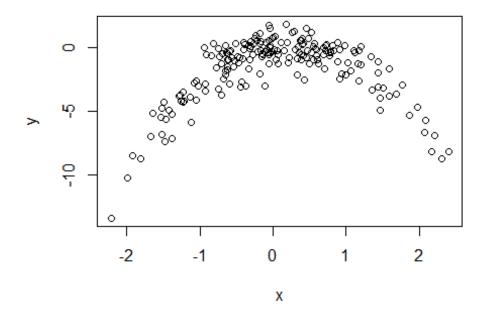
## **Abhishek HW4**

```
# This question should be answered using the Default data set. In Chapter 4
on classification, we used logistic regression to predict the probability of
default using income and balance. Now we will estimate the test error of this
logistic regression model using the validation set approach. Do not forget to
set a random seed before beginning your analysis.
# (a) Fit a logistic regression model that predicts default using income and
balance.
library(ISLR)
attach(Default)
head(Default)
##
     default student
                       balance
                                  income
## 1
                 No 729.5265 44361.625
         No
## 2
          No
                Yes 817.1804 12106.135
## 3
                 No 1073.5492 31767.139
          No
## 4
         No
                 No 529.2506 35704.494
## 5
                 No 785.6559 38463.496
         No
## 6
         No
                Yes 919.5885 7491.559
set.seed(1)
log_fit <- glm(default ~ income + balance, data = Default, family =</pre>
"binomial")
summary(log fit)
##
## Call:
## glm(formula = default ~ income + balance, family = "binomial",
##
      data = Default)
##
## Deviance Residuals:
##
      Min
                 10
                     Median
                                   3Q
                                           Max
## -2.4725 -0.1444 -0.0574 -0.0211
                                        3.7245
##
## Coefficients:
                 Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -1.154e+01 4.348e-01 -26.545 < 2e-16 ***
               2.081e-05 4.985e-06 4.174 2.99e-05 ***
## income
               5.647e-03 2.274e-04 24.836 < 2e-16 ***
## balance
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 2920.6 on 9999 degrees of freedom
```

```
## Residual deviance: 1579.0 on 9997 degrees of freedom
## AIC: 1585
## Number of Fisher Scoring iterations: 8
# (b) Using the validation set approach, estimate the test error of this
model. You need to perform the following steps:
# i. Split the sample set into a training set and a validation set.
indices <- sample(nrow(Default), nrow(Default)*0.5)</pre>
train <- Default[indices,]</pre>
test <- Default[-indices,]</pre>
# ii. Fit a logistic regression model using only the training data set.
log fit <- glm(default ~ income + balance,family = binomial,data=train)</pre>
# iii. Obtain a prediction of default status for each individual in the
validation set using a threshold of 0.5.
log proba <- predict(log fit, test, type="response")</pre>
log predict <- ifelse(log proba > 0.5, "Yes", "No")
# iv. Compute the validation set error, which is the fraction of the
observations in the validation set that are misclassified.
table(test$default, log predict, dnn=c("Actual", "Predicted"))
##
         Predicted
## Actual
            No Yes
                 28
##
      No 4805
                 52
##
      Yes 115
mean(log_predict != test$default)
## [1] 0.0286
# (c) Repeat the process in (b) three times, using three different splits of
the observations into a training set and a validation set. Comment on the
results obtained.
indices <- sample(nrow(Default), nrow(Default)*0.9)</pre>
train <- Default[indices,]</pre>
test <- Default[-indices,]</pre>
log_fit <- glm(default ~ income + balance,family = binomial,data=train)</pre>
log_proba <- predict(log_fit, test, type="response")</pre>
log_predict <- ifelse(log_proba > 0.5, "Yes", "No")
table(test$default, log_predict, dnn=c("Actual", "Predicted"))
```

```
Predicted
## Actual No Yes
                9
##
      No 958
##
      Yes 22 11
mean(log predict != test$default)
## [1] 0.031
indices <- sample(nrow(Default), nrow(Default)*0.7)</pre>
train <- Default[indices,]</pre>
test <- Default[-indices,]</pre>
log_fit <- glm(default ~ income + balance,family = binomial,data=train)</pre>
log_proba <- predict(log_fit, test, type="response")</pre>
log predict <- ifelse(log proba > 0.5, "Yes", "No")
table(test$default, log_predict, dnn=c("Actual", "Predicted"))
         Predicted
##
## Actual
            No Yes
##
      No 2889
                  16
##
            69
                  26
      Yes
mean(log predict != test$default)
## [1] 0.02833333
indices <- sample(nrow(Default), nrow(Default)*0.4)</pre>
train <- Default[indices,]</pre>
test <- Default[-indices,]</pre>
log_fit <- glm(default ~ income + balance,family = binomial,data=train)</pre>
log_proba <- predict(log_fit, test, type="response")</pre>
log_predict <- ifelse(log_proba > 0.5, "Yes", "No")
table(test$default, log_predict, dnn=c("Actual", "Predicted"))
         Predicted
##
## Actual
            No Yes
      No 5775
##
                  16
##
      Yes 142
                  67
mean(log_predict != test$default)
## [1] 0.02633333
# We see that the test error rate's are variable.
# (d) Consider another Logistic regression model that predicts default using
income, balance and student (qualitative). Estimate the test error for this
model using the validation set approach. Does including the qualitative
variable student lead to a reduction of test error rate?
indices <- sample(nrow(Default), nrow(Default)*0.5)</pre>
train <- Default[indices,]</pre>
```

```
test <- Default[-indices,]</pre>
log_fit <- glm(default ~ income + balance + student, family = binomial,</pre>
data=train)
log_proba <- predict(log_fit, test, type="response")</pre>
log_predict <- ifelse(log_proba > 0.5, "Yes", "No")
table(test$default, log_predict, dnn=c("Actual", "Predicted"))
##
         Predicted
## Actual
            No Yes
##
      No 4829
                  22
      Yes 101
                  48
##
mean(log_predict != test$default)
## [1] 0.0246
# The addition of the "student" dummy variable doesn't lead to a reduction in
the test error rate.
# This question requires performing cross validation on a simulated data set.
# (a) Generate a simulated data set as follows:
       set.seed(1)
#
       x=rnorm(200)
       y=x-2*x^2+rnorm(200)
# In this data set, what is 🛽 and what is 🗗? Write out the model used to
generate the data in equation form (i.e., the true model of the data).
set.seed(1)
x = rnorm(200)
y = x-2*x^2+rnorm(200)
# n=200 and p=2, the model used is Y = X - 2X2 + \varepsilon rror.
# (b) Create a scatter plot of \mathbb{Z} vs \mathbb{Z}. Comment on what you find.
plot(x, y)
```



```
# There is a curved(non-linear) relationship.
# (c) Consider the following four models for the data set:
          i. 2 = 20 + 212 + 2
         ii. 2 = 20 + 212 + 222 + 2
#
        iii. 2 = 20 + 212 + 222 + 2323 + 2
#
#
         iv. 2 = 20 + 212 + 2222 + 2323 + 2424 + 2
  Compute the LOOCV errors that result from fitting these models.
library(boot)
set.seed(1)
data <- data.frame(x, y)</pre>
cv_error = rep(0,4)
for (i in 1:4){
glm_fit = glm(y\sim poly(x,i), data=data)
cv_error[i] = cv.glm(data, glm_fit)$delta[1]
}
cv_error
## [1] 6.037638 1.040922 1.039049 1.028604
# (d) Repeat (c) using another random seed, and report your results. Are your
results the same as what you got in (c)? Why?
library(boot)
set.seed(20)
data <- data.frame(x, y)</pre>
```

```
cv error = rep(0,4)
for (i in 1:4){
glm_fit = glm(y\sim poly(x,i), data=data)
cv_error[i] = cv.glm(data, glm_fit)$delta[1]
cv_error
## [1] 6.037638 1.040922 1.039049 1.028604
# The results above are identical to the results obtained in (c). While
performing LOOCV multiple times will always yield the same results, because
we split based on one observation each time.
# (e) Which of the models in (c) has the smallest LOOCV error? Is this what
you expected? Explain your answer.
# The LOOCV error is minimum for the 2nd order polynomial function. This is
expected since we saw in (b) that the relation between "x" and "y" is
quadratic.
# (f) Now we use 5-fold CV for the model selection. Compute the CV errors
that result from fitting the four models. Which model has the smallest CV
error? Are the results consistent with LOOCV?
library(boot)
set.seed(1)
kcv error = rep(0,4)
data <- data.frame(x, y)</pre>
for (i in 1:4) {
glm_fit = glm(y\sim poly(x,i), data=data)
kcv_error[i] = cv.glm(data, glm_fit, K=5)$delta[1]
kcv_error
## [1] 5.888437 1.036580 1.039166 1.015776
# The 5-fold cross validation error is minimum for the 2nd order polynomial
function. Also, the results are in consistent with LOOCV.
# (q) Repeat (f) using 10-fold CV. Are the results the same as 5-fold CV?
library(boot)
set.seed(1)
kcv_error = rep(0,4)
data <- data.frame(x, y)</pre>
for (i in 1:4) {
glm_fit = glm(y\sim poly(x,i), data=data)
kcv error[i] = cv.glm(data, glm fit, K=10)$delta[1]
kcv_error
```

```
## [1] 5.968520 1.043498 1.035259 1.018977
```

# There is not much of a difference compared to error obtained in (f).