

R Notebook

Imports

```
library("dplyr")
```

```
## Warning: replacing previous import 'lifecycle::last_warnings' by  
## 'rlang::last_warnings' when loading 'pillar'
```

```
##  
## 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("DataExplorer")
```

```
## Warning: package 'DataExplorer' was built under R version 4.1.3
```

```
library("caret")
```

```
## Warning: package 'caret' was built under R version 4.1.3
```

```
## Loading required package: ggplot2
```

```
## Loading required package: lattice
```

```
library("randomForest")
```

```
## Warning: package 'randomForest' was built under R version 4.1.3
```

```
## randomForest 4.7-1.1
```

```
## Type rfNews() to see new features/changes/bug fixes.
```

```
##  
## Attaching package: 'randomForest'
```

```
## The following object is masked from 'package:ggplot2':  
##  
##     margin
```

```
## The following object is masked from 'package:dplyr':  
##  
##     combine
```

```
library("Hmisc")
```

```
## Warning: package 'Hmisc' was built under R version 4.1.3
```

```
## Loading required package: survival
```

```
##  
## Attaching package: 'survival'
```

```
## The following object is masked from 'package:caret':  
##  
##     cluster
```

```
## Loading required package: Formula
```

```
##  
## Attaching package: 'Hmisc'
```

```
## The following objects are masked from 'package:dplyr':  
##  
##     src, summarize
```

```
## The following objects are masked from 'package:base':  
##  
##     format.pval, units
```

```
library("car")
```

```
## Warning: package 'car' was built under R version 4.1.3
```

```
## Loading required package: carData
```

```
## Warning: package 'carData' was built under R version 4.1.3
```

```
##  
## Attaching package: 'car'
```

```
## The following object is masked from 'package:dplyr':  
##  
##     recode
```

```
library("glmnet")
```

```
## Warning: package 'glmnet' was built under R version 4.1.3
```

```
## Loading required package: Matrix
```

```
## Warning: package 'Matrix' was built under R version 4.1.3
```

```
## Loaded glmnet 4.1-4
```

```
library("pROC")
```

```
## Warning: package 'pROC' was built under R version 4.1.3
```

```
## Type 'citation("pROC")' for a citation.
```

```
##
```

```
## Attaching package: 'pROC'
```

```
## The following objects are masked from 'package:stats':
```

```
##
```

```
##      cov, smooth, var
```

```
library("gbm")
```

```
## Warning: package 'gbm' was built under R version 4.1.3
```

```
## Loaded gbm 2.1.8.1
```

Setting Up

```
df_train <- read.csv("joint_numeric.csv", header = TRUE , na.strings = c("na", "NA"),  
                    stringsAsFactors = FALSE, sep = ",")
```

```
df_test <- read.csv("joint_test_numeric.csv", header = TRUE , na.strings = c("na", "NA"),  
                   stringsAsFactors = FALSE, sep = ",")
```

```
df_test[is.na(df_test)] = 0
```

```
df_train$X<-NULL
```

```
df_test$X<-NULL
```

```
df_train$fraudulent<-as.factor(df_train$fraudulent)
```

```
df_test$fraudulent<-as.factor(df_test$fraudulent)
```

```
df_train$department_n_first_personp<-NULL
```

```
#colnames(df_train)[colSums(is.na(df_train)) > 0]
```

```
#colnames(df_test)[colSums(is.na(df_test)) > 0]
```

```
df_train = subset(df_train, select = c('has_company_logo', 'has_questions', 'fraudulent', 'has_department'))
df_test = subset(df_test, select = c('has_company_logo', 'has_questions', 'fraudulent', 'has_department'))

##IMPORTANTTTT
train_df <- df_train %>% select_if(function(col) length(unique(col))>1)
test_df <- df_test %>% select_if(function(col) length(unique(col))>1)
```

Original Logistic Regression Model

```
# Log reg with everything
set.seed(123)
fraud_glm0 <- glm(fraudulent~., family=binomial, data=df_train)
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
#summary(fraud_glm0)
```

```
#originally got this warning
```

```
#Warning: glm.fit: algorithm did not converge
```

```
#Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
# I found out it was due to ... Singularity means that your predictor variables are linearly dependent,
```

Correlation Plots

```
without_df = subset(train_df, select = -c(fraudulent))
```

```
#sum(is.na(train_df))
```

```
res2 <- rcorr(as.matrix(train_df))
```

```
corrdisp <- cor(without_df, method="s")
```

```
# find indices of highly correlated attributes
```

```
highlycorrelated <- findCorrelation(corrdisp, cutoff= 0.98)
```

```
#highlycorrelated
```

```
#count(highlycorrelated) --> 69 (with 0.5)
```

```
#without_df[highlycorrelated]
```

```
edit2_df = subset(train_df, select = -c(highlycorrelated))
```

```
edit3_df = subset(test_df, select= -c(highlycorrelated))
```

Revised Logistic

```
# from the correlation plot I noticed the below columns has a perfect colinearity so I got rid of one of them
set.seed(123)
```

```
#This one surpasses all warnings after ridding of all perfect colinearity
```

```
fraud_glm5 <- glm(fraudulent~., family=binomial, data=edit2_df)
```

```
#summary(fraud_glm5)
```

```
# Revising model before testing
fraud_glm6 = glm(fraudulent~
  . - region_cat_SW - has_industry -department_sent_vader
  - department_n_second_personp -company_profile_n_chars
  - company_profile_n_hashtags -company_profile_n_uq_words
  -company_profile_n_first_person -company_profile_n_prepositions
  -department_n_charsperword -description_n_nonasciis -description_sent_afinn
  -description_sent_vader -description_n_first_person -description_n_first_personp
  -requirements_n_hashtags -requirements_sent_vader -requirements_n_second_personp
  - benefits_sent_bing -benefits_sent_syuzhet -industry_top -employment_type_Full.time
  -title_customer -title_teacher -title_assistant -required_experience_Entry.level
  -required_experience_Mid.Senior.level -fn_Finance -fn_Production
  - required_education_Professional - has_department ,
  data = edit2_df, family = binomial)
#summary(fraud_glm6)
```

Analysis of Model

```
predictTrain = predict(fraud_glm6, type = "response")
table(edit2_df$fraudulent, predictTrain >= 0.5)
```

```
##
##      FALSE  TRUE
##  0 13518    86
##  1   476   224
```

```
accuracy = (244 + 13518) / nrow(edit2_df)
sensitivity = 244 / (244 + 476)
specificity = 13518 / (13518 + 86)
#sensitivity
#specificity

cat("accuracy: ", accuracy)
```

```
## accuracy: 0.9621085
```

```
threshold=0.5
predicted_values<-ifelse(predict(fraud_glm6,type="response")>threshold,1,0)
actual_values<-fraud_glm6$y
conf_matrix<-table(predicted_values,actual_values)
#conf_matrix
#sensitivity(conf_matrix)
#specificity(conf_matrix)
```

Applying Test

```
predictTest = predict(fraud_glm6, type = "response", newdata = edit3_df)

# no preference over error t = 0.5
edit3_df$fraudulent = as.numeric(predictTest >= 0.5)
table(edit3_df$fraudulent)
```

```
##
##      0      1
## 3508   68
```

```
predicted_probabilities <- predict(fraud_glm5,
                                   newdata=edit2_df,
                                   type="response")

class_preds <- ifelse(predicted_probabilities >= 0.5, 1, 0)

# Make a table of predictions vs. actual
result_table <- table(class_preds,
                      edit2_df$fraudulent)

#result_table

confusionMatrix(data = factor(class_preds),
                 reference = factor(edit2_df$fraudulent),
                 positive = "1")
```

```
## Confusion Matrix and Statistics
##
##              Reference
## Prediction      0      1
##              0 13508   447
##              1    96   253
##
##              Accuracy : 0.962
##              95% CI : (0.9588, 0.9651)
##              No Information Rate : 0.9511
##              P-Value [Acc > NIR] : 1.519e-10
##
##              Kappa : 0.4649
##
##  Mcnemar's Test P-Value : < 2.2e-16
##
##              Sensitivity : 0.36143
##              Specificity : 0.99294
##              Pos Pred Value : 0.72493
##              Neg Pred Value : 0.96797
##              Prevalence : 0.04894
##              Detection Rate : 0.01769
##              Detection Prevalence : 0.02440
##              Balanced Accuracy : 0.67719
##
##              'Positive' Class : 1
##
```