ASSIGNMENT-7

```
Question-1)
#Question 1
train = read.csv("C:\fy\text{YUsers}\fy\text{KIRAN KONDISETTI}\fy\text{Desktop}\fy\text{Train.csv}")
test = read.csv('C:\forall YUsers\forall YKIRAN KONDISETTI\forall YDesktop\forall YTest.csv')
test$Age[is.na(test$Age)] = mean(test$Age, na.rm=TRUE)
train$Age[is.na(train$Age)] = mean(train$Age, na.rm=TRUE)
levels(train$Embarked)
levels(train$Embarked)[1] = 'S'
train$Embarked <- sapply(as.character(train$Embarked), switch, 'C' = 0, 'Q' = 1, 'S' = 2)
test$Embarked <- sapply(as.character(test$Embarked), switch, 'C' = 0, 'Q' = 1, 'S' = 2)
train$Sex <- ifelse(train$Sex == 'male', 1, 0)
test$Sex <- ifelse(test$Sex == 'male', 1, 0)
#fix(train)
#logistic regression
attach(train)
LR
              glm(
                        factor(Survived)~factor(Pclass)
                                                                    Sex+
                                                                              Age+SibSp+
Parch+factor(Embarked), data= train, family= 'binomial')
LR.probs = predict(LR, type= 'response', newdata= test)
LR.probs
LR.probs = ifelse(LR.probs > 0.5, 1, 0)
```

```
table(LR.probs,test$Survived)
mean(LR.probs == test$Survived)
#LDA
library(MASS)
              lda(factor(Survived)~
                                      factor(Pclass)
                                                                     Age+SibSp+
LDA
                                                            Sex+
                                                       +
Parch+factor(Embarked) , data= train)
LDA.pred = predict(LDA, newdata= test)
LDA.class = LDA.pred\$class
LDA.class
table(LDA.class, test$Survived )
mean(LDA.class != test$Survived)
#QDA
library(MASS)
          qda(factor(Survived)~factor(Pclass)
                                                  Sex+ Age+SibSp+
                                                                        Parch
factor(Embarked), data= train, family= binomial)
QDA.pred = predict(QDA, newdata= test)
QDA.class = QDA.pred$class
QDA.class
table(QDA.class, test$Survived )
mean(QDA.class != test$Survived)
```

```
.#KNN
library(FNN)
#train$Age <- ifelse(train$Age<18, 1, 0)</pre>
#test$Age <- ifelse(test$Age<18, 1, 0)</pre>
train$Age <- ifelse(train$Age<18, 1, 0)
test$Age <- ifelse(test$Age<18, 1, 0)
train1 = train[, -c(1,2,4,9,10,11)]
test1 = test[, -c(1,2,4,9,10,11)]
train.Y = train$Survived
train1
set.seed(12)
knn1 <- knn(train1, test1, train.Y, k=1)
table(knn1, test$Survived)
mean(knn1 != test$Survived)
knn2 = knn(train1, test1, train.Y, k=2)
table(knn2, test$Survived)
mean(knn2 != test$Survived)
knn3 = knn(train1, test1, train.Y, k=3)
table(knn3, test$Survived)
mean(knn3!= test$Survived)
knn4 = knn(train1, test1, train.Y, k=4)
```

```
table(knn4, test$Survived)
mean(knn4!= test$Survived)
knn5 = knn(train1, test1, train.Y, k=5)
table(knn5, test$Survived)
mean(knn5 != test$Survived)
> table(LR.probs,test$Survived)
LR.probs
           0
        0 144 29
1 25 69
> mean(LR.probs == test$Survived)
[1] 0.7977528_
> table(LDA.class, test$Survived )
LDA.class 0
         0 146 30
         1 23 68
> mean(LDA.class != test$Survived)
 [1] 0.1985019
> table(QDA.class, test$Survived )
QDA.class 0
         0 130
                20
         1 39 78
> mean(QDA.class != test$Survived)
[1] 0.2209738
> table(knn1, test$Survived)
knn1 0
   0 137
          33
    1 32 65
> mean(knn1 != test$Survived)
[1] 0.2434457
> table(knn2, test$Survived)
knn2
      0
            1
   0 146 39
1 23 59
> mean(knn2 != test$Survived)
[1] 0.2322097
> table(knn3, test$Survived)
knn3 0
   13 0 1
0 149 34
1 20 64
> mean(knn3 != test$Survived)
[1] 0.2022472
> table(knn4, test$Survived)
knn4
    0 146
          37
    1 23
          61
```

Ans- The suitable value of K for KNN model is 3, which as a misclassification rate of 0.18. The TP of Logistic Regression, LDA, QDA and KNN(K=3) is 144, 146, 130 and 149 respectively. The FP for the models is 25, 23, 39, 20 respectively. True positive value tells us the number of positive classifications classified as positive. True negative value tells us the number of negative classifications classified as positive.

```
Question2)
#question 2
train = read.csv("C:\footnote{Y}\text{Users}\footnote{Y}\text{KIRAN KONDISETTI}\footnote{Y}\text{Desktop}\footnote{Y}\text{Train.csv"})
test = read.csv('C:\footnote{Y}\text{Users}\footnote{Y}\text{KIRAN KONDISETTI}\footnote{Y}\text{Desktop}\footnote{Y}\text{Test.csv'})
test\footnote{Age}[is.na(test\footnote{Age})] = mean(test\footnote{Age}, na.rm=TRUE)
train\footnote{Age}[is.na(train\footnote{Age})] = mean(train\footnote{Age}, na.rm=TRUE)
levels(train\footnote{Embarked})
levels(train\footnote{Embarked})[1] = 'S'
install.packages('naniar')
library(naniar)
levels(test\footnote{Cabin})
levels(test\footnote{Cabin})[1] = 'NA'
attach(test)
```

```
formula1
                as.formula('factor(Survived)~factor(Pclass)
                                                                  Sex+ Age+SibSp+
Parch+factor(Embarked)+Cabin')
LR C = glm(formula1,data= test, family='binomial')
LR C.probs = predict(LR C)
LR_C.probs = ifelse(LR_C.probs > 0.5, 1, 0)
mean(LR C.probs!= test$Survived)
LR C1
                 glm(factor(Survived)~factor(Pclass)
                                                                Sex+
                                                                          Age+SibSp+
Parch+factor(Embarked),data= test, family= 'binomial')
LR\_C1.probs = predict(LR\_C1)
LR_C1.probs = ifelse(LR_C1.probs > 0.5, 1, 0)
mean(LR C1.probs!= test$Survived)
> mean(LR_C.probs!= test$Survived)
[1] 0.1310861
> mean(LR_C1.probs!= test$Survived)
[1] 0.1685393
```

Ans) Training and testing are done on the test data set, since levels present in train and test data set are different. Error named – 'different levels present' is displayed when the model is trained on train data and tested on test data. This is the reason test data set is used to train and test the data. The misclassification of the model is 0.131, when cabin was included and 0.1685, when cabin is not included. This tells us that cabin feature has significance when it is included with the other predictors in the model.

```
Question-3) #question3
```

Ans- The Adjusted odds ratio is calculated using the model, hence it's a multi-variate function. Unadjusted odd ratio is calculated for each variable hence it is a Uni-variate function. The adjusted odd ratio for P-class2, P-class3, sex, embarked C and embarked Q is 0.45, 0.12, 0.07174, 1.48 and 1.51. This tells us that when we increase P-class2 by 1 unit there will be a decrease in the response by a factor 0.45, when we increase sex by 1 unit there will be a decrease in the response by a factor 0.07174. And, when we increase Embarked C and Embarked Q by 1 unit there will be an increase in the response by a factor 1.48 and 1.51 respectively.

```
Question-4)

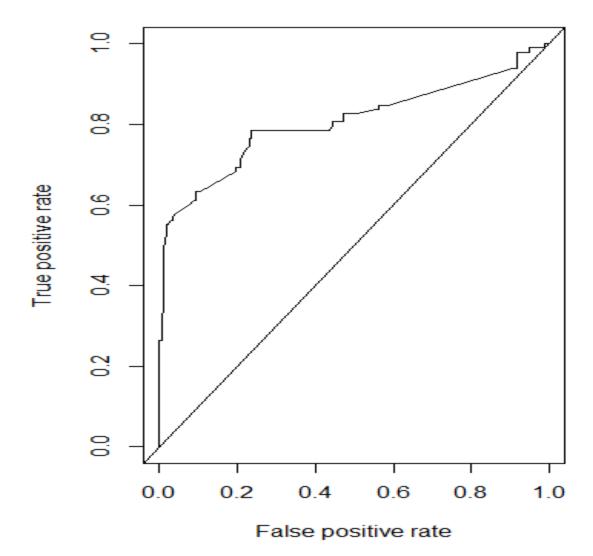
#question 4

formula = as.formula('factor(Survived)~factor(Pclass) + Sex+ Age+SibSp+
Parch+factor(Embarked)')

LR_0.5 = glm(formula, data= train, family= 'binomial')
```

```
LR.probs 0.5 = predict(LR 0.5, type= 'response', newdata= test)
LR.probs 0.5 = ifelse(LR.probs 0.5 > 0.5, 1, 0)
table(LR.probs 0.5,test$Survived)
mean(LR.probs 0.5!= test$Survived)
LR_0.2 = glm(formula, data= train, family= 'binomial')
LR.probs 0.2 = predict(LR 0.2, type= 'response', newdata= test)
LR.probs 0.2 = ifelse(LR.probs 0.2 > 0.2, 1, 0)
table(LR.probs 0.2,test$Survived)
mean(LR.probs 0.2!= test$Survived)
LR_0.8= glm(formula, data= train, family= 'binomial')
LR.probs 0.8 = predict(LR 0.8, type= 'response', newdata= test)
LR.probs 0.8 = ifelse(LR.probs 0.8 > 0.8, 1, 0)
table(LR.probs_0.8,test$Survived)
mean(LR.probs 0.8!= test$Survived)
> table(LR.probs_0.5,test$Survived)
LR.probs_0.5
                   1
29
            1 25 69
 mean(LR.probs_0.5!= test$Survived)
> table(LR.probs_0.2,test$Survived)
LR.probs_0.2 0 1
            0 97 16
            1 72 82
> mean(LR.probs_0.2!= test$Survived)
[1] 0.329588
```

```
> table(LR.probs_0.8,test$Survived)
Ans- Threshold value of 0.5 is appropriate for survival prediction. The misclassification
rate for threshold 0.5 is 0.20.
Question-5)
#question 5
install.packages('ROCR')
library(ROCR)
pred= predict(LR, type= 'response', newdata= test)
pr = prediction(pred,test$Survived)
prf<- performance(pr, measure='tpr', x.measure='fpr')</pre>
plot(prf)
abline(0,1)
prf
auc_ROCR <- performance(pr, measure = "auc")</pre>
auc<- auc_ROCR@y.values[[1]]</pre>
auc
> auc
[1] 0.8060621
```



```
Question-6)
#question 6
#question 6
library(FNN)
train = read.csv("C:\fy\text{YUsers\fy\text{YKIRAN KONDISETTI\fy\text{Pesktop\fy\fy\text{Train.csv"}}}
test = read.csv('C:\forall YUsers\forall KIRAN KONDISETTI\forall YDesktop\forall Test.csv')
test$Age[is.na(test$Age)] = mean(test$Age, na.rm=TRUE)
train$Age[is.na(train$Age)] = mean(train$Age, na.rm=TRUE)
levels(train$Embarked)
levels(train$Embarked)[1] = 'S'
train$Embarked <- sapply(as.character(train$Embarked), switch, 'C' = 0, 'Q' = 1, 'S' = 2)
test$Embarked <- sapply(as.character(test$Embarked), switch, 'C' = 0, 'Q' = 1, 'S' = 2)
train$Sex <- ifelse(train$Sex == 'male', 1, 0)
test$Sex <- ifelse(test$Sex == 'male', 1, 0)
LR 7
                 glm(factor(Survived)~factor(Pclass)
                                                                             Age+SibSp+
                                                                   Sex+
Parch+factor(Embarked)+Parch+Fare,data = train, family= 'binomial')
summary(LR_7)
train2 = train[, -c(1,2,4,8,9,10,11,12)]
test2 = test[, -c(1,2,4,8,9,10,11,12)]
train.Y1= train$Survived
train1
set.seed(12)
```

```
knn11 <- knn(train2, test2, train.Y1, k=1)
table(knn11, test$Survived)
mean(knn11!= test$Survived)
knn21 = knn(train2, test2, train.Y1, k=2)
table(knn21, test$Survived)
mean(knn21 != test$Survived)
knn31 = knn(train2, test2, train.Y1, k=3)
table(knn31, test$Survived)
mean(knn31 != test$Survived)
knn41 = knn(train2, test2, train.Y1, k=4)
table(knn41, test$Survived)
mean(knn41!= test$Survived)
knn51 = knn(train2, test2, train.Y1, k=5)
table(knn51, test$Survived)
mean(knn51 != test$Survived)
Output-
table(knn11, test$Survived)
knn11
         0
     0 135
            35
       34 63
  mean(knn11 != test$Survived)
[1] 0.258427
> table(knn21, test$Survived)
knn21
     0 143 27
1 26 71
> mean(knn21 != test$Survived)
[1] 0.1985019
> table(knn31, test$Survived)
```

```
31 0 1
0 139 29
knn31
    1 30 69
> mean(knn31 != test$survived)
[1] 0.2209738
> table(knn41, test$Survived)
      0
knn41
    0 148 33
    1 21 65
> mean(knn41 != test$Survived)
[1] 0.2022472
> table(knn51, test$Survived)
knn51 0 1
0 145 36
    1 24 62
 mean(knn51 != test$Survived)
[1] 0.2247191
>
```

Ans) The significant features can be found out using p-values of the predictors from Logistic regression model and they are P-class, sex, age, SibSp. Using these features KNN model is built. The model performs better than the previous model since the misclassification error for K=2 is 0.198.

R-CODE-

```
#Question 1
train = read.csv("C:\footnote{Y}\text{Users}\footnote{Y}\text{KIRAN KONDISETTI}\footnote{Y}\text{Desktop}\footnote{Y}\text{Train.csv"})
test = read.csv('C:\footnote{Y}\text{Users}\footnote{Y}\text{KIRAN KONDISETTI}\footnote{Y}\text{Desktop}\footnote{Y}\text{Test.csv'})
test\footnote{Age[is.na(test\footnote{Age})] = mean(test\footnote{Age, na.rm}=TRUE)}
train\footnote{Age[is.na(train\footnote{Age})] = mean(train\footnote{Age, na.rm}=TRUE)}
levels(train\footnote{Embarked})
levels(train\footnote{Embarked})[1] = 'S'
train\footnote{Embarked} <- sapply(as.character(train\footnote{Embarked}), switch, 'C' = 0, 'Q' = 1, 'S' = 2)</pre>
```

```
test$Embarked <- sapply(as.character(test$Embarked), switch, 'C' = 0, 'Q' = 1, 'S' = 2)
train$Sex <- ifelse(train$Sex == 'male', 1, 0)
test$Sex <- ifelse(test$Sex == 'male', 1, 0)
#fix(train)
#logistic regression
attach(train)
LR
             glm(
                      factor(Survived)~factor(Pclass)
                                                                Sex+
                                                                         Age+SibSp+
Parch+factor(Embarked), data= train, family= 'binomial')
LR.probs = predict(LR, type= 'response', newdata= test)
LR.probs
LR.probs = ifelse(LR.probs > 0.5, 1, 0)
table(LR.probs,test$Survived)
mean(LR.probs == test$Survived)
#LDA
library(MASS)
                                        factor(Pclass)
                                                                         Age+SibSp+
LDA
               lda(factor(Survived)~
                                                                Sex+
                                                          +
Parch+factor(Embarked) , data= train)
LDA.pred = predict(LDA, newdata= test)
LDA.class = LDA.pred\$class
LDA.class
```

```
table(LDA.class, test$Survived)
mean(LDA.class != test$Survived)
#QDA
library(MASS)
QDA = qda(factor(Survived)~factor(Pclass) + Sex+ Age+SibSp+ Parch +
factor(Embarked), data= train, family= binomial)
QDA.pred = predict(QDA, newdata= test)
QDA.class = QDA.pred$class
QDA.class
table(QDA.class, test$Survived )
mean(QDA.class != test$Survived)
.#KNN
library(FNN)
#train$Age <- ifelse(train$Age<18, 1, 0)</pre>
#test$Age <- ifelse(test$Age<18, 1, 0)</pre>
train$Age <- ifelse(train$Age<18, 1, 0)</pre>
test$Age <- ifelse(test$Age<18, 1, 0)
train1= train[, -c(1,2,4,9,10,11)]
test1 = test[, -c(1,2,4,9,10,11)]
train.Y = train$Survived
```

```
train1
set.seed(12)
knn1 <- knn(train1, test1, train.Y, k=1)
table(knn1, test$Survived)
mean(knn1 != test$Survived)
knn2 = knn(train1, test1, train.Y, k=2)
table(knn2, test$Survived)
mean(knn2 != test$Survived)
knn3 = knn(train1, test1, train.Y, k=3)
table(knn3, test$Survived)
mean(knn3 != test$Survived)
knn4 = knn(train1, test1, train.Y, k=4)
table(knn4, test$Survived)
mean(knn4 != test$Survived)
knn5 = knn(train1, test1, train.Y, k=5)
table(knn5, test$Survived)
mean(knn5!= test$Survived)
#question 2
train = read.csv("C:\fy\text{YUsers\fy\text{YKIRAN KONDISETTI\fy\text{Pesktop\fy\fy\text{Train.csv"}}}
test = read.csv('C:\forall YUsers\forall KIRAN KONDISETTI\forall YDesktop\forall Test.csv')
test$Age[is.na(test$Age)] = mean(test$Age, na.rm=TRUE)
```

```
train$Age[is.na(train$Age)] = mean(train$Age, na.rm=TRUE)
levels(train$Embarked)
levels(train$Embarked)[1] = 'S'
install.packages('naniar')
library(naniar)
levels(test$Cabin)
levels(test$Cabin)[1] = 'NA'
attach(test)
               as.formula('factor(Survived)~factor(Pclass)
                                                                Sex+ Age+SibSp+
Parch+factor(Embarked)+Cabin')
LR_C = glm(formula1,data= test, family='binomial')
LR\_C.probs = predict(LR\_C)
LR C.probs = ifelse(LR C.probs > 0.5, 1, 0)
mean(LR_C.probs!= test$Survived)
                 glm(factor(Survived)~factor(Pclass)
                                                                       Age+SibSp+
LR C1
                                                              Sex+
                                                        +
Parch+factor(Embarked),data= test, family= 'binomial')
LR_C1.probs = predict(LR_C1)
LR_C1.probs = ifelse(LR_C1.probs > 0.5, 1, 0)
mean(LR_C1.probs!= test$Survived)
```

```
#question3
install.packages('vcd')
library(vcd)
attach(train)
#fix(train)
od = glm(factor(Survived)~factor(Pclass) + Sex+ factor(Embarked), data= train, family
= 'binomial')
x= od$coefficients
or = exp(x)
or
#question 4
               as.formula('factor(Survived)~factor(Pclass)
                                                            + Sex+ Age+SibSp+
Parch+factor(Embarked)')
LR_0.5 = glm(formula, data= train, family= 'binomial')
LR.probs_0.5 = predict(LR_0.5, type= 'response', newdata= test)
LR.probs_0.5 = ifelse(LR.probs_0.5 > 0.5, 1, 0)
table(LR.probs_0.5,test$Survived)
mean(LR.probs_0.5!= test$Survived)
LR_0.2 = glm(formula, data= train, family= 'binomial')
LR.probs_0.2 = predict(LR_0.2, type= 'response', newdata= test)
```

```
LR.probs_0.2 = ifelse(LR.probs_0.2 > 0.2, 1, 0)
table(LR.probs_0.2,test$Survived)
mean(LR.probs_0.2!= test$Survived)
LR_0.8= glm(formula, data= train, family= 'binomial')
LR.probs_0.8 = predict(LR_0.8, type= 'response', newdata= test)
LR.probs_0.8 = ifelse(LR.probs_0.8 > 0.8, 1, 0)
table(LR.probs_0.8,test$Survived)
mean(LR.probs_0.8!= test$Survived)
#question 5
install.packages('ROCR')
library(ROCR)
pred= predict(LR, type= 'response', newdata= test)
pr = prediction(pred,test$Survived)
prf<- performance(pr, measure='tpr', x.measure='fpr')</pre>
plot(prf)
abline(0,1)
prf
auc_ROCR <- performance(pr, measure = "auc")</pre>
auc<- auc_ROCR@y.values[[1]]</pre>
```

```
#question 6
library(FNN)
train = read.csv("C:\fy\text{YUsers}\fy\text{KIRAN KONDISETTI}\fy\text{Desktop}\fy\text{Train.csv}")
test = read.csv('C:\forall YUsers\forall XIRAN KONDISETTI\forall YDesktop\forall Test.csv')
test$Age[is.na(test$Age)] = mean(test$Age, na.rm=TRUE)
train$Age[is.na(train$Age)] = mean(train$Age, na.rm=TRUE)
levels(train$Embarked)
levels(train$Embarked)[1] = 'S'
train$Embarked <- sapply(as.character(train$Embarked), switch, 'C' = 0, 'Q' = 1, 'S' = 2)
test \$ Embarked <- \ sapply (as.character(test \$ Embarked), \ switch, \ 'C' = 0, \ 'Q' = 1, \ 'S' = 2)
train$Sex <- ifelse(train$Sex == 'male', 1, 0)
test$Sex <- ifelse(test$Sex == 'male', 1, 0)
LR 7
                  glm(factor(Survived)~factor(Pclass)
                                                                     Sex+
                                                                               Age+SibSp+
Parch+factor(Embarked)+Parch+Fare,data = train, family= 'binomial')
summary(LR_7)
train2 = train[, -c(1,2,4,7,9,10,11,12)]
test2 = test[, -c(1,2,4,7,9,10,11,12)]
train.Y1= train$Survived
train1
```

```
set.seed(143)
knn11 <- knn(train2, test2, train.Y1, k=1)
table(knn11, test$Survived)
mean(knn11 != test$Survived)
knn21 = knn(train2, test2, train.Y1, k=2)
table(knn21, test$Survived)
mean(knn21 != test$Survived)
knn31 = knn(train2, test2, train.Y1, k=3)
table(knn31, test$Survived)
mean(knn31 != test$Survived)
knn41 = knn(train2, test2, train.Y1, k=4)
table(knn41, test$Survived)
mean(knn41 != test$Survived)
knn51 = knn(train2, test2, train.Y1, k=5)
table(knn51, test$Survived)
mean(knn51 != test$Survived)
```