

#Admission Dataset

#load packages

library("MLmetrics") #metric for model

library(readxl)#to read excel

library(plyr)#split and put data

library(DescTools) #For Mode

library(dplyr) #For Pipe Operator

library(ggplot2) #For QQ Plot

library(moments) #For Skewness & Kurtosis

library(rstatix) #For Welch ANOVA Test

library(Hmisc) #For rcorr Function & Missing Values Treatment

library(QuantPsyc) #For lm.beta function

library(ggpubr) #For advanced QQ Plots

library(aod) #For wald.test function in logistic regression

library(fmsb)#For NagelkerkeR2 function in logistic regression

library(caret) #For Data manipulation

library(imputeMissings) #For imputing missing Values

library(purrr) #For Missing Values

library(naivebayes) #For naive bayes

library(rpart.plot) #For DT Graph

library(psych) #descriptive stats

library(rpart) #for DT

library(e1071) #for SVM

library(knitr) #to save as pdf

library(gridExtra)#work with grid graphics

library(datasets)#dataset related operations

library(tinytex)# to convert to latex

```
#load and attach Admissions dataset
```

```
AdmissionData<- read_xlsx("C:\\Users\\sujoydutta\\Desktop\\Data analysis\\R\\Datasets_R\\Lesson  
8_Classification\\Admission data.xlsx")
```

```
str(AdmissionData)
```

```
#converting to dataframe
```

```
Add<-data.frame(AdmissionData)
```

```
str(Add)
```

```
#####
```

```
#converting variables to appropriate types
```

```
Add$admit<-as.factor(AdmissionData$admit)
```

```
Add$rank<-as.factor(AdmissionData$rank)
```

```
Add$gre<-as.numeric(AdmissionData$gre)
```

```
Add$gpa<-as.numeric(AdmissionData$gpa)
```

```
str(Add)
```

```
#####
```

```
# check for missing values
```

```
map(Add, ~sum(is.na(.)))
```

```
#Checking Skewness
```

```
sapply(Add[(2:3)], "skewness")
```

```
#boxplot to see outliers
```

```
boxplot(Add$gre)
```

```
boxplot(Add$gpa)
```

```
#getting the outliers
```

```
outA<-boxplot(Add$gre)$out
```

```
outB<-boxplot(Add$gpa)$out
```

```
#removing the outliers
```

```
Addnew<- Add
```

```
Addnew<-Addnew[-which(Add$gre %in% outA),]
```

```
Addnew<-Addnew[-which(Add$gpa %in% outB),]
```

```
str(Addnew)
```

```
#####
```

```
#categorizing gre
```

```
Addnew<- transform(Addnew,GRE_lvl = ifelse(gre<440,"low gre",ifelse(gre<580,"medium gre","high gre")))
```

```
str(Addnew)
```

```
#summary stats
```

```
summary(Addnew)
```

```
describe(Addnew)
```

```
#normality of new dataset
```

```
skewness(Addnew$gre)
```

```
kurtosis(Addnew$gre)
```

```
skewness(Addnew$gpa)
```

```
kurtosis(Addnew$gpa)
```

```
#correlation of gpa and gre
```

```
ggplot(Addnew, aes(x =gpa, y = gre)) +
```

```

geom_line() +
labs(title = "GPA and GRA relation",
      x = "GPA",
      y = "GRA")
corr.test(Addnew$gpa,Addnew$gre,method = "pearson", use = "complete.obs")

```

```

# GRE_lvl distribution
ggplot(Addnew, aes(x = GRE_lvl)) +
  geom_density(fill = "indianred3") +
  labs(title = "Participants by GRE_lvl")

```

```

# seeing the race distribution

```

```

plotdata <- Addnew %>%
  count(Race) %>%
  arrange(desc(Race)) %>%
  mutate(prop = round(n*100/sum(n), 1),
         lab.ypos = cumsum(prop) - 0.5*prop)

plotdata$label <- paste0(plotdata$Race, "\n",
                        round(plotdata$prop), "%")

```

```

ggplot(plotdata,
  aes(x = "",
      y = prop,
      fill =Race)) +
  geom_bar(width = 1,
           stat = "identity",
           color = "red") +
  geom_text(aes(y = lab.ypos, label = label),
           color = "yellow") +

```

```

coord_polar("y",
            start = 0,
            direction = -1) +
theme_void() +
theme(legend.position = "FALSE") +
labs(title = "Race Distribution")
# seeing GRE levels distribution by Rank
ggplot(Addnew,
       aes(x = rank,
           fill = GRE_lvl)) +
geom_bar(position = "dodge")

#To plot GRE per Social economic level
plot(Addnew$ses,Addnew$gre, xlab = "Number of prospects", ylab = "duration per prospect", col =
"blue")

# Calculate test statistics
add_aov <- aov(Addnew$gre~factor(Addnew$ses))
summary(add_aov)

#seeing which gender gets more admission
ggplot(Addnew, aes(x = Gender_Male, fill = admit)) +
geom_bar(position = "fill") +
theme_classic()+
labs(x = "Gender",y = "Count",fill = "Admission")

# Seeing if high gpa leads to person taking the admission?
boxplot(Addnew$gpa~factor(Addnew$admit),
       xlab = "Did they take the admission?", ylab = "GPA")

```

```

# Calculate test statistics

add_aov <- aov(Addnew$gpa~factor(Addnew$admit))

summary(add_aov)


str(Addnew)

#####

#Data partition

set.seed(123)

Train <- createDataPartition(Addnew$admit, p=0.7, list=FALSE)

training <- Addnew[ Train, ]

testing <- Addnew[ -Train, ]


#####

#Model1- Logistic Regression

#create the model using train()

Model1 <- train(admit~., data=training, method="glm", family="binomial")

summary(Model1)


#exponentiated coefficients

exp(coef(Model1$finalModel))


# prediction for test data

predadmit<-predict(Model1, newdata=testing)


#fitness metrics for validation

confusionMatrix(predadmit, testing$admit, mode = "everything", positive="1")


#predicting probability for new values

newvalues<-data.frame(gre=800,gpa=4,ses=2,GRE_lvl='medium
gre',ses=as.factor(2),Gender_Male=1,Race=1,rank=as.factor(1))

```

```
pred<-predict(Model1, newdata=newvalues,type="prob")
```

```
pred
```

```
#####
```

```
#Model2- SVM
```

```
#create the model using train()
```

```
Model2<-svm(admit~.,data=training)
```

```
summary(Model2)
```

```
# prediction for test data
```

```
predadmit<-predict(Model2, newdata=testing)
```

```
#fitness metrics for validation
```

```
confusionMatrix(predadmit, testing$admit, mode = "everything", positive="1")
```

```
#predicting probability for new values
```

```
newvalues<-data.frame(gre=900,gpa=3.8,ses=2,GRE_lvl='high  
gre',Gender_Male=1,Race=2,rank=as.factor(3))
```

```
pred<-predict(Model2, newdata=newvalues,type="prob")
```

```
pred
```

```
#Model3- Decision Tree
```

```
#create the model using train()
```

```
Model3 <- train(admit~., data=training, method="rpart",parms= list(split="information"))
```

```
summary(Model3)
```

```
# prediction for test data
```

```
predadmit<-predict(Model3, newdata=testing)
```

```
#fitness metrics for validation
```

```
confusionMatrix(predadmit, testing$admit, mode = "everything", positive="1")
```

```
#predicting probability for new values
```

```
newvalues<-data.frame(gre=650,gpa=4,ses=1,GRE_lvl='medium  
gre',Gender_Male=1,Race=3,rank=as.factor(4))
```

```
pred<-predict(Model2, newdata=newvalues,type="prob")
```

```
pred
```

```
#converting to pdf
```

```
knitr::stitch('Admissiondata.r')
```

```
setwd("C:\\Users\\sujoydutta\\Desktop\\Data analysis\\R\\R scripts")
```

```
dir()
```

```
"Admissiondata.R"
```

```
knitr::stitch('Admissiondata.R')
```