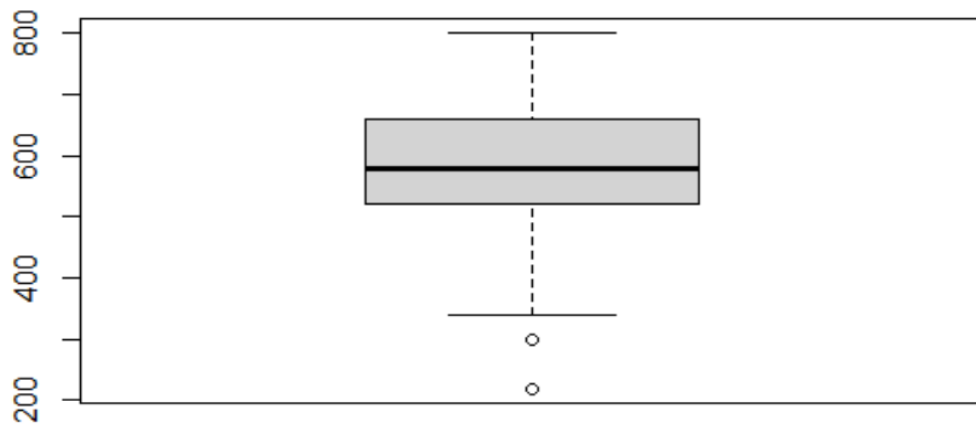
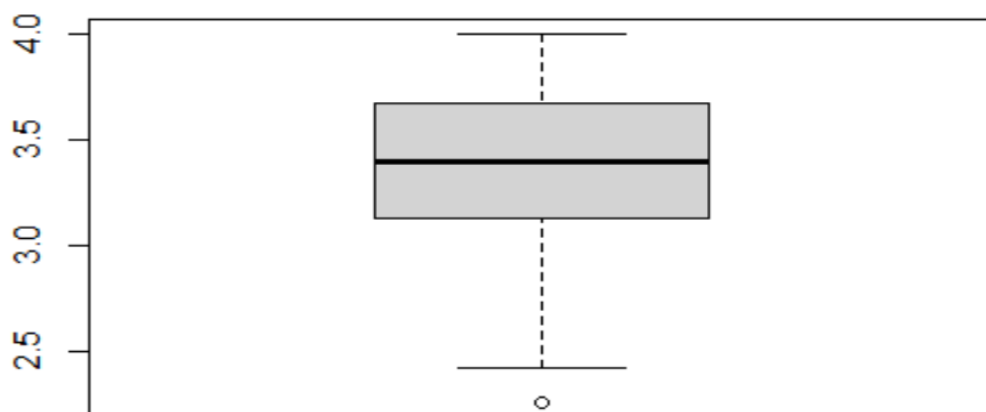


SCREEN SHOTS – COLLEGE ADMISSION PROJECT

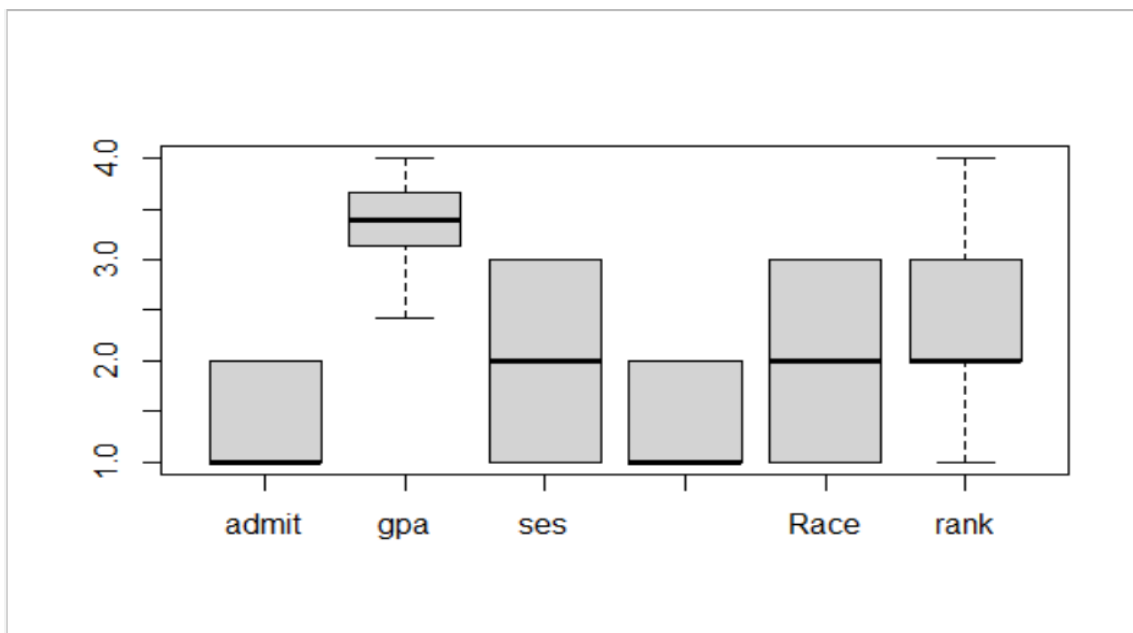
Outlier in GRE:



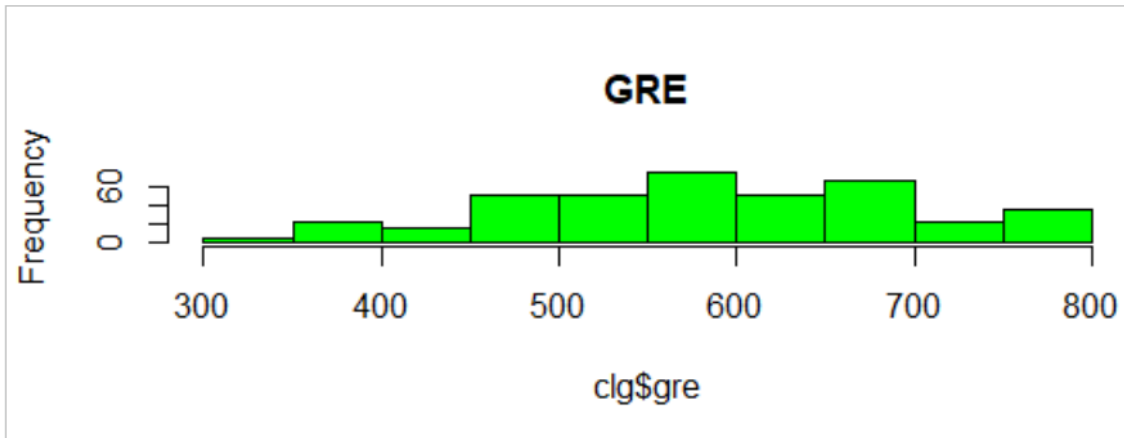
Outlier in GPA:



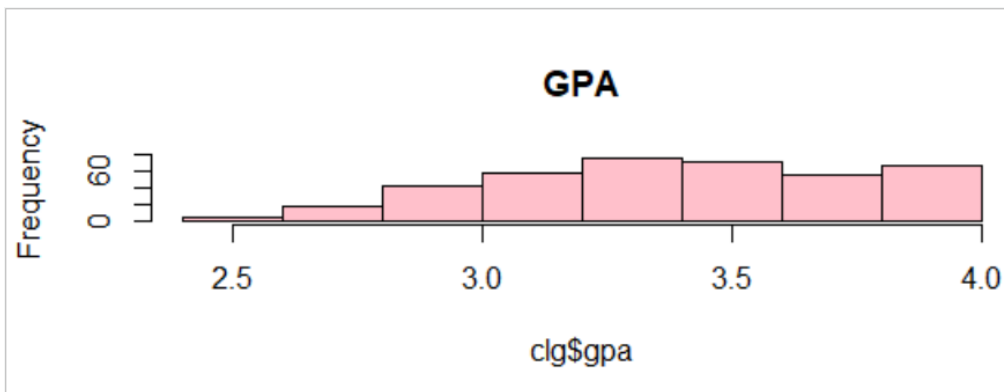
After removing outliers:



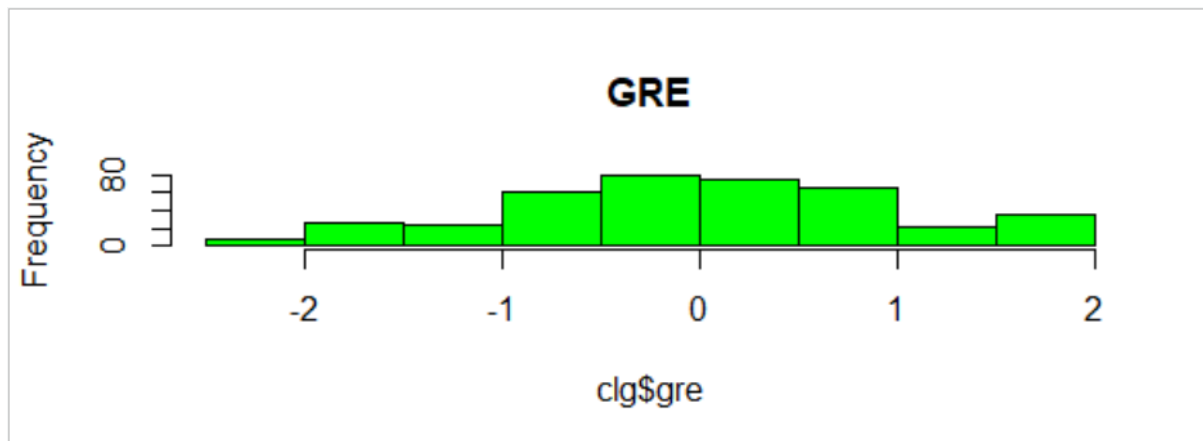
Distribution of GRE:



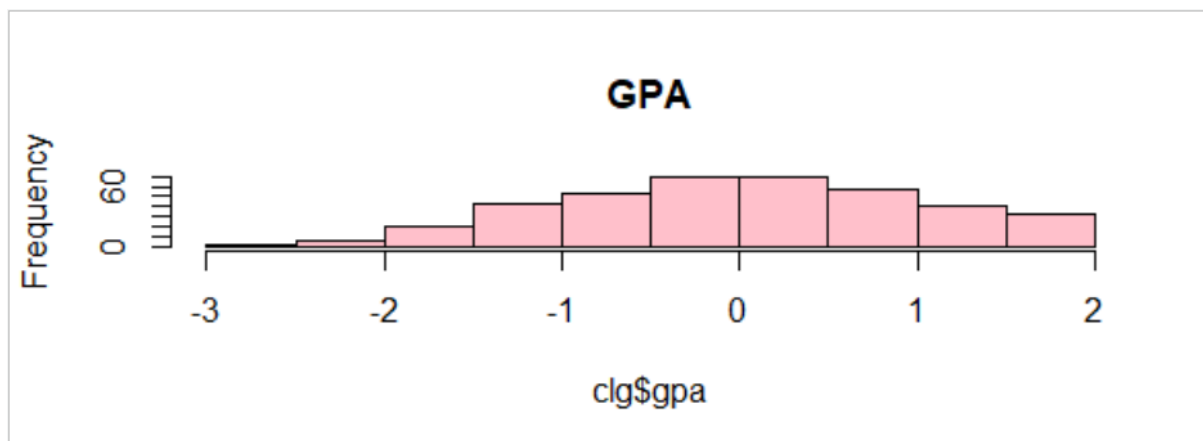
Distribution of GPA:



After scaling GRE:



After scaling GPA:



Logistic regression: Accuracy & confusion matrix

```
Console Terminal x Jobs x
R 4.1.2 · F:/R_programming/
(Intercept) -0.7885 0.1264 -6.238 4.44e-10 ***
gpa 0.4042 0.1362 2.968 0.00299 **
gre 0.3046 0.1356 2.247 0.02463 *
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

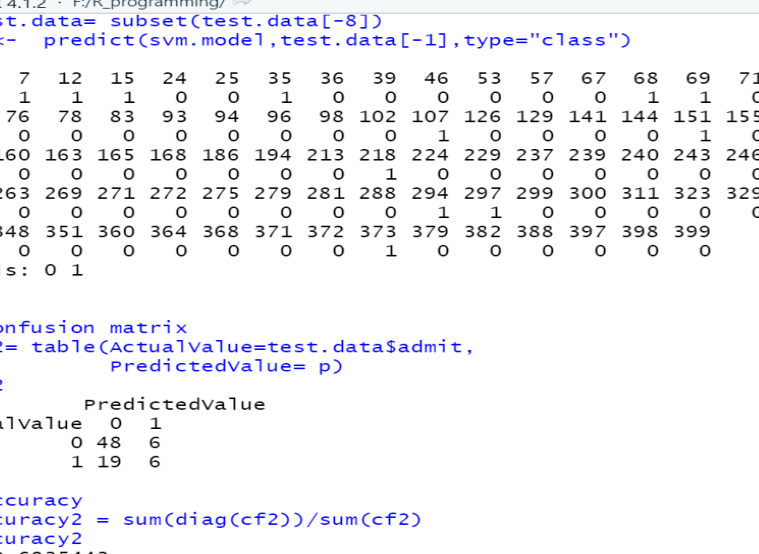
(Dispersion parameter for binomial family taken to be 1)

Null deviance: 396.00 on 315 degrees of freedom
Residual deviance: 375.07 on 313 degrees of freedom
AIC: 381.07

Number of Fisher scoring iterations: 4

> admitPredict = predict(model2, test.data, type = "response")
> test.data$admit1 = ifelse(admitPredict > 0.5, 1, 0)
>
> #confusion matrix
> cf1 = table(ActualValue = test.data$admit,
+            PredictedValue = test.data$admit1)
> #accuracy
> accuracy1 = sum(diag(cf1)) / sum(cf1)
> accuracy1
[1] 0.6708861
> cf1
      PredictedValue
ActualValue 0 1
0 47 7
1 19 6
> |
```

SVM model : Accuracy and confusion matrix



```
R 4.1.2 · F:/R_programming/
> test.data= subset(test.data[-8])
> p <- predict(svm.model,test.data[-1],type="class")
> p
  6    7   12   15   24   25   35   36   39   46   53   57   67   68   69   71
0    1    1    1    0    0    1    0    0    0    0    0    0    1    1    0
73   76   78   83   93   94   96   98  102  107  126  129  141  144  151  155
0    0    0    0    0    0    0    0    0    1    0    0    0    0    1    0
156 160 163 165 168 186 194 213 218 224 229 237 239 240 243 246
0    0    0    0    0    0    0    0    1    0    0    0    0    0    0    0
260 263 269 271 272 275 279 281 288 294 297 299 300 311 323 329
0    0    0    0    0    0    0    0    1    1    0    0    0    0    0    0
335 348 351 360 364 368 371 372 373 379 382 388 397 398 399
0    0    0    0    0    0    0    0    1    0    0    0    0    0    0
Levels: 0 1
>
>
> #confusion matrix
> cf2= table(ActualValue=test.data$admit,
+           PredictedValue= p)
> cf2
      PredictedValue
ActualValue 0      1
0      48      6
1      19      6
>
> #accuracy
> accuracy2 = sum(diag(cf2))/sum(cf2)
> accuracy2
[1] 0.6835443
>
```

KNN model: Accuracy and confusion matrix

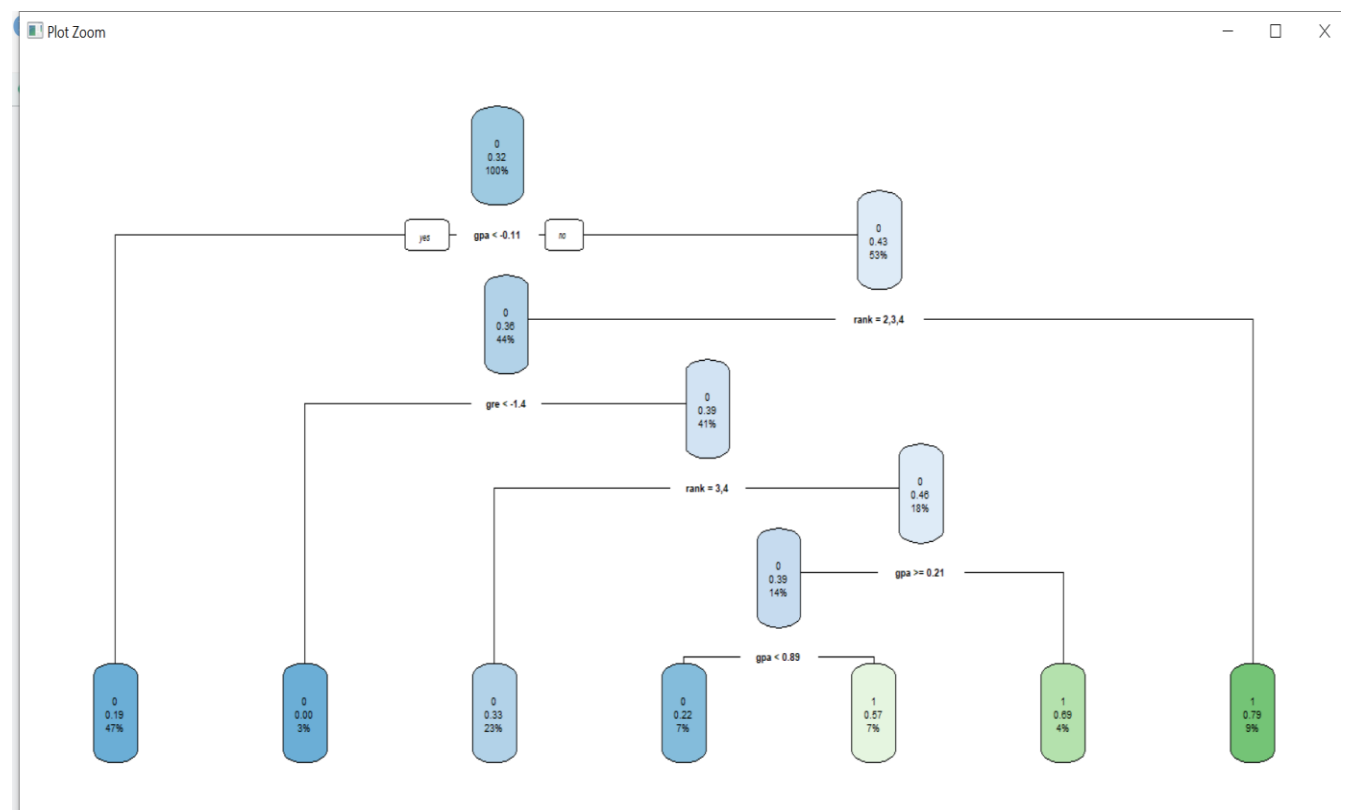
```
R 4.1.2 · F:/R_programming/ ↗  
> accuracy2 = sum(diag(cf2))/sum(cf2)  
> accuracy2  
[1] 0.6835443  
> library(caTools)  
> library(class)  
>  
> knn.data=knn(train.data, test.data, train.data$admit, k=19)  
> knn.data  
[1] 1 0 0 1 0 1 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 1 0 0 0 0 0 1  
[32] 0 0 0 1 0 0 0 0 0 0 1 0 0 1 0 0 1 0 0 0 1 0 0 0 0 0 0 0 1 0 0 0 0 0 0 1 0 0 0 0 0 0  
[63] 0 0 1 0 0 0 0 0 0 0 1 1 0 0 1 0 0 0 0 0  
Levels: 0 1  
>  
> # Confusion Matrix  
> cf3 =table(test.data$admit, knn.data)  
> cf3  
      knn.data  
    0   1  
0  54   0  
1   9  16  
>  
> #Accuracy  
> accuracy3 =sum(diag(cf3))/sum(cf3)  
> accuracy3  
[1] 0.8860759  
>  
> misClassError = mean(knn.data != test.data$admit)  
> print(paste('Accuracy =', 1-misClassError))  
[1] "Accuracy = 0.886075949367089"  
>
```

Naïve bayes: Accuracy and confusion matrix

```
Console Terminal Jobs
R 4.1.2 · F:/R_programming/
Gender_Male Race rank
6      1      1      2
7      1      2      1
12     0      2      1
15     1      1      1
24     0      1      4
25     0      2      2
>
> View(p)
>
> #prediction
> p1 = predict(NB_Model,test.data)
Warning message:
predict.naive_bayes(): more features in the newdata are provided as t
here are probability tables in the object. Calculation is performed b
ased on features to be found in the tables.
>
> #confusion matrix
> cf4 = table(p1, test.data$admit)
> cf4

p1    0  1
0  49  20
1   5   5
>
> #accuracy
> accuracy4 =sum(diag(cf4))/sum(cf4)
> accuracy4
[1] 0.6835443
>
```

Decision Tree:



Categorisation of data:

