15], heart update[, 16])
# compute probabilities manually (below); or, use Ida() in package MASS with predict()
propensity.risk <- exp(da.reg$scores[,2])/(exp(da.reg$scores[,1])+exp(da.reg$scores[,2]))
data.frame(Actual=heart update$TenYearCHD,
da.reg$classification, da.reg$scores, propensity.risk=propensity.risk)
confusionMatrix(da.reg$classification, as.factor(heart_update$TenYearCHD))
## Linear Discriminant Analysis
set.seed(2)
train.index <- sample(c(1:dim(heart update)[1]), dim(heart update)[1]*0.6)
train.df <- heart update[train.index, ]
valid.df <- heart_update[-train.index, ]</pre>
library(caret)
library(randomForest)
library(AUC)
library(MASS)
model.LDA <- lda(TenYearCHD~., data=train.df, na.action="na.omit")
model.LDA
pc <- predict(model.LDA, na.roughfix(valid.df))</pre>
summary(pc$class)
xtab <- table(pc$class, valid.df$TenYearCHD)
caret::confusionMatrix(xtab, positive = "1")
pb <- NULL
pb <- pc$posterior
pb <- as.data.frame(pb)
colnames(pb) \leftarrow c("X", "Y")
pred.LDA <- data.frame(valid.df$TenYearCHD, pb$Y)</pre>
colnames(pred.LDA) <- c("target", "score")
pred.LDA$target <- as.factor(pred.LDA$target)</pre>
lift.LDA <- lift(target ~ score, data = pred.LDA, cuts=10, class="1")
xyplot(lift.LDA, main="LDA - Lift Chart", type=c("l", "g"), lwd=2
    , scales=list(x=list(alternating=FALSE,tick.number = 10)
             ,y=list(alternating=FALSE,tick.number = 10)))
## Support Vector Machine
library(caret)
intrain <- createDataPartition(y = heart_update$TenYearCHD, p= 0.7, list = FALSE)
training <- heart update[intrain,]
testing <- heart_update[-intrain,]
training[["TenYearCHD"]] = factor(training[["TenYearCHD"]])
trctrl <- trainControl(method = "repeatedcy", number = 10, repeats = 3)
svm_Linear <- train(TenYearCHD ~., data = training, method = "svmLinear", trControl=trctrl, preProcess
= c("center", "scale"), tuneLength = 10)
svm Linear
test_pred <- predict(svm_Linear, newdata = testing)
test pred
confusionMatrix(table(test_pred, testing$TenYearCHD))
grid <- expand.grid(C = c(0,0.01, 0.05, 0.1, 0.25, 0.5, 0.75, 1, 1.25, 1.5, 1.75, 2,5))
```

```
svm_Linear_Grid <- train(TenYearCHD ~., data = training, method = "svmLinear",
trControl=trctrl, preProcess = c("center", "scale"), tuneGrid = grid, tuneLength = 10)
svm_Linear_Grid
plot(svm_Linear_Grid)
test_pred_grid <- predict(svm_Linear_Grid, newdata = testing)
test_pred_grid
confusionMatrix(table(test_pred_grid, testing$TenYearCHD))</pre>
```