library(e1071) library(caret) library(caTools) library(rpart) library(rpart.plot) library(kernlab) library(readr) library(ggplot2)

#### Working directory

getwd()

### Extracting data from csv file

df\_admission <- read.csv('College\_admission.csv')

class(df\_admission) dim(df\_admission) View(df\_admission) names(df\_admission)

#### Finding the missing values

sum(is.na(df\_admission))

#### Finding outliers using boxplot

boxplot(df\_admission\$gre, main="College admission ", xlab ="Gre Score", ylab ="Gre", horizontal=TRUE) summary(df\_admission)

#### Creating copy of the dataframe for further analysis

admin <- data.frame(df\_admission)

#### Finding the quartile to remove outliers

Q <- quantile(admin\$gre,probs=c(.25, .75), na.rm = FALSE) iqr<- IQR(admin\$gre) up <- Q[2]+1.5 iqr # Upper Range low<- Q[1]-1.5 iqr # Lower Range admin <- subset(admin, admin\$gre > low & admin\$gre < up) View(admin) dim(admin)

#### Boxplot showing no outliers

boxplot(admin\$gre, main="College admission", xlab = "Gre Score", ylab = "Gre", horizontal=TRUE)

## Find the structure of the data set and if required, transform the numeric data type to factor and vice-versa.

str(admin)

### Finding whether data is normally distributed or not

mean(admin\$gre) sd(admin\$gre)

d\_norm <- dnorm(admin\$gre, mean = 590.8081, sd = 111.7971) d\_norm plot(density(d\_norm, adjust = 10))# The density plot talks that dataframe is normally distributed

#### Use variable reduction techniques to identify significant variables

### Finding correlation between dependent variable and independent variable

View(cor(admin))

### Dropping insignificant column

# Run logistic model to determine the factors that influence the admission process of a student

#### (Drop insignificant variables)

#### Separating the categorical variable and converting into factor

rank\_cat<-admin[,-c(1,2,3)] rank\_cat <- as.factor(rank\_cat) View(rank\_cat)

#### Creating dummy variable

dummy\_rank <- data.frame(model.matrix(~rank\_cat-1)) View(dummy\_rank)

#### Combing the dummy columns and the other columns

admin\_final <- cbind(admin[,c(1,2,3)],dummy\_rank) View(admin\_final)

#### Spliting the data into train and test

```
set.seed(123)
indices = sample.split(admin_final$admit, SplitRatio = 0.7)
train = admin_final[indices,]
test = admin_final[!(indices),]
dim(train) dim(test)

Model Building

Logistic model
model_1 = glm(admit ~ ., data = train, family = "binomial")
summary(model_1)
View(test)
```

#### **Test Data**

#### Probabilities prediction of admit variable for test data

```
test_pred = predict(model_1, type = "response", newdata = test)
test_pred
test$prob <- test_pred
View(test)</pre>
```

#### Using the probability cutoff of 50%.

```
test_pred_admit <- factor(ifelse(test_pred >= 0.50, "Yes", "No")) test_actual_admit <- factor(ifelse(test$admit==1,"Yes","No")) test_actual_admit test_pred_admit table(test_actual_admit,test_pred_admit)
```

 $test\_conf <- confusion Matrix (test\_pred\_admit, test\_actual\_admit, positive = "Yes") \ test\_confusion Matrix (test\_pred\_admit, test\_actual\_admit, positive = "Yes") \ test\_actual\_admit, positive = "Yes"$ 

### Based on specificity we can say that it is good in predicting class 0

```
admin_tree <- admin

prop.table(table(admin_tree$admit))

table(admin_tree$admit)

set.seed(123) split.indices <- sample(nrow(admin_tree), nrow(admin_tree)*0.7, replace = F) train <- admin_tree[split.indices,] test <- admin_tree[-split.indices,]

View(train)
```

#### **Decision Tree**

tree.model <- rpart(admit ~ ., # formula data = train, # training data method = "class") # classification not regression

#### display decision tree

prp(tree.model)

#### make predictions on the test set

tree.predict <- predict(tree.model, test, type = "class") tree.predict

#### evaluate the results

confusionMatrix(tree.predict, as.factor(test\$admit), positive = "1")

#### Based on specificity above model is good in prediction of class 0

**SVM Model** 

#### Converting dependent variable into factor

```
admin_svm <- admin_final
admin_svm$admit <- as.factor(admin_svm$admit)
set.seed(123)
indices = sample.split(admin_svm$admit, SplitRatio = 0.7)
train = admin_svm[indices,]
test = admin_svm[!(indices),]
dim(train)
dim(test)
model_svm<- ksvm(admit ~ ., data = train,scale = FALSE, C=1)
```

#### Predicting the model results

```
evaluate_1<- predict(model_svm, test)
levels(as.factor(evaluate_1))
```

#### Confusion Matrix - Finding accuracy, Sensitivity and specificity

confusionMatrix(evaluate\_1, as.factor(test\$admit))

#### Based on sensitivity above model is good in prediction of class 1

Based on the three model Logistic Regression, Decision tree and SVM model

## We can conclude that Logistic regression and SVM model are better fit model

### Accuracy, Sensitivity and Specificity is very good as compared to Decision tree

#### **Descriptive Analysis**

admin1 <- admin admin1 <- within(admin1, { gre\_cat <- NA # need to initialize variable gre\_cat[gre <= 440] <- "Low" gre\_cat[gre > 440 & gre <= 580] <- "Middle" gre\_cat[gre > 580] <- "High" } ) str(admin1) dim(admin1) admin1\$gre\_cat <- factor(admin1\$gre\_cat, levels = c("High", "Middle", "Low"))

str(admin1\$gre\_cat)

 $sum\_Desc <- aggregate(admit \sim gre\_cat, admin1, FUN = sum) \ length\_Desc <- aggregate(admit \sim gre\_cat, admin1, FUN = length) \ probability\_table <- cbind(sum\_Desc, recs = length\_Desc[,2])$ 

 $probability\_table\_final <- transform (probability\_table\_probability\_admission = admit/recs) \ probability\_table\_final <- transform (probability\_table\_prob$ 

#### Creating the point chart

ggplot(probability\_table\_final, aes(x = gre\_cat, y = probability\_admission))+geom\_point() table(admin1\$admit, admin1\$gre\_cat) knitr::stitch('Assignment.R') Sys.which("pdflatex") pdflatex ""