HW4

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February 13, 2018

4 - 6

(a)

$$\begin{split} \text{P}\{\text{Y=receive an A}\} &= \frac{e^{\hat{\beta}_0 + \hat{\beta}_1 X_1 + \hat{\beta}_2 X_2}}{1 + e^{\hat{\beta}_0 + \hat{\beta}_1 X_1 + \hat{\beta}_2 X_2}} = \frac{e^{-6 + 0.05 X_1 + X_2}}{1 + e^{-6 + 0.05 X_1 + X_2}} \\ \text{P}\{\text{Y}|X_1 = 40, X_2 = 3.5\} &= \frac{e^{-6 + 0.05 \times 40 + 3.5}}{1 + e^{-6 + 0.05 \times 40 + 3.5}} \approx 0.37754 \end{split}$$

(b)

$$0.5 = \frac{e^{-6+0.05X_1+3.5}}{1+e^{-6+0.05X_1+3.5}}$$
$$X_1 = 50$$

4 - 13

```
library(MASS)
boston <- Boston
crim.median <- median(boston$crim)</pre>
boston$crimclass <- as.numeric(boston$crim > crim.median)
boston <- subset(boston, select = -crim)</pre>
splt <- floor(dim(boston)[1]*0.75)</pre>
train <- 1:splt
test <- (splt+1):dim(boston)[1]</pre>
boston.train <- boston[train,]</pre>
boston.test <- boston[test,]</pre>
crim.test <- boston$crimclass[test]</pre>
model.1 <- glm(crimclass ~ ., data = boston, family = binomial, subset = train)</pre>
summary(model.1)
##
## Call:
## glm(formula = crimclass ~ ., family = binomial, data = boston,
       subset = train)
##
## Deviance Residuals:
  Min 1Q Median
                                     3Q
                                             Max
## -2.3643 -0.2584 -0.0315 0.1406
                                          3.4545
##
## Coefficients:
                  Estimate Std. Error z value Pr(>|z|)
## (Intercept) -42.799118 7.890475 -5.424 5.82e-08 ***
```

```
## zn
               -0.066836
                           0.036975 -1.808 0.070668 .
## indus
                          0.051755 -1.710 0.087346 .
               -0.088478
## chas
               1.023592 0.753010 1.359 0.174041
## nox
               59.170886
                           9.555240 6.193 5.92e-10 ***
## rm
               -0.676176
                           0.816074 -0.829 0.407347
                          0.012974 0.667 0.504905
               0.008651
## age
                0.654216  0.232571  2.813  0.004909 **
## dis
## rad
                0.621347
                           0.183638 3.384 0.000716 ***
## tax
               -0.001433
                           0.003760 -0.381 0.703168
## ptratio
                0.485265
                           0.141215
                                      3.436 0.000590 ***
## black
               -0.009549
                           0.006112 -1.562 0.118195
                0.068709
                           0.054148
                                      1.269 0.204474
## lstat
## medv
                0.202732
                           0.080266 2.526 0.011546 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 500.3 on 378 degrees of freedom
## Residual deviance: 182.5 on 365 degrees of freedom
## AIC: 210.5
##
## Number of Fisher Scoring iterations: 8
prob.1 <- predict(model.1, boston.test, type = "response")</pre>
pred.1 <- rep(0, length(prob.1))</pre>
pred.1[prob.1 > 0.5] = 1
mean(pred.1 != crim.test)
## [1] 0.07874016
library(bestglm)
## Warning: package 'bestglm' was built under R version 3.4.3
## Loading required package: leaps
## Warning: package 'leaps' was built under R version 3.4.3
boston.for.bestglm <- within(boston, {</pre>
 y <- crimclass
  crimclass <- NULL
})
res.bestglm <- bestglm(Xy = boston.for.bestglm, family = binomial, IC = "AIC", method = "exhaustive")
## Morgan-Tatar search since family is non-gaussian.
summary(res.bestglm$BestModel)
##
## Call:
## glm(formula = y ~ ., family = family, data = Xi, weights = weights)
##
## Deviance Residuals:
                1Q Median
##
      Min
                                  3Q
                                          Max
## -2.4197 -0.1840 -0.0004 0.0022
                                       3.4087
##
## Coefficients:
```

```
Estimate Std. Error z value Pr(>|z|)
## (Intercept) -31.441272 6.048989 -5.198 2.02e-07 ***
               -0.082567
                          0.031424 -2.628 0.00860 **
               43.195824
                         6.452812
                                     6.694 2.17e-11 ***
## nox
## age
                0.022851
                          0.009894
                                     2.310 0.02091 *
                0.634380 0.207634
                                    3.055 0.00225 **
## dis
## rad
                0.718773
                         0.142066
                                     5.059 4.21e-07 ***
## tax
               -0.007676
                          0.002503 -3.066 0.00217 **
## ptratio
               0.303502
                          0.109255
                                     2.778 0.00547 **
## black
               -0.012866
                          0.006334
                                    -2.031 0.04224 *
## medv
                0.112882
                          0.034362
                                     3.285 0.00102 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 701.46 on 505 degrees of freedom
## Residual deviance: 216.22 on 496 degrees of freedom
## AIC: 236.22
##
## Number of Fisher Scoring iterations: 9
model.2 <- glm(crimclass ~ . - indus - chas - rm - lstat, data = boston, family = binomial,</pre>
              subset = train)
summary(model.2)
##
## Call:
## glm(formula = crimclass ~ . - indus - chas - rm - lstat, family = binomial,
##
      data = boston, subset = train)
##
## Deviance Residuals:
      Min
              1Q
                    Median
                                 3Q
                                         Max
## -2.3146 -0.3041 -0.0390 0.1653
                                      3.3830
##
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) -37.078705 6.924853 -5.354 8.58e-08 ***
                         0.032505 -2.126 0.03354 *
## zn
              -0.069092
## nox
              48.551533
                         7.461723
                                     6.507 7.68e-11 ***
## age
                0.011921
                          0.010410
                                     1.145 0.25215
## dis
                0.571056
                          0.211047
                                     2.706 0.00681 **
## rad
                0.651362 0.163957
                                     3.973 7.10e-05 ***
## tax
               -0.002903
                          0.003510 -0.827 0.40817
## ptratio
                0.377143
                           0.121314
                                     3.109 0.00188 **
               -0.009162
                          0.006100 -1.502 0.13314
## black
## medv
                0.114768
                          0.036731
                                     3.125 0.00178 **
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 500.30 on 378 degrees of freedom
## Residual deviance: 190.08 on 369 degrees of freedom
## AIC: 210.08
```

```
##
## Number of Fisher Scoring iterations: 8
prob.2 <- predict(model.2, boston.test, type = "response")
pred.2 <- rep(0, length(prob.2))
pred.2[prob.2 > 0.5] = 1
mean(pred.2 != crim.test)
```

```
## [1] 0.07874016
```

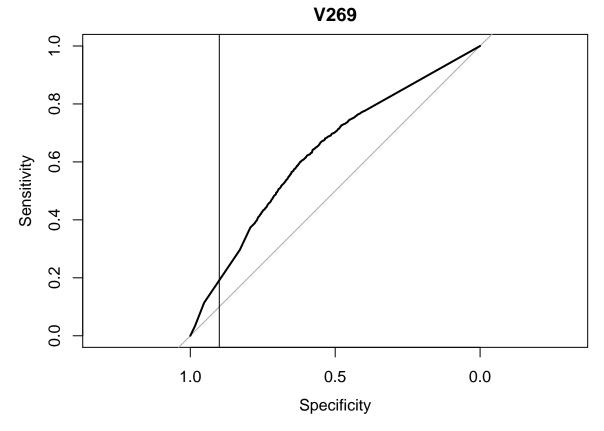
Create a new column called "crimclass". If crime rate is above the median, crimclass is 1, and 0 otherwise. 75% of dataset is training data and 25% is test dataset. The error rate when using all predictors is 7.874016%. Using "bestglm" package we find the best logistic regression model selected from all predictors. With 9 of the predictors, the new logistic regression model also have 7.874016% error rate.

MNIST

Problem 1

```
load("mnist all.RData")
library(pROC)
## Warning: package 'pROC' was built under R version 3.4.3
## Type 'citation("pROC")' for a citation.
##
## Attaching package: 'pROC'
## The following objects are masked from 'package:stats':
##
##
       cov, smooth, var
index <- (train$y == 0 \mid train<math>$y == 1)
df <- train$x[index,]</pre>
df.y <- train$y[index]</pre>
df <- as.data.frame(df)</pre>
df$y <- df.y
var(df[,269])
## [1] 12159.13
model.3 <- glm(y ~ V269, data = df, family = binomial)
summary(model.3)
##
## Call:
## glm(formula = y ~ V269, family = binomial, data = df)
##
## Deviance Residuals:
##
       Min
                  1Q
                       Median
                                     3Q
                                             Max
## -1.4761 -1.0599
                       0.9072
                                 1.0202
                                          1.3606
##
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.420932
                            0.027898 -15.09 <2e-16 ***
```

```
## V269
                0.004315
                           0.000167
                                      25.83 <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 17504 on 12664 degrees of freedom
## Residual deviance: 16814 on 12663 degrees of freedom
## AIC: 16818
##
## Number of Fisher Scoring iterations: 4
df$pred <- predict(model.3, type = "response")</pre>
myroc <- roc(df$y, df$pred)</pre>
plot(myroc, main = "V269")
abline(v = 0.9)
```



```
x = c(mytable[2,2]/sum(mytable[2,]), mytable[1,2]/sum(mytable[1,]))
names(x) <- c("trueP", "falseP")
x</pre>
```

trueP falseP ## 0.11391279 0.04761101

Variable is Pixel No. 269

Logistic regression equation:

$$P\{y\} = \frac{e^{\hat{\beta}_0 + \hat{\beta}_1 V_{269}}}{1 + e^{\hat{\beta}_0 + \hat{\beta}_1 V_{269}}} = \frac{e^{-0.420932 + 0.004315 V_{269}}}{1 + e^{-0.420932 + 0.004315 V_{269}}}$$

When the fraction of false positives is 0.1,

$$\frac{0.1 - 0.04761101}{0.1708594 - 0.04761101} \times (0.2963512 - 0.11391279) + 0.11391279 = 0.1914616$$

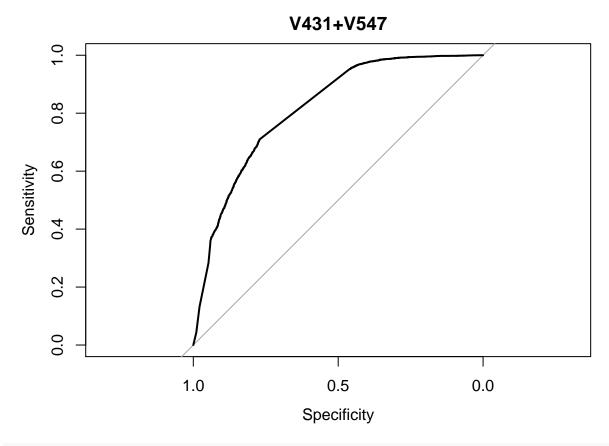
the fraction of true positives is approximately 0.19

Problem 2

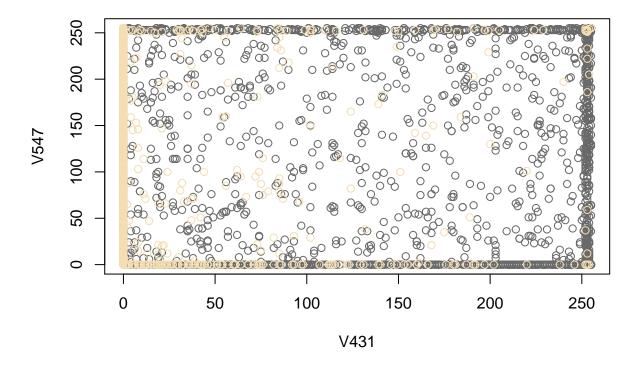
```
var(df[,431])
## [1] 7030.701
var(df[,547])
## [1] 12423.97
cor(df[,431], df[,547])
## [1] -0.02477022
model.4 <- glm(y ~ V431 + V547, data = df, family = binomial)
summary(model.4)
##
## Call:
## glm(formula = y ~ V431 + V547, family = binomial, data = df)
##
## Deviance Residuals:
##
                1Q
                     Median
                                  3Q
                                          Max
                     0.6079
                                       3.3224
## -1.8892 -1.2011
                              0.8417
##
## Coefficients:
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.0555008 0.0284174
                                      1.953
                                             0.0508 .
             -0.0219321 0.0006467 -33.914
                                              <2e-16 ***
## V431
              0.0060600 0.0002017 30.045
## V547
                                             <2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
```

```
## Null deviance: 17504 on 12664 degrees of freedom
## Residual deviance: 12956 on 12662 degrees of freedom
## AIC: 12962
##
## Number of Fisher Scoring iterations: 6

df$pred <- predict(model.4, type = "response")
myroc <- roc(df$y, df$pred)
plot(myroc, main = "V431+V547")</pre>
```



```
auc(df$y, df$pred)
## Area under the curve: 0.8166
plot(df$V431[df$y == 0], df$V547[df$y == 0], col = "dimgrey", xlab = "V431", ylab = "V547")
points(df$V431[df$y == 1], df$V547[df$y == 1], col = "wheat")
```



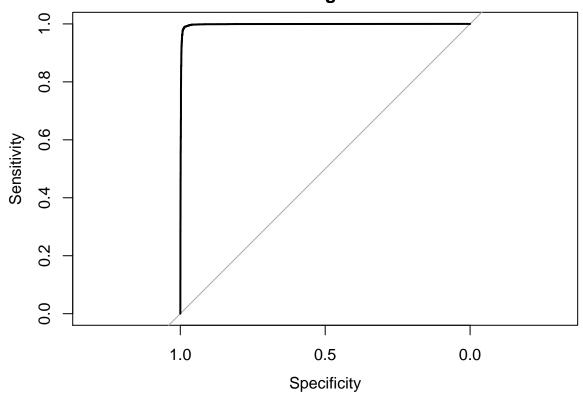
Classifier ssing Pixel No.431 and No.547 is good. From the scatterplot, major points at bottomleft are 1s and those at upperright are 0s. The training accuracy is 81.66%.

Problem 3

```
variance <- c()</pre>
index <- 1:784
for (i in index){
  variance <- c(variance, var(df[,i]))</pre>
df.var <- data.frame(index, variance)</pre>
df.var <- df.var[order(-variance),]</pre>
head(df.var, 10)
##
       index variance
## 407
          407 15407.78
## 435
          435 15163.62
## 379
          379 14902.05
## 463
          463 14224.23
## 462
          462 14008.45
   352
          352 13953.82
##
   351
          351 13670.39
##
   380
          380 13664.89
   490
          490 13651.83
##
## 434
          434 13541.13
```

```
model.5 \leftarrow glm(y \sim V351 + V352 + V379 + V380 + V407 + V434 + V435 + V462 + V463 + V490, data = df,
              family = binomial)
summary(model.5)
##
## Call:
## glm(formula = y \sim V351 + V352 + V379 + V380 + V407 + V434 + V435 +
      V462 + V463 + V490, family = binomial, data = df)
##
## Deviance Residuals:
##
      Min
               10
                   Median
                                3Q
                                        Max
## -3.9196 -0.0958 0.0320
                           0.0535
                                     3.2821
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -5.381427
                         0.173351 -31.043 < 2e-16 ***
## V351
              0.002624
                         0.001865
                                  1.407 0.159352
## V352
              0.001963
                        0.001781
                                  1.102 0.270501
## V379
              ## V380
              0.008275
                         0.001852 4.468 7.90e-06 ***
                         0.002554 2.083 0.037295 *
## V407
              0.005319
## V434
              0.007240
                        0.001893 3.823 0.000132 ***
## V435
              0.006364 0.002467 2.579 0.009904 **
## V462
              0.006390 0.002615 2.444 0.014531 *
              ## V463
## V490
             -0.001590 0.001871 -0.850 0.395532
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 17504.4 on 12664 degrees of freedom
## Residual deviance: 1130.5 on 12654 degrees of freedom
## AIC: 1152.5
##
## Number of Fisher Scoring iterations: 8
df$pred <- predict(model.5, type = "response")</pre>
myroc <- roc(df$y, df$pred)</pre>
plot(myroc, main = "10 with the Largest Variances")
```

10 with the Largest Variances



auc(myroc)

Area under the curve: 0.9977

The new ROC curve performs better than previous one. The training accuracy is 99.77%.