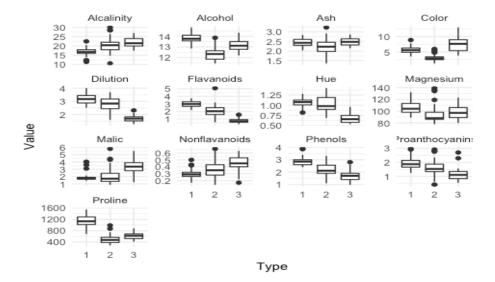
dehomework2

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Wine data-LDA, QDA and NB

```
# Data exploration: wine and other features
wine_train=read.csv("wine_train.csv")
wine_test=read.csv("wine_test.csv")
library(dplyr)
library(tidyr)
library(corrplot)
library(ggplot2)
head(wine_train)
    Type Alcohol Malic Ash Alcalinity Magnesium Phenols Flavanoids Nonflavanoids
## 1
       1 13.77 1.90 2.68
                                 17.1
                                                  3.00
                                                             2.79
                                           115
                                                                          0.39
## 2
           13.94 1.73 2.27
                                 17.4
                                           108
                                                  2.88
                                                             3.54
                                                                          0.32
       1
## 3
           13.75 1.73 2.41
                                 16.0
                                            89
                                                  2.60
                                                             2.76
                                                                          0.29
       1
## 4
          12.85 1.60 2.52
                                 17.8
                                            95
                                                  2.48
                                                             2.37
                                                                          0.26
       1
## 5
          13.63 1.81 2.70
                                 17.2
                                           112
                                                  2.85
                                                             2.91
                                                                          0.30
       1
                                 19.1
## 6
       1 13.58 1.66 2.36
                                           106
                                                  2.86
                                                             3.19
                                                                          0.22
## Proanthocyanins Color Hue Dilution Proline
## 1
              1.68 6.30 1.13 2.93
                                         1375
## 2
              2.08 8.90 1.12
                                  3.10
                                          1260
## 3
              1.81 5.60 1.15
                                  2.90
                                         1320
## 4
              1.46 3.93 1.09
                                  3.63
                                         1015
## 5
              1.46 7.30 1.28
                                  2.88
                                         1310
## 6
              1.95 6.90 1.09
                                  2.88
                                         1515
# Wide data
wine train$Type=factor(wine train$Type)
wine test$Type=factor(wine test$Type)
dat_exp=gather(wine_train, key="Variable", value="Value",-c("Type"))
ggplot(dat_exp) + geom_boxplot(aes(x = Type, y = Value)) + facet_wrap(.~Variable, scale
s = "free y") + theme_minimal()
```



Data Exploration

Based on the boxplots above, all features differ more or less across different wine types. Among them, Alchohol, Color, Dilution, Flavanoids, Alcalinity, Nonflavanoids, Phenois, Proline and Proanthocyanins are especially potential features that can help distinguish wine of different types.

LDA, QDA and Naive Bayes

```
# Lda modeL
wine_lda=lda(Type~.,data=wine_train)
wine_lda_train_predict=predict(wine_lda, wine_train)$class
wine_lda_test_predict=predict(wine_lda, wine_test)$class
train_error_lda=mean(wine_lda_train_predict!=wine_train$Type)
test_error_lda=mean(wine_lda_test_predict!=wine_test$Type)

# test error and training error for Lda
test_error_lda
## [1] 0.01818182
```

The testing error of LDA model is 0.018.

```
# qda model
wine_qda=qda(Type~.,data=wine_train)
wine_qda_train_predict=predict(wine_qda, wine_train)$class
wine_qda_test_predict=predict(wine_qda, wine_test)$class
train_error_qda=mean(wine_qda_train_predict!=wine_train$Type)
test_error_qda=mean(wine_qda_test_predict!=wine_test$Type)

# test error and training error for qda
test_error_qda
## [1] 0.03636364
```

The testing error for QDA model is 0.036.

```
library(e1071)
library(foreign)
```

```
library(boot)
library(class)
library(ISLR)

# Nb classifier
wine_nb=naiveBayes(Type~.,data=wine_train)
predict_nb_train=predict(wine_nb, wine_train)
predict_nb_test=predict(wine_nb, wine_test)
train_error_nb=mean(predict_nb_train!=wine_train$Type)
test_error_nb=mean(predict_nb_test!=wine_test$Type)

# error for nb classifier
train_error_nb
## [1] 0.01626016

test_error_nb
## [1] 0.03636364
```

The testing error for NbClassifier is 0.036.

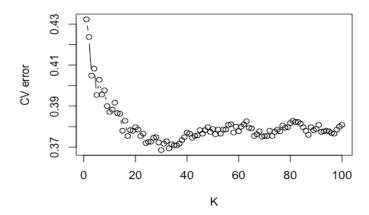
KNN and Cross Validation

Use cross-validation to select the best k and use the test data to evaluate the performance of the selected model. Show the training, cross-validation and test errors for each choice of k and report your findings.

```
theft train=read.csv("theft train.csv")
theft_test=read.csv("theft_test.csv")
theft_test$theft=factor(theft_test$theft)
theft_train$theft=factor(theft_train$theft)
# Leave one out CV function
Kfold_cv<-function(k_fold,knn,train,train_label){</pre>
  fold size=floor(nrow(train)/k fold)
  cv error=rep(0,k fold)
  for (i in 1:k_fold){
    if (i!=k_fold){
      id=((i-1)*fold size+1):(i*fold size)
    else {id=((i-1)*fold_size+1):nrow(train)
      }
    cv train=train[-id,]
    cv test=train[id,]
    mean cv train=colMeans(cv train)
    sd_cv_train=apply(cv_train,2,sd)
    cv train=scale(cv train, center=mean cv train, scale=sd cv train)
    cv test=scale(cv test, center=mean cv train, scale=sd cv train)
    pred_knn=knn(cv_train, cv_test, train_label[-id],k=knn)
    cv error[i]=mean(pred knn!=train label[id])
   return(mean(cv error))
```

```
set.seed(2020)
theft_train_X=theft_train[,-3]
theft_train_label=theft_train$theft
theft_test_X=theft_test[,-3]
theft_test_label=theft_test$theft
# 10 fold
k fold=10
knn=1:100
cv.errors=rep(1,100)
train.errors=rep(1,100)
test.errors=rep(1,100)
train_mean=colMeans(theft_train_X)
train_sd=apply(theft_train_X,2,sd)
theft_train_scaled=scale(theft_train_X,center=train_mean, scale=train_sd)
theft_test_scaled=scale(theft_test_X,center =train_mean, scale=train_sd)
set.seed(2020)
for (i in 1:100){
  cv.errors[i]=Kfold_cv(k fold, knn=knn[i], train=theft train X, train label = theft tr
ain_label)
}
set.seed(2020)
for (i in 1:100){
  knn_model1=knn(theft_train_scaled, theft_train_scaled, theft_train_label, knn[i])
  train.errors[i]=mean(knn_model1!=theft_train_label)
  knn_model2=knn(theft_train_scaled, theft_test_scaled, theft_train_label, knn[i])
  test.errors[i]=mean(knn_model2!=theft_test[,3])
}
set.seed(2020)
# which minimizes the cv error
min(cv.errors)
## [1] 0.3685714
which(cv.errors==min(cv.errors))
## [1] 30
# which minimized the test error
min(test.errors)
## [1] 0.368
which(test.errors==min(test.errors))
## [1] 26
plot(cv.errors~knn,type='b',main = '10-Fold CV error v.s. choice of k in KNN',xlab = 'K
',ylab = 'CV error')
```

10-Fold CV error v.s. choice of k in KNN

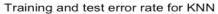


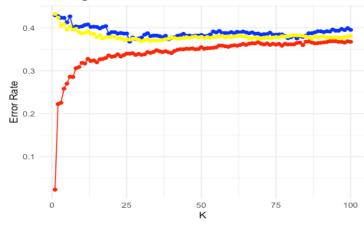
```
#test error at k=30
results=knn(theft_train_scaled, theft_test_scaled, theft_train_label, 30)
mean(results!=theft_test_label)
## [1] 0.378
```

The plot shows the CV error based on the 10-fold cross validation and lowest CV error (0.368) is obtained at k=30.

Fitting the best model selected by 10-fold cv on the test data, we get an error rate of 37.8%.

```
k=c(1:100)
errors2 = data.frame(train.errors, test.errors, cv.errors, k)
ggplot(errors2, aes(x = k)) +
   geom_line(aes(y = train.errors), col = "red") + geom_point(aes(y = train.errors), col
   = "red") +
   geom_line(aes(y = test.errors), col = "blue") + geom_point(aes(y = test.errors), col
   = "blue") +
   geom_line(aes(y = cv.errors), col = "yellow") + geom_point(aes(y = cv.errors), col =
"yellow") +
   ylab("Error Rate") + xlab("K") + ggtitle("Training and test error rate for KNN") + th
   eme_minimal()
```





The red line is the training error line, and it is always the lowest for whichever k we chose and this is expected because we are applying model to the in-sample data and thus the error rate is underestimated.

The test errors are mostly slightly higer than cv errors, but very close to each other. The k which minimizes cv error does not minimize test error. However, since we already used the cv estimation of error rate to select the model, it tends to overestimate the performance of the model. It would be therefore, use the test error (which is 0.378) at k=30 to report the performance.

CV for glm

```
weeklv=Weeklv
head(Weekly)
##
    Year Lag1
                Lag2
                       Lag3
                              Lag4
                                    Lag5
                                            Volume Today Direction
## 1 1990 0.816 1.572 -3.936 -0.229 -3.484 0.1549760 -0.270
Down
## 3 1990 -2.576 -0.270 0.816 1.572 -3.936 0.1598375 3.514
                                                               Up
## 4 1990 3.514 -2.576 -0.270 0.816 1.572 0.1616300 0.712
                                                               Up
## 5 1990 0.712 3.514 -2.576 -0.270 0.816 0.1537280 1.178
                                                               Up
## 6 1990 1.178 0.712 3.514 -2.576 -0.270 0.1544440 -1.372
                                                              Down
colnames(weekly)
## [1] "Year"
                 "Lag1"
                             "Lag2"
                                        "Lag3"
                                                   "Lag4"
                                                              "Lag5"
## [7] "Volume"
                 "Today"
                             "Direction"
weekly$Direction=as.character(weekly$Direction)
weekly$Direction[weekly$Direction=="Up"]="1"
weekly$Direction[weekly$Direction=="Down"]="0"
head(weekly)
                        Lag3
                              Lag4
                                            Volume Today Direction
    Year
           Lag1
                Lag2
                                     Lag5
## 1 1990 0.816 1.572 -3.936 -0.229 -3.484 0.1549760 -0.270
## 2 1990 -0.270 0.816 1.572 -3.936 -0.229 0.1485740 -2.576
                                                                 0
## 3 1990 -2.576 -0.270 0.816 1.572 -3.936 0.1598375 3.514
                                                                1
## 4 1990 3.514 -2.576 -0.270 0.816 1.572 0.1616300 0.712
                                                                1
## 5 1990 0.712 3.514 -2.576 -0.270 0.816 0.1537280 1.178
                                                                1
## 6 1990 1.178 0.712 3.514 -2.576 -0.270 0.1544440 -1.372
                                                                 0
weekly$Direction=factor(weekly$Direction)
```

Logit regression: Dir~lag1+lag2

```
glm.a=glm(Direction~Lag1+Lag2, family = binomial,data=weekly)
summary(glm.a)
##
## glm(formula = Direction ~ Lag1 + Lag2, family = binomial, data = weekly)
##
## Deviance Residuals:
    Min 10 Median
                              30
                                    Max
## -1.623 -1.261 1.001
                          1.083
                                  1.506
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.22122 0.06147 3.599 0.000319 ***
```

The result shows that the second lag has a significant positive effect on the direction of the stock return, while the first lag does not: The higer the percentage return for 2 weeks previous, the more likely the return on this week is positive.

The result also shows the model is probably not a good choice. The residual deviance can be regarded as a measure of goodness-of-nit, but it is highly significant (statistic=1488, df=1086), indicating that the model does not fit well.

Logit Regression excluding the first observation

```
weekly.1=weekly[-1,]
glm.b=glm(Direction~Lag1+Lag2, family = binomial,data=weekly.1)
summary(glm.b)
##
## Call:
### glm(formula = Direction ~ Lag1 + Lag2, family = binomial, data = weekly.1)
##
## Deviance Residuals:
      Min 1Q Median 3Q
##
                                        Max
## -1.6258 -1.2617 0.9999 1.0819 1.5071
##
## Coefficients:
   Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 0.22324 0.06150 3.630 0.000283 ***
## Lag1 -0.03843 0.02622 -1.466 0.142683
## Lag2
             0.06085 0.02656 2.291 0.021971 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 1494.6 on 1087 degrees of freedom
## Residual deviance: 1486.5 on 1085 degrees of freedom
## AIC: 1492.5
##
## Number of Fisher Scoring iterations: 4
```

The result roughly remains the same. The lag 2 is still significant, but with estimated coefficient increased from 0.06025 to 0.06085. The lag 1 still remains insignificant.

The null deviance and residual deviance all underwent a tiny decrease (by 2) and AIC as well. Generally speaking, nothing changed significantly in this model that excluded the first observation.

Predict the first observation

When the threhold is 0.5, the prediction on the first observation is wrong. (The prediction is "up" while the real label is "down")

```
set.seed(2020)
Loocv.errors=rep(0,nrow(weekly))
mse=rep(0,nrow(weekly))
for (i in (1:nrow(weekly))){
 train=weekly[-i,]
 test=weekly[i,]
  glm.cv=glm(Direction~Lag1+Lag2, data=train,family=binomial)
  predict=inv.logit(predict(glm.cv,test))
  mse[i]=(predict-as.numeric(as.character(test$Direction)))^2
  if (predict>0.5) {predict=1}
  else {predict=0}
  Loocv.errors[i]=(predict!=test$Direction)
mean(Loocv.errors)
## [1] 0.4499541
in_sample=inv.logit(predict(glm.a,weekly))
for (i in (1:length(in_sample))){
 if (in_sample[i]>0.5) {in_sample[i]=1}
  else {in sample[i]=0}
mean(in_sample!=weekly$Direction)
## [1] 0.4444444
```

The cv estimation of error rate is about 0.45 while the estimated error rate whithout cv (directly apply the model to the in-sample data and do the prediction) is 0.44. We see that 0.44 slightly overestimates the performance of the model and cv error rate should provide a more accurate evaluation of the model performance.

The error rate is pretty high, close to 0.5, indicating the model is not an optimal one and it needs improvement, such as adding more lags, polynomial terms etc.