# **Logistic Regression**

## Titanic\_train dataset 📇

#### Updating HTML index of packages in '.Library'

### Clean Data -> Drop NA

```
titanic_train <- na.omit(titanic_train)
```

```
#split data
set.seed(99)
n <- nrow(titanic_train)
id <- sample(1:n, size=n*0.5) #50% train 50% test
train_data <- titanic_train[id,]
test_data <- titanic_train[-id,]</pre>
```

#### **Create Predicted and Evaluate Model**

```
#Train model
train_model <- qlm(Survived~Pclass + Age + Sex + SibSp,
                     data = train_data,
                     family = "binomial")
prob_train <- predict(train_model, type = "response")</pre>
train_data$pred_Survived <- ifelse(prob_train>=0.5,1,0)
##test model
prob_test <- predict(train_model, newdata = test_data, type = "response")</pre>
test_data$pred_Survived <- ifelse(prob_test >=0.5,1,0)
##confusion metrix
con_metrix_train <- table(train_data$pred_Survived,train_data$Survived,</pre>
                     dnn = c("Predicted", "Actual"))
con_metrix_test <- table(test_data$pred_Survived,test_data$Survived,</pre>
                     dnn = c("Predicted", "Actual"))
##Model Evaluation Train
acc_train <-(con_metrix_train[1,1]+con_metrix_train[2,2])/</pre>
              sum(con_metrix_train)
precision_train <- (con_metrix_train[2,2]/</pre>
               (con_metrix_train[2,1]+con_metrix_train[2,2]))
recall_train <- (con_metrix_train[2,2]/</pre>
               (con_metrix_train[1,2]+con_metrix_train[2,2]))
f1_train <- 2*((precision_train*recall_train)/(precision_train+recall_train))
##Model Evaluation Test
acc_test <- (con_metrix_test[1,1]+con_metrix_test[2,2])/sum(con_metrix_test)</pre>
precision_test <- (con_metrix_test[2,2]/(con_metrix_test[2,1]+con_metrix_test[2,2]</pre>
recall_test <- (con_metrix_test[2,2]/(con_metrix_test[1,2]+con_metrix_test[2,2]))</pre>
f1_test <- 2*((precision_test*recall_test)/(precision_test+recall_test))
```

```
Model_name = c("Train model", "Test model"),
 Accuracy = c(acc_train,acc_test),
 Precision = c(precision_train,precision_test),
 Recall = c(recall_train, recall_test),
 F1 = c(f1\_train, f1\_test)
)
cat("Hypothesis test of Multiple Regression:\n")
print(summary(train_model))
cat("-----\n",
"Label of Values: 1 was Survived, 0 was Died\n", "Confusion Matrix of Train:\n")
print(con_metrix_train)
cat("Confusion Matrix of Test:\n")
print(con_metrix_test)
cat("Accuracy:\n")
print(df_accuracy)
graph <- df_accuracy %>%
 qather(Accuracy:F1,
        key = "type",
        value = "RMSE")
ggplot(graph, aes(x = type, y = RMSE, color = Model_name, group = Model_name)) +
 geom\_line(size = 4) +
 coord\_cartesian(ylim = c(0.73, 0.83)) +
 scale_color_manual(values = c("#F5C4C4","#ED9591")) +
 theme_minimal()+
 labs(title = "Model Evaluation",x = "Type of Accuracy",y = "Value")
Hypothesis test of Multiple Regression:
Call:
glm(formula = Survived ~ Pclass + Age + Sex + SibSp, family = "binomial",
   data = train_data)
Deviance Residuals:
   Min
             1Q
                 Median
                             3Q
                                     Max
-2.1447 -0.6103 -0.3305 0.6229
                                  2.6132
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
(Intercept) 6.87162 0.87671 7.838 4.58e-15 ***
          -1.56949 0.22642 -6.932 4.16e-12 ***
Pclass
           -0.06410 0.01214 -5.281 1.28e-07 ***
Age
Sexmale
          -2.72375 0.30931 -8.806 < 2e-16 ***
           SibSp
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

df\_accuracy <- data.frame(</pre>

