

Homework 3

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```
library(tidyverse) # data manipulation

## -- Attaching packages ----- tidyverse 1.3.0 --

## v ggplot2 3.3.3      v purrr 0.3.4
## v tibble 3.0.6       v dplyr 1.0.4
## v tidyr 1.1.2        v stringr 1.4.0
## v readr 1.4.0        v forcats 0.5.1

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()

library(dplyr)
library(ISLR) # for data
library(janitor) # clean names

##
## Attaching package: 'janitor'

## The following objects are masked from 'package:stats':
##
##   chisq.test, fisher.test

library(AppliedPredictiveModeling) # better plots
library(caret) # modeling

## Loading required package: lattice

##
## Attaching package: 'caret'

## The following object is masked from 'package:purrr':
##
##   lift

library(corrplot) # correlation plots

## corrplot 0.84 loaded
```

```
library(pROC) # ROC curve
```

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

```
##
```

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

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

```
##
```

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

```
library(MASS) # LDA
```

```
##
```

```
## Attaching package: 'MASS'
```

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

```
##
```

```
##      select
```

```
##(a)
```

```
# Import data
```

```
data(Weekly)
```

```
weekly <-
```

```
  Weekly %>%
```

```
  dplyr::select(-Today)
```

```
# Feature plots
```

```
theme1 <- transparentTheme(trans = .4)
```

```
trellis.par.set(theme1)
```

```
featurePlot(x = weekly[, 1:7],
```

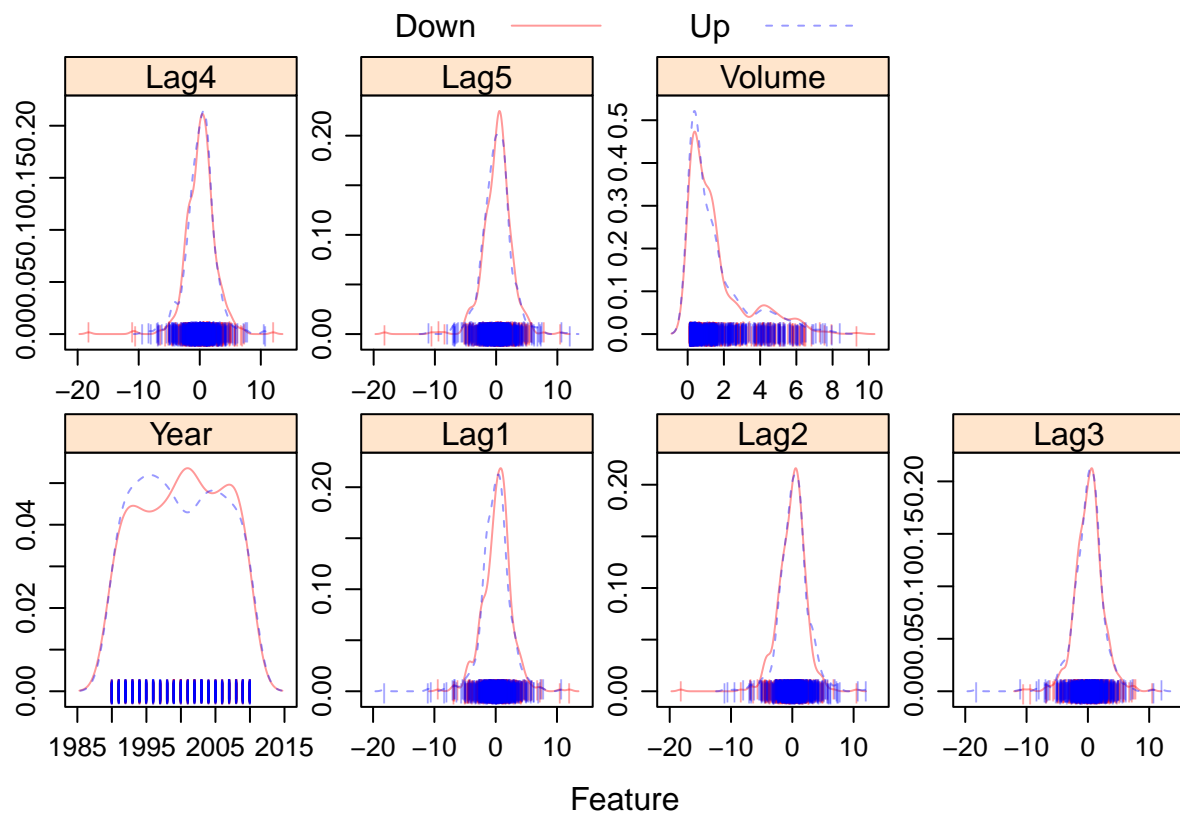
```
            y = weekly$Direction,
```

```
            scales = list(x = list(relation = "free"),
```

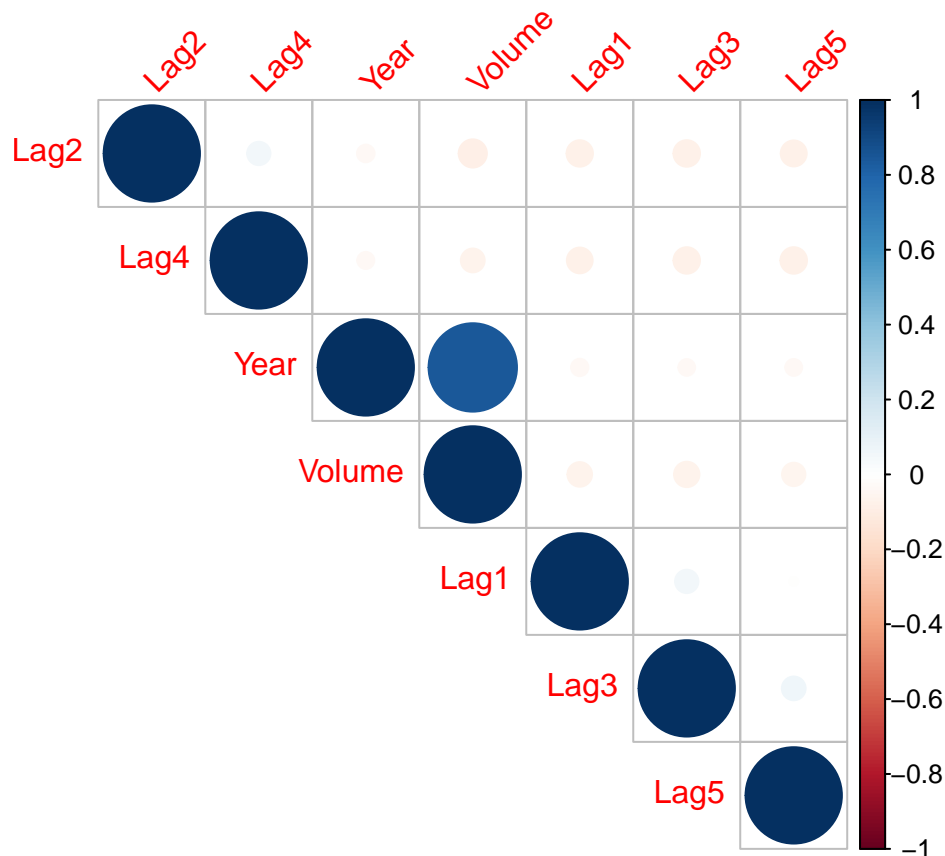
```
                           y = list(relation = "free")),
```

```
            plot = "density", pch = "|",
```

```
            auto.key = list(columns = 2))
```



```
# Correlation plot
corrplot(cor(weekly[, -8]), tl.srt = 45, order = 'hclust', type = 'upper')
```



The feature plots show that the distribution of the response classes (Up or Down) highly overlap for each feature. The correlation plot shows that there is not much pairwise correlation between the features.

(b)

```
#training set
train_df =
  weekly %>%
  filter(Year < 2009) %>%
  dplyr::select(-Year)

#test set
test_df = anti_join(weekly, train_df)
```

```
## Joining, by = c("Lag1", "Lag2", "Lag3", "Lag4", "Lag5", "Volume", "Direction")
```

```
contrasts(weekly$Direction)
```

```
##      Up
## Down  0
## Up    1
```

```

#perform logistic regression
glm_train <- glm(Direction ~ .,
                 data = train_df,
                 family = binomial(link = "logit"))
summary(glm_train)

##
## Call:
## glm(formula = Direction ~ ., family = binomial(link = "logit"),
##      data = train_df)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.7186  -1.2498   0.9823   1.0841   1.4911
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  0.33258    0.09421   3.530 0.000415 ***
## Lag1        -0.06231    0.02935  -2.123 0.033762 *
## Lag2         0.04468    0.02982   1.499 0.134002
## Lag3        -0.01546    0.02948  -0.524 0.599933
## Lag4        -0.03111    0.02924  -1.064 0.287241
## Lag5        -0.03775    0.02924  -1.291 0.196774
## Volume      -0.08972    0.05410  -1.658 0.097240 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 1354.7  on 984  degrees of freedom
## Residual deviance: 1342.3  on 978  degrees of freedom
## AIC: 1356.3
##
## Number of Fisher Scoring iterations: 4

```

It seems that the Lag1 predictor is statistically significant.

```

#compute the confusion matrix and overall fraction of correct predictions using the test data
test.pred.prob <- predict(glm_train, newdata = test_df,
                          type = "response")

test.pred <- rep("Down", length(test.pred.prob))
test.pred[test.pred.prob>0.5] <- "Up"

confusionMatrix(data = as.factor(test.pred),
                 reference = test_df$Direction,
                 positive = "Up")

## Confusion Matrix and Statistics
##
##              Reference
## Prediction Down Up

```

```
##      Down   31 44
##      Up     12 17
##
##              Accuracy : 0.4615
##              95% CI : (0.3633, 0.562)
##      No Information Rate : 0.5865
##      P-Value [Acc > NIR] : 0.9962
##
##              Kappa : -3e-04
##
##      McNemar's Test P-Value : 3.435e-05
##
##              Sensitivity : 0.2787
##              Specificity : 0.7209
##              Pos Pred Value : 0.5862
##              Neg Pred Value : 0.4133
##              Prevalence : 0.5865
##              Detection Rate : 0.1635
##      Detection Prevalence : 0.2788
##              Balanced Accuracy : 0.4998
##
##      'Positive' Class : Up
##
```

The confusion matrix indicates that the fraction of correct classification is 0.4615. Because the No Information Rate value (0.5865) is higher than the Accuracy value, it would be hard to conclude that the classification is entirely valid. The large P-value also indicates that the accuracy is not significantly better than the No Information Rate. In addition, the Kappa Value that is close to 0 indicates that the classification performance is not good. The sensitivity (0.2787) is low while the specificity (0.7209) is relatively high.

(c)

```
glm_train2 <- glm(Direction ~ Lag1 + Lag2,
                  data = train_df,
                  family = binomial(link = "logit"))

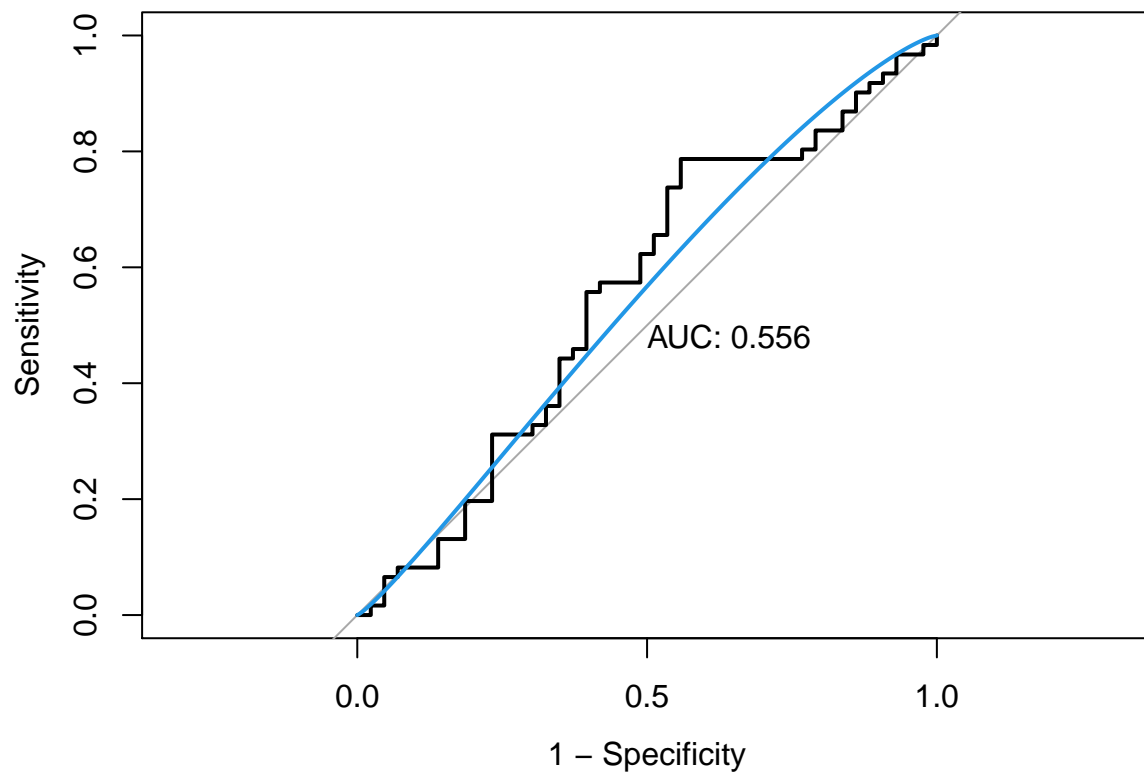
test.pred.prob <- predict(glm_train2, newdata = test_df,
                          type = "response")
test.pred <- rep("neg", length(test.pred.prob))
test.pred[test.pred.prob>0.5] <- "pos"

roc.glm <- roc(test_df$Direction, test.pred.prob)

## Setting levels: control = Down, case = Up

## Setting direction: controls < cases

plot(roc.glm, legacy.axes = TRUE, print.auc = TRUE)
plot(smooth(roc.glm), col = 4, add = TRUE)
```



AUC is 0.518.

The

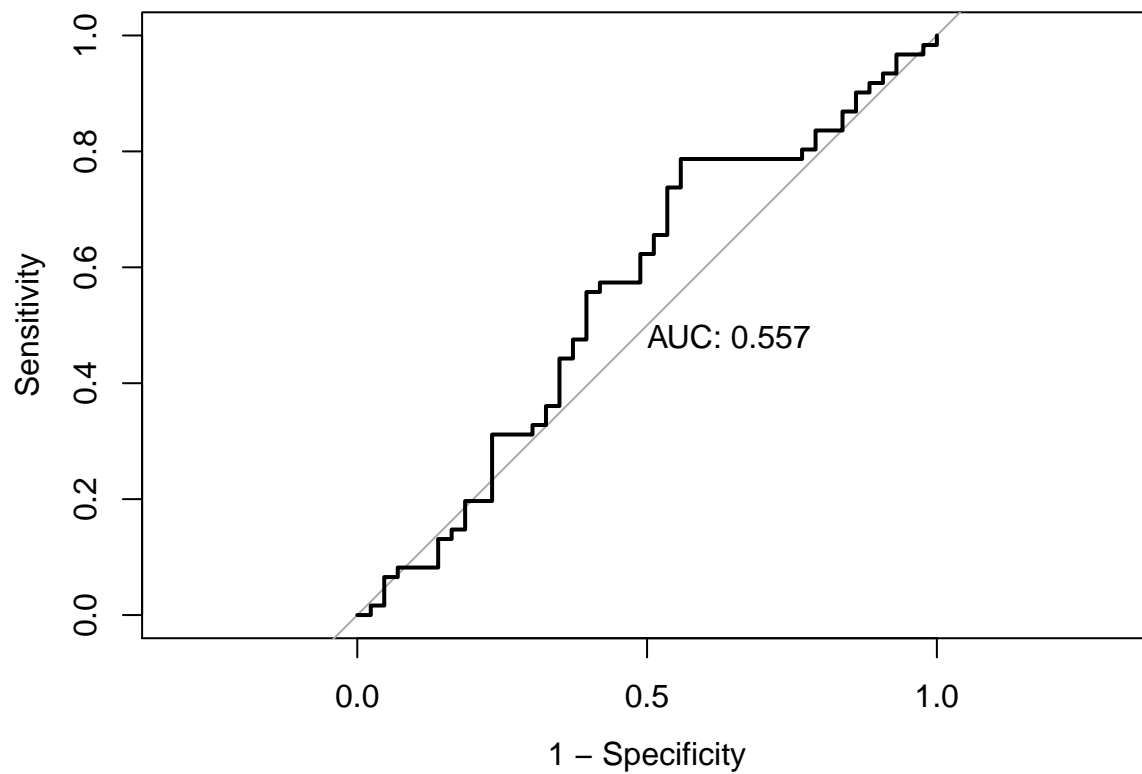
(d)

```
# LDA
lda.fit <- lda(Direction~ Lag1 + Lag2,
               data = train_df)
lda.pred <- predict(lda.fit, newdata = test_df)
roc.lda <- roc(test_df$Direction, lda.pred$posterior[,2])
```

```
## Setting levels: control = Down, case = Up
```

```
## Setting direction: controls < cases
```

```
plot(roc.lda, legacy.axes = TRUE, print.auc = TRUE)
```

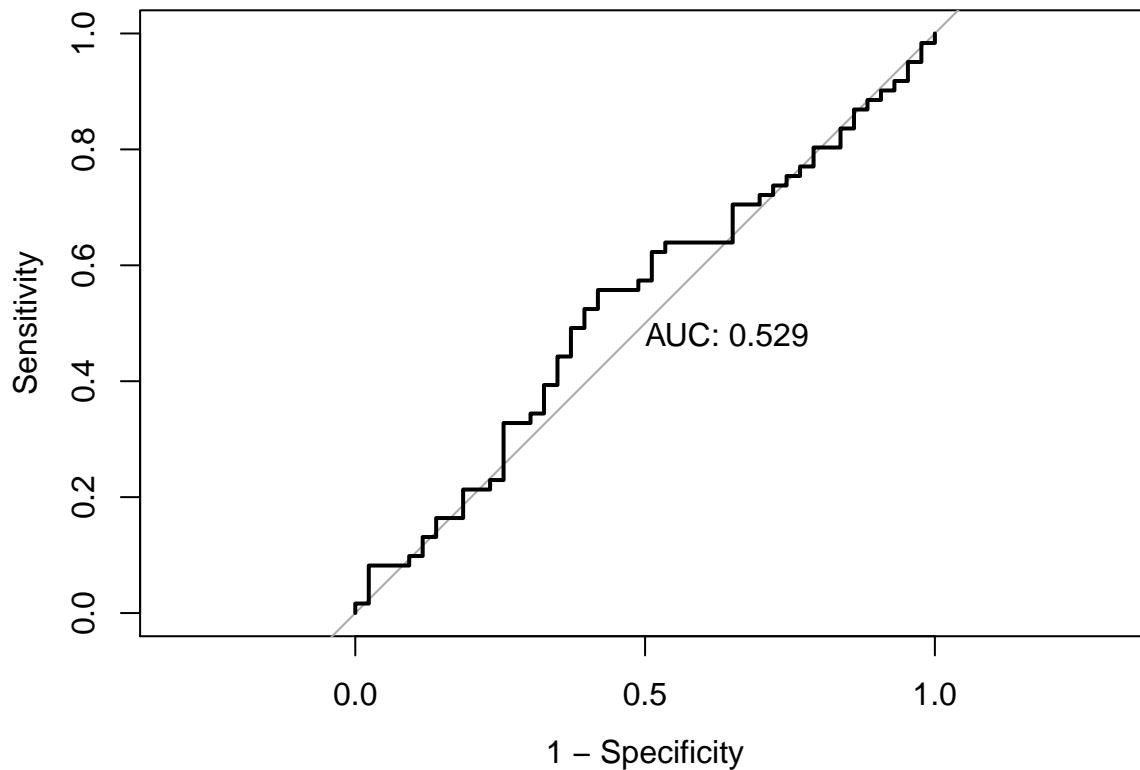


```
# QDA
qda.fit <- qda(Direction ~ Lag1 + Lag2,
               data = train_df)
qda.pred <- predict(qda.fit, newdata = test_df)
roc.qda <- roc(test_df$Direction, qda.pred$posterior[,2])
```

```
## Setting levels: control = Down, case = Up
```

```
## Setting direction: controls > cases
```

```
plot(roc.qda, legacy.axes = TRUE, print.auc = TRUE)
```

```
auc <- c(roc.lda$auc[1], roc.qda$auc[1])
```

The AUC from LDA is 0.55661 and the AUC from QDA is 0.5288.

(e)

```
ctrl <- trainControl(method = "repeatedcv",
                     repeats = 5,
                     summaryFunction = twoClassSummary,
                     classProbs = TRUE)

model.knn <- train(x = train_df[,1:2],
                  y = train_df$Direction,
                  method = "knn",
                  preProcess = c("center", "scale"),
                  tuneGrid = data.frame(k = seq(1, 499, by = 5)),
                  trControl = ctrl)
```

```
## Warning in train.default(x = train_df[, 1:2], y = train_df$Direction, method
## = "knn", : The metric "Accuracy" was not in the result set. ROC will be used
## instead.
```

```
model.knn$bestTune
```

```
##      k
## 100 496
```

```
pred_knn = predict.train(model.knn, newdata = test_df, type = 'prob')
roc_knn <- roc(test_df$Direction, pred_knn[,2])
```

```
## Setting levels: control = Down, case = Up
```

```
## Setting direction: controls < cases
```

```
plot.roc(roc_knn, legacy.axes = TRUE, print.auc = TRUE)
```

