3. ROC Curve

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1. Read the SPAM dataset

```
spam <- read.table("spambase/spambase.data",sep=",")</pre>
spam.names <- c(read.table("spambase/spambase.names",sep=":",skip=33,nrows=53,as.is=TRUE)[,1],</pre>
                 "char freq #",
                 read.table("spambase/spambase.names",sep=":",skip=87,nrows=3,as.is=TRUE)[,1],
                 "spam.01")
names(spam) <- spam.names</pre>
n<-dim(spam)[1]</pre>
p < -dim(spam)[2] - 1
spam.01 \leftarrow spam[,p+1]
spam.vars <- as.matrix(spam[,1:p])</pre>
cat(paste("n = ", n, ', p = ', p,sep=""))
## n = 4601, p = 57
cat(paste("Proportion of spam e-mails =", round(mean(spam.01), 2), sep=""))
## Proportion of spam e-mails =0.39
glm.spam <- glm(spam.01 ~ spam.vars,family=binomial)</pre>
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
# summary(qlm.spam)
```

2. Train-Test partioning of the data

The manual way to create a train-test split is shown below.

Note the composition of the data, which can be divided as follows:

4601 initial data split into train (2/3) -> 3068 and test (1/3) -> 1533 spam_train (2/3) -> 1209 and nospam_train (2/3) -> 1859 spam_test (1/3) -> 604 and nospam_test (1/3) -> 929

```
set.seed(1234)
spamIndex <- which(spam.01 %in% c(1))
nospamIndex <- which(spam.01 %in% c(0))
spamTrain <- sample(spamIndex, size = ceiling(2/3*length(spamIndex)))</pre>
```

```
nospamTrain <- sample(nospamIndex, size = ceiling(2/3*length(nospamIndex)))
trainIndex <- union(nospamTrain, spamTrain)

trainData <- spam[trainIndex,]
testData <- spam[-trainIndex,]

n <- dim(trainData)[1]
p <- dim(trainData)[2] - 1

Y <- scale(trainData[,p+1], center=FALSE, scale=FALSE)
X <- scale(trainData[,1:p], center=TRUE, scale=TRUE)</pre>
```

3. Classification rules

We now consider three methods to obtain classification rules: logistic regression via maximum likelihood, logistic regression via Lasso, and k-nearest neighbors.

Starting with logistic regression via maximum likelihood:

```
glm.logistic <- glm(spam.01 ~ ., data = as.data.frame(trainData), family = binomial())

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

predictions_glm <- predict(glm.logistic, newdata = testData[,1:p], type = "response")

#summary(glm.logistic)

Lasso GLM:
library(glmnet)

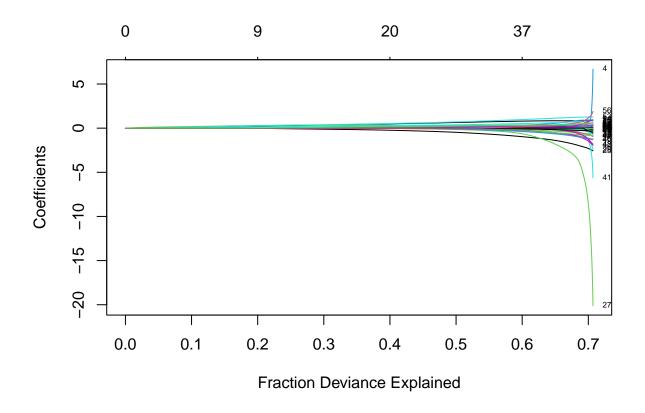
## Loading required package: Matrix

## Loaded glmnet 4.1-8

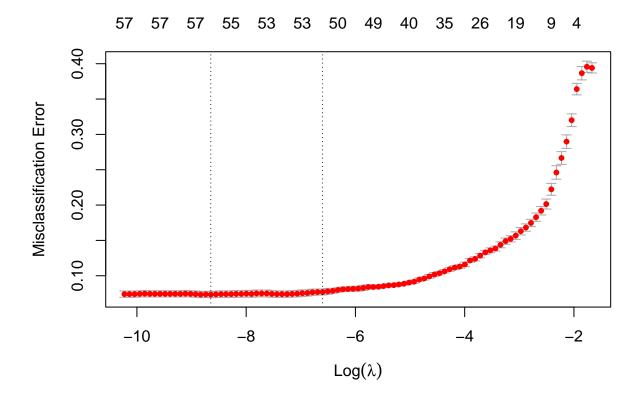
set.seed(1234)

fit_lasso <- glmnet(X, Y, family = "binomial")
  cvfit_lasso = cv.glmnet(X, Y, family = "binomial", type.measure = "class")

plot(fit_lasso, xvar = "dev", label = TRUE)</pre>
```



plot(cvfit_lasso)



```
final_fit_lasso <- glmnet(trainData[,1:p], trainData$spam.01, family = "binomial", lambda = cvfit_lasso
newx <- model.matrix(testData$spam.01 ~ ., data=testData[,1:p])
predictions_lasso <- predict(final_fit_lasso, newx = newx[,-1], type = "response")</pre>
```

k-nearest neighbours:

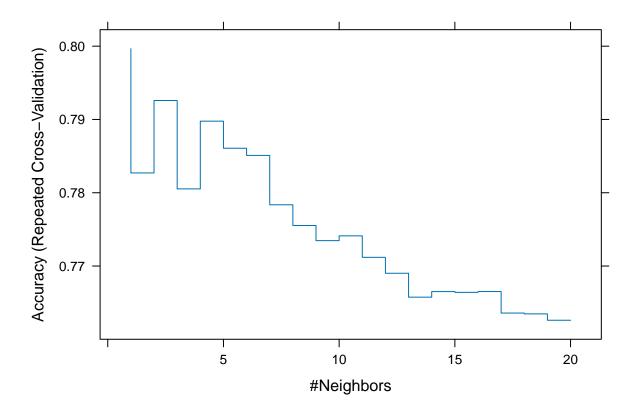
k ## 1 1

The best performing value for k was k = 1.

knn_fit

```
## k-Nearest Neighbors
##
## 3068 samples
## 57 predictor
```

```
2 classes: '0', '1'
##
##
## No pre-processing
## Resampling: Cross-Validated (10 fold, repeated 3 times)
## Summary of sample sizes: 2761, 2762, 2761, 2761, 2761, 2761, ...
## Resampling results across tuning parameters:
##
##
        Accuracy
                   Kappa
##
     1 0.7996562 0.5797007
##
     2 0.7827053 0.5446942
##
     3 0.7925855 0.5657786
##
     4 0.7805263 0.5399740
##
     5 0.7897664 0.5583766
##
     6 0.7860715 0.5501987
##
     7 0.7850936 0.5479650
##
     8 0.7783583 0.5336719
##
     9 0.7755314 0.5277080
##
    10 0.7734659 0.5233219
##
    11 0.7741191 0.5236551
     12 0.7711886 0.5180970
##
##
    13 0.7690142 0.5137125
##
    14 0.7657523 0.5067844
##
    15 0.7665148 0.5082876
##
    16 0.7664087 0.5087999
##
    17 0.7665176 0.5089476
##
    18 0.7635836 0.5025951
##
     19 0.7634732 0.5025083
##
     20 0.7626032 0.5010018
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was k = 1.
plot(knn_fit, print.thres = 0.5, type="S")
```



```
predictions_knn <- predict(knn_fit, newdata = testData)
testData$spam.01 <- as.factor(testData$spam.01)
confusionMatrix(predictions_knn, testData$spam.01)</pre>
```

```
## Confusion Matrix and Statistics
##
##
             Reference
  Prediction
                    1
##
##
            0 777 149
##
            1 152 455
##
                  Accuracy : 0.8037
##
                    95% CI: (0.7829, 0.8233)
##
##
       No Information Rate: 0.606
       P-Value [Acc > NIR] : <2e-16
##
##
                     Kappa: 0.5892
##
##
    Mcnemar's Test P-Value: 0.9082
##
##
##
               Sensitivity: 0.8364
               Specificity: 0.7533
##
##
            Pos Pred Value: 0.8391
            Neg Pred Value: 0.7496
##
##
                Prevalence: 0.6060
            Detection Rate: 0.5068
##
```

```
## Detection Prevalence : 0.6040
## Balanced Accuracy : 0.7948
##
## 'Positive' Class : 0
##
```

4. ROC Curves

Using the test data we find the following ROC curves:

```
library(pROC)

## Warning: package 'pROC' was built under R version 4.3.3

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

##

## Attaching package: 'pROC'

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

##

## cov, smooth, var

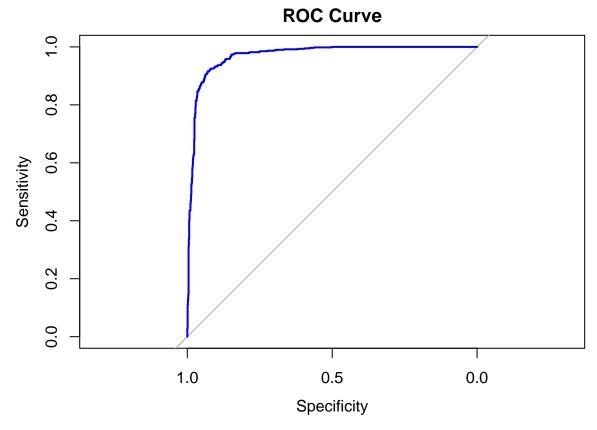
## glm ROC

roc_curve_glm <- roc(testData[,p+1], predictions_glm)

## Setting levels: control = 0, case = 1

## Setting direction: controls < cases

plot(roc_curve_glm, main = "ROC Curve", col = "blue", title="Classic GLM")</pre>
```



```
# Lasso ROC
roc_curve_lasso <- roc(testData[,p+1], predictions_lasso)

## Setting levels: control = 0, case = 1

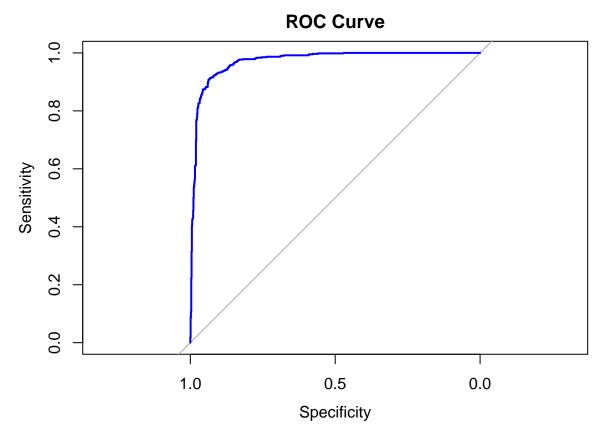
## Warning in roc.default(testData[, p + 1], predictions_lasso): Deprecated use a

## matrix as predictor. Unexpected results may be produced, please pass a numeric

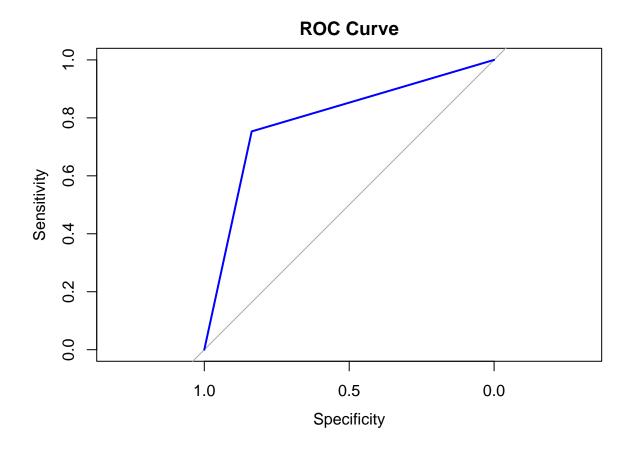
## vector.

## Setting direction: controls < cases

plot(roc_curve_lasso, main = "ROC Curve", col = "blue", title="Lasso GLM")</pre>
```



```
# knn ROC
roc_curve_knn <- roc(testData[,p+1], as.numeric(as.character(predictions_knn)))
## Setting levels: control = 0, case = 1
## Setting direction: controls < cases
plot(roc_curve_knn, main = "ROC Curve", col = "blue", title="Knn fit")</pre>
```



5. Misclassification Rate at $c=\frac{1}{2}$

```
predicted_classes_glm <- ifelse(predictions_glm >= 0.5, 1, 0)
misclassification_rate_glm <- mean(predicted_classes_glm != testData$spam.01)
cat(paste("missclassification rate for GLM using c = 1/2:",misclassification_rate_glm,sep=" "))
## missclassification rate for GLM using c = 1/2: 0.0802348336594912
predicted_classes_lasso <- ifelse(predictions_lasso >= 0.5, 1, 0)
misclassification_rate_lasso <- mean(predicted_classes_lasso != testData$spam.01)
cat(paste("missclassification rate for Lasso using c = 1/2:",misclassification_rate_lasso,sep=" "))
## missclassification rate for Lasso using c = 1/2: 0.076320939334638
predicted_classes_knn <- ifelse(as.numeric(as.character(predictions_knn)) >= 0.5, 1, 0)
misclassification_rate_knn <- mean(predicted_classes_knn != testData$spam.01)
cat(paste("missclassification rate for 1-nn using c = 1/2:",misclassification_rate_knn,sep=" "))
## missclassification rate for 1-nn using c = 1/2: 0.19634703196347
6. Calculating \ell_{val}
l_val_glm_store <- numeric(dim(testData)[1])</pre>
for (j in 1:dim(testData)[1]){
  aux1 = as.numeric(testData$spam.01[j]) * log(predictions_glm[j])
```

```
aux2 = (1-as.numeric(testData$spam.01[j])) * log(1-predictions_glm[j])
  l_val_glm_store[j] = aux1 + aux2
  if (is.nan(aux1) | is.nan(aux2) | l_val_glm_store[j] == -Inf ) {
    l_val_glm_store[j] = 0
  }
l_val_glm <- mean(l_val_glm_store)</pre>
###
l_val_lasso_store = numeric(dim(testData)[1])
for (j in 1:dim(testData)[1]){
  aux1 = as.numeric(testData$spam.01[j]) * log(predictions_lasso[j])
  aux2 = (1-as.numeric(testData$spam.01[j])) * log(1- predictions_lasso[j])
  1_val_lasso_store[j] = aux1 + aux2
  if (is.nan(aux1) | is.nan(aux2) | l_val_lasso_store[j] == -Inf | aux2 == Inf) {
    l_val_lasso_store[j] = 0
  }
}
l_val_lasso <- mean(l_val_lasso_store)</pre>
###
preds <- as.numeric(as.character(predictions_knn))</pre>
l_val_knn_store <- numeric(dim(testData)[1])</pre>
for (j in 1:dim(testData)[1]){
  aux1 = as.numeric(testData$spam.01[j]) * log(preds[j])
  aux2 = (1-as.numeric(testData$spam.01[j])) * log(1-preds[j])
  l_val_knn_store[j] = aux1 + aux2
  if (is.nan(aux1) | is.nan(aux2) | 1_val_knn_store[j] == -Inf | 1_val_knn_store[j] == Inf) {
    l_val_knn_store[j] = 0
}
l_val_knn <- mean(l_val_knn_store)</pre>
cat(paste("l_val for classical GLM",l_val_glm,sep=" "))
## 1_val for classical GLM -4.98083188638305
cat(paste("l_val for Lasso GLM",l_val_lasso,sep=" "))
## 1_val for Lasso GLM -5.65131876027708
cat(paste("l_val for 1-nn",l_val_knn,sep=" "))
## 1_val for 1-nn 0
```