Data Exploration

Andy Shen

10/27/2020

Traditional Techniques

```
rm(list = ls())
library(MASS) #lda, qda
library(class) #knn
library(tidyverse)
library(caret)
find_score <- function(conf_mat) {</pre>
  print(sum(diag(conf_mat) * c(1, 20, 20)))
  final <- sum(diag(conf_mat) * c(1, 20, 20)) / sum(apply(conf_mat, 2, sum) * c(1, 20, 20))
  final
}
setwd("/Users/andyshen/Desktop/Git/Stats-101C-F20/Midterm Project")
train <- read.csv("training.csv", stringsAsFactors = TRUE)</pre>
test <- read.csv("test.csv", stringsAsFactors = TRUE)</pre>
train$class <- factor(train$class)</pre>
levels(train$class) <- c("NG", "OG", "TSG")</pre>
test$class <- factor(test$class)</pre>
levels(test$class) <- c("NG", "OG", "TSG")</pre>
set.seed(5732)
samp <- sample(1:nrow(train), floor(0.8 * nrow(train)), replace = FALSE)</pre>
train1 <- train[samp, ]</pre>
test_train <- train[-samp, ]</pre>
```

 $Family Member Count, RVIS_percentile, N_Missense, intolerant_pNull, Gene_age, pLOF_Z score \\VEST_score$

LDA

```
lda.mod <- lda(
  class ~ FamilyMemberCount + RVIS_percentile + N_Missense +
    intolerant_pNull + Gene_age + pLOF_Zscore, data = train1
)
preds <- predict(lda.mod, test_train, type = "response")$posterior
preds <- apply(preds, 1, which.max) - 1
tbl <- table(preds, test_train$class)
ter <- sum(diag(tbl)) / sum(tbl)
tbl</pre>
```

```
## preds NG OG TSG
## 0 559 21 22
## 1 2 5 0
## 2 10 4 13
```

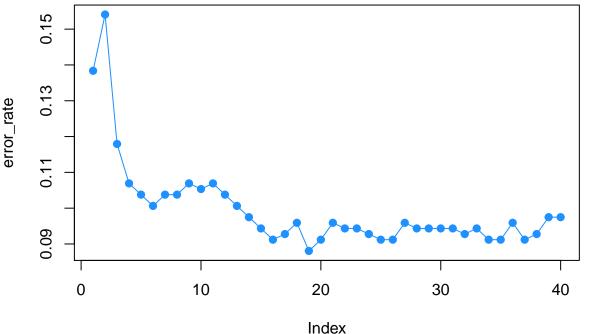
Test error rate is 0.093.

QDA

Test error rate is 0.164

```
qda.mod <- qda(
 class ~ FamilyMemberCount + RVIS_percentile + N_Missense +
    intolerant_pNull + Gene_age + pLOF_Zscore, data = train1
)
preds <- predict(qda.mod, test_train, type = "response")$posterior</pre>
preds <- apply(preds, 1, which.max) - 1</pre>
tbl <- table(preds, test_train$class)</pre>
ter <- sum(diag(tbl)) / sum(tbl)</pre>
tbl
##
## preds NG OG TSG
##
       0 503 12 10
##
       1 35 14 10
##
       2 33 4 15
```

```
KNN
train1k <- train1 %>% dplyr::select(-class)
test_traink <- test_train %>% dplyr::select(-class)
for(col in 1:ncol(train1k)) {
  train1k[, col] <- train1k[, col] / max(train1k[, col])</pre>
  test_traink[, col] <- test_traink[, col] / max(test_traink[, col])</pre>
} #standardizing. ncol(test) == ncol(train)
trainx <- train1k %>% dplyr::select(
  FamilyMemberCount, RVIS_percentile, N_Missense, intolerant_pNull, Gene_age, pLOF_Zscore
trainy <- train1$class</pre>
testx <- test_traink %>% dplyr::select(
  FamilyMemberCount, RVIS_percentile, N_Missense, intolerant_pNull, Gene_age, pLOF_Zscore
testy <- test_train$class</pre>
rows <- 40
knn_mat <- matrix(NA, nrow = rows, ncol = length(testy))</pre>
error_rate <- rep(NA, rows)
for(i in 1:rows) {
  knn_mat[i,] <- knn(trainx, testx, trainy, k = i)</pre>
  tbl <- table("actual" = testy, "predicted" = knn_mat[i,])</pre>
  error_rate[i] <- 1 - (sum(diag(tbl)) / sum(tbl))</pre>
}
plot(error_rate, type = "1", col = "dodgerblue")
points(error_rate, pch = 19, col = "dodgerblue")
     0.15
```



```
best <- which.min(error_rate)</pre>
best_tbl <- table("actual" = testy, "predicted" = knn_mat[best,])</pre>
```

best_tbl predicted ## 3 ## actual 1 2 ## NG 566 3 2 ## OG 23 1 TSG 24 8 ## 3 ter <- sum(diag(best_tbl)) / sum(best_tbl)</pre>

Test error rate is 0.088. This is for K=19.

Using caret

```
tc <- trainControl(
  method = "cv", number = 5, classProbs = TRUE, savePredictions = TRUE
) # setting up training technique</pre>
```

```
LDA
LDAfit <- caret::train(</pre>
  class ~ FamilyMemberCount + RVIS_percentile + N_Missense +
    intolerant_pNull + Gene_age + pLOF_Zscore,
  data = train1, method = "lda",
  preProc = c("center", "scale"),
  trControl = tc
preds <- predict(LDAfit, test_train)</pre>
tbl <- table(preds, test_train$class)</pre>
acc <- find_score(tbl)</pre>
## [1] 919
tbl
##
## preds NG OG TSG
     NG 559 21 22
##
##
     OG
         2 5
                  0
##
     TSG 10
              4 13
acc
## [1] 0.4911812
QDA
QDAfit <- caret::train(</pre>
  class ~ FamilyMemberCount + RVIS_percentile + N_Missense +
    intolerant_pNull + Gene_age + pLOF_Zscore,
 data = train1, method = "qda",
 preProc = c("center", "scale"),
```

```
class ~ FamilyMemberCount + RVIS_percentile + N_Missense +
   intolerant_pNull + Gene_age + pLOF_Zscore,
   data = train1, method = "qda",
   preProc = c("center", "scale"),
   trControl = tc
)
preds <- predict(QDAfit, test_train)
tbl <- table(preds, test_train$class)
acc <- find_score(tbl)
## [1] 1083</pre>
```

```
## ## preds NG OG TSG ## NG 503 12 10 ## OG 35 14 10 ## TSG 33 4 15
```

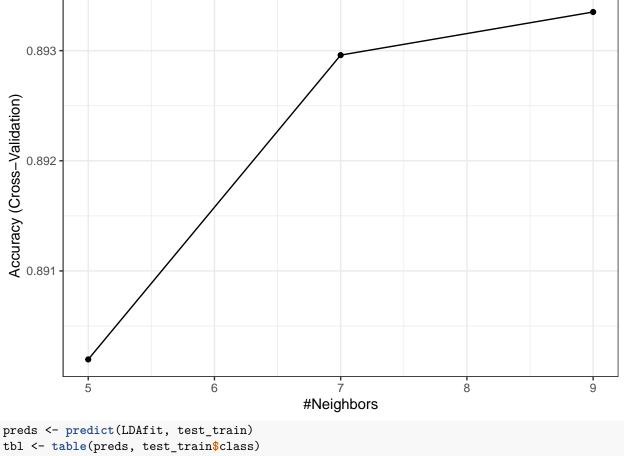
```
acc
```

[1] 0.5788348

Logistic Regression

```
LRfit <- caret::train(</pre>
  class ~ FamilyMemberCount + RVIS_percentile + N_Missense +
    intolerant_pNull + Gene_age + pLOF_Zscore,
  data = train1, method = "glm", family = "binomial",
  preProc = c("center", "scale"),
  trControl = tc
## Warning: model fit failed for Fold1: parameter=none Error in method$fit(x = x, y = y, wts = wts, par
    glm models can only use 2-class outcomes
\#\# Warning: model fit failed for Fold2: parameter=none Error in method$fit(x = x, y = y, wts = wts, parameter=none)
    glm models can only use 2-class outcomes
## Warning: model fit failed for Fold3: parameter=none Error in method$fit(x = x, y = y, wts = wts, par
    glm models can only use 2-class outcomes
## Warning: model fit failed for Fold4: parameter=none Error in method$fit(x = x, y = y, wts = wts, par
    glm models can only use 2-class outcomes
## Warning: model fit failed for Fold5: parameter=none Error in method$fit(x = x, y = y, wts = wts, par
    glm models can only use 2-class outcomes
## Warning in nominalTrainWorkflow(x = x, y = y, wts = weights, info = trainInfo, :
## There were missing values in resampled performance measures.
## Something is wrong; all the Accuracy metric values are missing:
##
       Accuracy
                     Kappa
## Min.
         : NA
                 Min. : NA
## 1st Qu.: NA
                 1st Qu.: NA
## Median : NA
                 Median : NA
## Mean
         :NaN
                 Mean
                       :NaN
## 3rd Qu.: NA
                 3rd Qu.: NA
## Max. : NA
                 Max. : NA
## NA's :1
                 NA's
## Error: Stopping
KNN
```

```
KNNfit <- caret::train(
  class ~ FamilyMemberCount + RVIS_percentile + N_Missense +
    intolerant_pNull + Gene_age + pLOF_Zscore, data = train1,
    method = "knn", preProc = c("center", "scale"), trControl = tc
)
ggplot(KNNfit) + theme_bw()</pre>
```



```
tbl <- table(preds, test_train$class)
acc <- find_score(tbl)</pre>
## [1] 919
tbl
##
                 OG TSG
## preds NG
##
      NG
          559
                 21
                      22
      OG
             2
                  5
##
##
      TSG
           10
                  4
                      13
```

[1] 0.4911812

acc