## White\_MLR\_Final.R

## nebojsahrnjez

## 2021-12-01

```
library(readr)
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 ------ tidyverse 1.3.1 --
## v ggplot2 3.3.5 v purrr 0.3.4
## v tibble 3.1.6 v stringr 1.4.0
## v tidyr 1.1.4 v forcats 0.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
library(corrplot)
## corrplot 0.92 loaded
library(moments)
library(car)
## Loading required package: carData
## Attaching package: 'car'
```

```
## The following object is masked from 'package:purrr':
##
##
      some
## The following object is masked from 'package:dplyr':
##
##
      recode
library(ggplot2)
library(ggrepel)
library(gridExtra)
## Attaching package: 'gridExtra'
## The following object is masked from 'package:dplyr':
##
      combine
#Libraries from exploratory analysis
library(cvTools)
## Loading required package: lattice
## Loading required package: robustbase
library(MASS)
##
## Attaching package: 'MASS'
## The following object is masked from 'package:dplyr':
##
##
      select
#Libraries for this script
white <- read_csv("winequality-white.csv")</pre>
## Rows: 4898 Columns: 12
## -- Column specification ------
## Delimiter: ","
## dbl (12): fixed acidity, volatile acidity, citric acid, residual sugar, chlo...
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
sum(is.na(white))
## [1] 0
white <- na.omit(white)</pre>
#Reading in the data
dfw <- as.data.frame(white)</pre>
dfw \leftarrow dfw[-2782,]
dfw2 <- subset(dfw, select = -density)</pre>
#Creating the dataframes to be used
MLR.results <- data.frame(matrix(ncol=2,nrow=0,</pre>
                  dimnames=list(NULL, c("Model", "Classification Accuracy %"))))
#Empty data frame for results
set.seed(100)
test <- sample(1:nrow(dfw), size = nrow(dfw)/5)</pre>
train <- (-test)</pre>
#Training and Test sets for dfw
dfw.train <- dfw[train,]</pre>
dfw.test <- dfw[test,]</pre>
#Create a test and training data set, 20/80 split
w.mlm.train <- lm(
  quality ~.,
  data = dfw.train
#Create linear regression with all predictors using the training dataset
w.mlm.predict <- predict(w.mlm.train, newdata = dfw.test)</pre>
w.mlm.predict.rounded <- round(w.mlm.predict, digits = 0)</pre>
#Round the predicted to a integer so it can be compared to the test set for
# classification
(con_mat <- table(w.mlm.predict.rounded, dfw.test$quality))</pre>
##
## w.mlm.predict.rounded 3 4
                                    5 6
                                                 8
                               2
                                   1
                                        0
##
                           0
                                             Ω
##
                        5
                            1 16 128 52
                        6
##
                           3 18 157 357 131 20
##
                        7 0 0 3 32 38 14
```

```
w.mlm.acc <- round(mean(w.mlm.predict.rounded==dfw.test$quality)*100,digits = 2)</pre>
#Create the confusion matrix and calculate the proportion correct
MLR.results[1,] <- c("dfw MLR w/ Train/Test", w.mlm.acc)</pre>
#dfw train/test multi linear regression
test <- sample(1:nrow(dfw2), size = nrow(dfw2)/5)</pre>
train <- (-test)
dfw2.train <- dfw2[train,]</pre>
dfw2.test <- dfw2[test,]</pre>
#Create a test and training data set, 20/80 split
w.mlm.train2 <- lm(
  quality ~.,
  data = dfw2.train
#Create linear regression with all predictors using the training dataset
w.mlm.predict2 <- predict(w.mlm.train2, newdata = dfw2.test)</pre>
w.mlm.predict.rounded2 <- round(w.mlm.predict2, digits = 0)</pre>
#Round the predicted to a integer so it can be compared to the test set for
# classification
(con_mat <- table(w.mlm.predict.rounded2, dfw2.test$quality))</pre>
##
## w.mlm.predict.rounded2
                            3
                                 4
                                     5
                                          6
                                              7
                                                  8
                                                      9
                             0
                                      3
                                         1
##
                             0 12 135 59
                                              2
                                                      0
##
                             4 18 141 340 134 17
                         6
##
                                                       2
                                     5 52 46
                                                  5
w.mlm.acc2 <- round(mean(w.mlm.predict.rounded2==dfw2.test$quality)*100,
                     digits = 2)
#Create the confusion matrix and calculate the proportion correct
MLR.results[2,] <- c("dfw2 MLR w/ Train/Test", w.mlm.acc2)
#dfw2 train/test multi linear regression
k <- 10 #number of folds
folds <- cvFolds(nrow(dfw), K=k)</pre>
folds2 <- cvFolds(nrow(dfw2), K=k)</pre>
w.mlm.cv.class <- matrix(NA,k,1, dimnames=list(NULL, paste(1)))</pre>
w.mlm.cv.class2 <- matrix(NA,k,1, dimnames=list(NULL, paste(1)))</pre>
```

```
#Preparing both datasets for cross-validation
for(i in 1:k){
  tr.mlr <- dfw[folds$subsets[folds$which != i],]</pre>
  te.mlr <- dfw[folds$subsets[folds$which == i],]</pre>
  w.mlm <- lm(quality~., data = tr.mlr)</pre>
  w.mlm.pred <- predict(w.mlm, newdata = te.mlr)</pre>
  w.mlm.cv.class[i] <- mean(round(w.mlm.pred, digits = 0)==te.mlr$quality)</pre>
w.mlm.cv.class
##
## [1,] 0.4959184
## [2,] 0.5040816
## [3,] 0.5469388
## [4,] 0.5285714
## [5,] 0.5265306
## [6,] 0.5061224
## [7,] 0.5387755
## [8,] 0.5235174
## [9,] 0.5092025
## [10,] 0.4989775
w.mlm.cv.class <- mean(w.mlm.cv.class)</pre>
print(paste("The average outputs correctly predicted is",
            round(w.mlm.cv.class*100,digits =2),"%",sep=" "))
## [1] "The average outputs correctly predicted is 51.79 %"
MLR.results[3,] <- c("dfw MLR w/ 10-fold CV", round(w.mlm.cv.class*100,
                                                       digits=2))
#dfw cross-validated Multiple Linear Regression
for(i in 1:k){
  tr.mlr2 <- dfw2[folds2$subsets[folds2$which != i],]</pre>
  te.mlr2 <- dfw2[folds2$subsets[folds2$which == i],]</pre>
  w.mlm2 <- lm(quality~., data = tr.mlr2)</pre>
  w.mlm.pred2 <- predict(w.mlm2, newdata = te.mlr2)</pre>
  w.mlm.cv.class2[i] <- mean(round(w.mlm.pred2, digits = 0)==te.mlr2$quality)</pre>
w.mlm.cv.class2
##
## [1,] 0.5163265
```

```
## [2,] 0.5244898
## [3,] 0.5571429
## [4,] 0.5142857
## [5,] 0.4938776
## [6,] 0.4897959
## [7,] 0.5040816
## [8,] 0.5480573
## [9,] 0.5214724
## [10,] 0.4846626
w.mlm.cv.class2 <- mean(w.mlm.cv.class2)</pre>
print(paste("The average outputs correctly predicted is",
            round(w.mlm.cv.class2*100,digits =2), "%", sep=" "))
## [1] "The average outputs correctly predicted is 51.54 %"
MLR.results[4,] <- c("dfw2 MLR w/ 10-fold CV", round(w.mlm.cv.class2*100,
                                                       digits=2))
#dfw2 cross-validated Multiple Linear Regression
dfw$quality <- factor(dfw$quality, ordered = TRUE)</pre>
dfw2$quality <- factor(dfw2$quality, ordered = TRUE)</pre>
#Making the response variable a factor for ordinal logistic regression
test <- sample(1:nrow(dfw), size = nrow(dfw)/5)</pre>
train <- (-test)
dfw.train <- dfw[train,]</pre>
dfw.test <- dfw[test,]</pre>
test <- sample(1:nrow(dfw2), size = nrow(dfw2)/5)</pre>
train <- (-test)</pre>
dfw2.train <- dfw2[train,]</pre>
dfw2.test <- dfw2[test,]</pre>
#Re-create training and test set with new factored response variable
w.olr <- polr(quality~., data = dfw.train, Hess = TRUE)
w.olr.pred <- predict(w.olr, newdata = dfw.test)</pre>
w.olr.pred <- as.numeric(as.character(unlist(w.olr.pred)))</pre>
dfw.test$quality <- as.numeric(as.character(unlist(dfw.test$quality)))</pre>
w.olr.class <- mean(w.olr.pred == dfw.test$quality)</pre>
MLR.results[5,] <- c("dfw OLR w/ Training/Test", round(w.olr.class*100,
                                                        digits=2))
```

#dfw OLR w/ Training/Test Set

```
w.olr2 <- polr(quality~., data = dfw2.train, Hess = TRUE)
w.olr.pred2 <- predict(w.olr2, newdata = dfw2.test)</pre>
w.olr.pred2 <- as.numeric(as.character(unlist(w.olr.pred2)))</pre>
dfw2.test$quality <- as.numeric(as.character(unlist(dfw2.test$quality)))</pre>
w.olr.class2 <- mean(w.olr.pred2 == dfw2.test$quality)</pre>
MLR.results[6,] <- c("dfw2 OLR w/ Training/Test", round(w.olr.class2*100,
                                                           digits=2))
#dfw2 OLR w/ Training/Test Set
folds <- cvFolds(nrow(dfw), K=k)</pre>
folds2 <- cvFolds(nrow(dfw2), K=k)</pre>
w.olr.cv.class <- matrix(NA,k,1, dimnames=list(NULL, paste(1)))</pre>
w.olr.cv.class2 <- matrix(NA,k,1, dimnames=list(NULL, paste(1)))</pre>
\# Re-create the folds and empty classification matrix for OLR
for(i in 1:k){
  tr.olr <- dfw[folds$subsets[folds$which != i],]</pre>
  te.olr <- dfw[folds$subsets[folds$which == i],]</pre>
  w.olr.cv <- polr(quality~., data = tr.olr, Hess = TRUE)</pre>
  w.olr.cv.pred <- predict(w.olr.cv, newdata = te.olr)</pre>
  w.olr.cv.pred <- as.numeric(as.character(unlist(w.olr.cv.pred)))</pre>
  te.olr$quality <- as.numeric(as.character(unlist(te.olr$quality)))</pre>
  w.olr.cv.class[i] <- mean(w.olr.cv.pred==te.olr$quality)</pre>
w.olr.cv.class
##
## [1,] 0.5448980
## [2,] 0.5469388
## [3,] 0.5285714
## [4,] 0.5020408
## [5,] 0.5346939
## [6,] 0.4897959
## [7,] 0.5061224
## [8,] 0.5153374
## [9,] 0.5562372
## [10,] 0.5337423
w.olr.cv.class <- mean(w.olr.cv.class)</pre>
print(paste("The average outputs correctly predicted is",
            round(w.olr.cv.class*100,digits =2),"%",sep=" "))
```

```
## [1] "The average outputs correctly predicted is 52.58 %"
MLR.results[7,] <- c("dfw OLR w/ 10-fold CV", round(w.olr.cv.class*100,
                                                      digits=2))
#dfw OLR w/ 10-fold CV
for(i in 1:k){
  tr.olr2 <- dfw2[folds2$subsets[folds2$which != i],]</pre>
 te.olr2 <- dfw2[folds2$subsets[folds2$which == i],]</pre>
 w.olr.cv2 <- polr(quality~., data = tr.olr2, Hess = TRUE)</pre>
  w.olr.cv.pred2 <- predict(w.olr.cv2, newdata = te.olr2)</pre>
  w.olr.cv.pred2 <- as.numeric(as.character(unlist(w.olr.cv.pred2)))</pre>
 te.olr2$quality <- as.numeric(as.character(unlist(te.olr2$quality)))</pre>
 w.olr.cv.class2[i] <- mean(w.olr.cv.pred2==te.olr2$quality)</pre>
w.olr.cv.class2
##
## [1,] 0.4959184
## [2,] 0.5346939
## [3,] 0.5244898
## [4,] 0.5612245
## [5,] 0.5265306
## [6,] 0.5265306
## [7,] 0.5122449
## [8,] 0.5235174
## [9,] 0.5296524
## [10,] 0.5378323
w.olr.cv.class2 <- mean(w.olr.cv.class2)</pre>
print(paste("The average outputs correctly predicted is",
            round(w.olr.cv.class2*100,digits =2), "%", sep=" "))
## [1] "The average outputs correctly predicted is 52.73 \%"
MLR.results[8,] <- c("dfw2 OLR w/ 10-fold CV", round(w.olr.cv.class2*100,
                                                      digits=2))
#dfw2 OLR w/ 10-fold CV
MLR.results
##
                          Model Classification. Accuracy..
```

53.63

53.32

51.79

## 1

## 2

## 3

dfw MLR w/ Train/Test

dfw MLR w/ 10-fold CV

dfw2 MLR w/ Train/Test

##	4	dfw2 MLR	w/ 10-fold CV	51.54
##	5	dfw OLR w/	Training/Test	51.48
##	6	dfw2 OLR w/	Training/Test	54.34
##	7	dfw OLR	w/ 10-fold CV	52.58
##	8	dfw2 OLR	w/ 10-fold CV	52.73