Titanic Survival Statistics

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Open library

```
library(titanic)
## Warning: package 'titanic' was built under R version 4.2.2
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
      filter, lag
## The following objects are masked from 'package:base':
##
##
      intersect, setdiff, setequal, union
library(tidyverse)
## -- Attaching packages ------ 1.3.2 --
## v ggplot2 3.3.6 v purrr
                           0.3.4
## v tibble 3.1.8 v stringr 1.4.1
                  v forcats 0.5.2
## v tidyr 1.2.1
## v readr
          2.1.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
Drop NA
titanic_train <- na.omit(titanic_train)</pre>
```

Split Data

```
set.seed(42)

n <- nrow(titanic_train)
id <- sample(1:n, size= n*0.7)
train_data <- titanic_train[id, ]
test_data <- titanic_train[-id, ]</pre>
```

Create model

```
model <- glm(Survived ~ Pclass + Age + Sex + SibSp, data = train_data, family = "binomial")
summary(model)
##
## Call:
## glm(formula = Survived ~ Pclass + Age + Sex + SibSp, family = "binomial",
      data = train_data)
##
## Deviance Residuals:
      Min
               1Q
                    Median
                                  3Q
                                         Max
## -2.9097 -0.6223 -0.3378 0.6033
                                       2.4133
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
                         0.69789 8.907 < 2e-16 ***
## (Intercept) 6.21585
              -1.49165
## Pclass
                          0.17867 -8.349 < 2e-16 ***
             -0.05004
                       0.01031 -4.851 1.23e-06 ***
## Age
## Sexmale
             -2.82454
                       0.27045 -10.444 < 2e-16 ***
                       0.15504 -2.616 0.00889 **
             -0.40560
## SibSp
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 673.56 on 498 degrees of freedom
##
## Residual deviance: 424.77 on 494 degrees of freedom
## AIC: 434.77
##
## Number of Fisher Scoring iterations: 5
```

Train Data

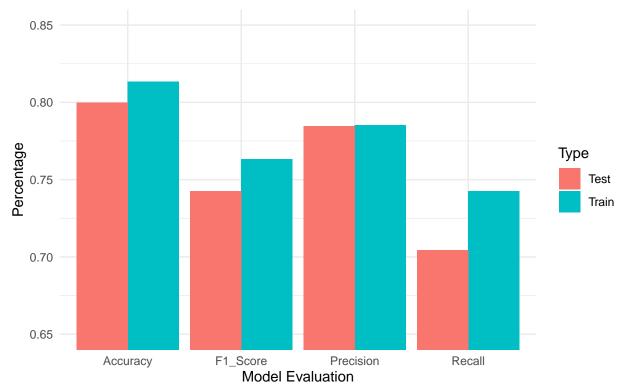
```
##Model evaluation
train_a \leftarrow (conM[1,1] + conM[2,2]) / sum(conM)
train_p \leftarrow conM[2,2] / (conM[2,2] + conM[2,1])
train_r \leftarrow conM[2,2]/(conM[1,2]+conM[2,2])
cat("Acculacy:",train_a,"\nPrecision:",train_p,"\nRecall",train_r)
## Acculacy: 0.8136273
## Precision: 0.7853403
## Recall 0.7425743
##Train F1
Train_f1 <- 2*((train_p*train_r)/(train_p+train_r))</pre>
cat("\nTrain F1 :", Train_f1)
## Train F1 : 0.7633588
Test Data
test_prob_su <- predict(model, newdata = test_data, type = "response")</pre>
test_data$pred_su <- ifelse(test_prob_su >= 0.5,1,0)
##Confuionmetric
conM1 <- table(test_data$pred_su, test_data$Survived,</pre>
               dnn = c("Predicted", "Actuall"))
##Model evaluation
test_a \leftarrow (conM1[1,1] + conM1[2,2]) / sum(conM1)
test_p \leftarrow conM1[2,2] / (conM1[2,2] + conM1[2,1])
test_r \leftarrow conM1[2,2]/(conM1[1,2]+conM1[2,2])
cat("Acculacy:",test_a,"\nPrecision:",test_p,"\nRecall",test_r)
## Acculacy: 0.8
## Precision: 0.7848101
## Recall 0.7045455
Test_f1 <- 2*((test_p*test_r)/(test_p+test_r))</pre>
cat("\nTest F1 :", Test_f1)
```

Plot model

Test F1 : 0.742515

Plot data

Train & Test Model Evaluation



Source: titanic from titanic package

The bar chart shows that the model performs better with train data. We can prove this theory based on the F1 score.