# Stat 427/627 Statistical Machine Learning

#### In-class Lab 5: Cross-Validation

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Recall the Auto data set in the ISLR2 package. This data frame has 392 observations on the following 9 variables.

- mpg: miles per gallon
- cylinders: Number of cylinders between 4 and 8
- displacement: Engine displacement (cu. inches)
- horsepower: Engine horsepower
- weight: Vehicle weight (lbs.)
- acceleration: Time to accelerate from 0 to 60 mph (sec.)
- year: Model year (modulo 100)
- origin: Origin of car (1. American, 2. European, 3. Japanese)
- name: Vehicle name

```
library(ISLR2)
colnames(Auto)
```

```
## [1] "mpg" "cylinders" "displacement" "horsepower" "weight"
## [6] "acceleration" "year" "origin" "name"
```

## 1 Validation set approach (review)

Randomly split the data set into training vs validation (or testing) set.

```
pct.train <- 0.5  # training percentage can vary, but should be > 0.5
n <- nrow(Auto)
Z <- sample(n, floor(pct.train*n))
auto.train <- Auto[Z, ]
auto.test <- Auto[-Z, ]

mpg.lm <- lm(mpg ~ weight + horsepower + acceleration, data=auto.train)

# Predict Y on the testing/validation set.
pred.test <- predict(mpg.lm, newdata=auto.test)</pre>
```

```
# Prediction MSE
mean((auto.test$mpg - pred.test)^2)
```

## [1] 18.11043

#### 2 Leave-One-Out Cross-Validation (LOOCV) for linear regression.

There are functions in different R package that can do LOOCV for various algorithms. For examples, lda() and qda() have a built-in option CV=TRUE that returns LOOCV classification. Function knn.cv() handles cross-validation for KNN.

We will work with cv.glm(data, glmfit, cost, K) in package boot for linear regression and logistic regression (later).

```
# install.package(boot)
library(boot)
```

Be sure to the entire date set for LOOCV. The algorithm will take care of the "split."

• glm() can also fit linear regression with family=gaussian (default).

```
mpg.glm <- glm(mpg ~ weight + horsepower + acceleration, family=gaussian, data=Auto)
mpg.glm
##
## Call: glm(formula = mpg ~ weight + horsepower + acceleration, family = gaussian,
##
       data = Auto)
##
## Coefficients:
##
   (Intercept)
                       weight
                                  horsepower acceleration
                    -0.005789
##
      45.678293
                                   -0.047496
                                                 -0.002066
## Degrees of Freedom: 391 Total (i.e. Null); 388 Residual
## Null Deviance:
## Residual Deviance: 6994 AIC: 2252
cv.error <- cv.glm(data=Auto, glmfit=mpg.glm)</pre>
names(cv.error)
## [1] "call"
                        "delta" "seed"
cv.error$delta
```

```
## [1] 18.25595 18.25542
```

cv.glm() saves the LOOCV prediction MSE  $(\frac{1}{n}\sum (Y_i-\widehat{Y}_{(-i)})^2)$  in delta.delta consists of 2 numbers:

- delta[1] is the estimated prediction error. - delta[2] is adjusted for the lost sample size due to cross-validation.

For model comparison purpose, one can use either version of delta as long as it is consistent in the analysis.

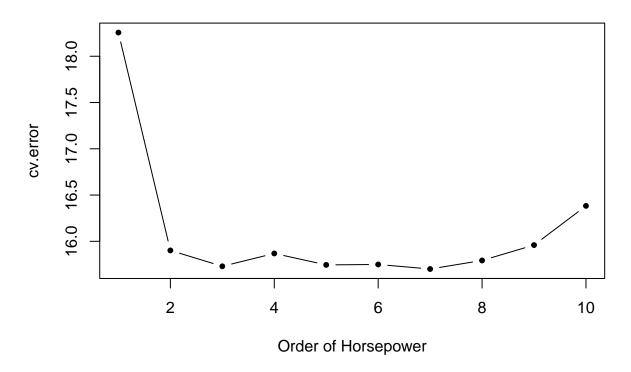
Consider models with different orders of horsepower as the predictor.

```
cv.error <- rep(0, 10)
# Use a loop to fit model with `horsepower` in 1, 2, ..., p orders.
for (p in 1:10) {
   glm.fit <- glm( mpg ~ weight + poly(horsepower,p) + acceleration, data=Auto)</pre>
```

```
cv.error[p] = cv.glm( Auto, glm.fit )$delta[1]
}
cv.error

## [1] 18.25595 15.90163 15.72995 15.86879 15.74517 15.74989 15.70073 15.79314
## [9] 15.95933 16.38301
plot(cv.error, type="b", pch=20, xlab="Order of Horsepower", main="LOOCV")
```

#### **LOOCV**

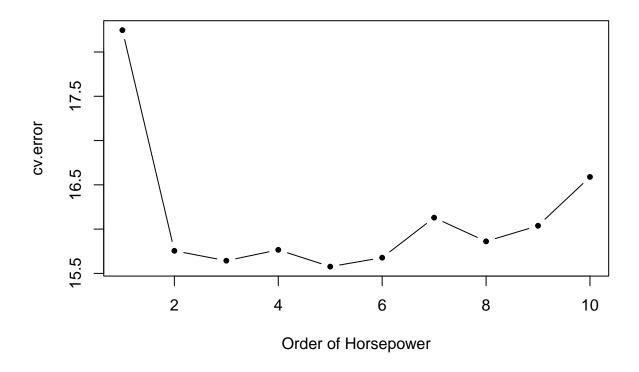


### 3 K-fold Cross-Validation for linear regression.

```
We can set the number of folds in cv.glm() using K=__ in the function. By default, K = n which is LOOCV.
cv.error <- rep(0, 10)
# Use a loop to fit model with `horsepower` in 1, 2, ..., p orders.
for (p in 1:10) {
    glm.fit <- glm( mpg ~ weight + poly(horsepower,p) + acceleration, data=Auto)
    cv.error[p] = cv.glm(Auto, glm.fit, K=10)$delta[1]
}
cv.error
## [1] 18.24636 15.75527 15.64321 15.76580 15.57649 15.67733 16.12964 15.86144</pre>
```

```
## [1] 16.24030 13.75327 13.04321 13.70300 13.37043 13.0733 10.12304 13.00144 ## [9] 16.03796 16.58903 plot(cv.error, type="b", pch=20, xlab="Order of Horsepower", main="10-fold CV")
```

#### 10-fold CV



Note that K-fold CV values change when you rerun the code, unless the random seed is set. Why?

#### 4 Cross-validation for logistic regression.

Recall that cv.glm() computes delta to estimate the prediction MSE  $(\frac{1}{n}\sum (Y_i - \widehat{Y}_{(-i)})^2)$ . When the response is 0-1,  $\widehat{Y}_{(-i)} = \widehat{\pi}_{(-i)}$ , where  $\pi = P(Y = 1)$ .

However, the correct classification rate and the error rate are more standard measures of classification accuracy. We can force <code>cv.glm()</code> to return these measures by introducing a suitable loss function. The prediction error rate will be calcuated and saved in <code>delta</code>.

Define a 0-1 loss function for classification. We will use cutoff value 0.5 for now. You can change it if needed.

```
loss <- function(Y, pred.p){
  return( mean( (Y==1 & pred.p < 0.5) | (Y==0 & pred.p >= 0.5) ) )
}
loss(c(1, 1), c(0.3, 0.6))
```

## [1] 0.5

Recall the Students' depression data.

```
depr <- read.csv(".../Data/depression_data.csv", header=T)
# Remove missing values in Diagnosis
depr <- na.omit(depr)
# Since we'll use logistic regression for this data set. Convert the response as 0-1 or a factor.</pre>
```

```
depr$Diagnosis <- 1*(depr$Diagnosis == 1)</pre>
names(depr)
## [1] "ID"
                           "Gender"
                                               "Guardian_status"
                                                                   "Cohesion_score"
## [5] "Depression_score" "Diagnosis"
depr.glm1 <- glm(Diagnosis ~ Gender + Guardian_status + Cohesion_score,</pre>
                 family = binomial, data=depr)
cv1 <- cv.glm(depr, depr.glm1, cost=loss) # By default, K=n, LOOCV</pre>
# Drop quardian?
depr.glm2 <- glm(Diagnosis ~ Gender + Cohesion_score,</pre>
                 family = binomial, data=depr)
cv2 <- cv.glm(depr, depr.glm2, cost=loss) # By default, K=n, LOOCV
cbind(WithGuardianER=cv1$delta, NoGuardianER=cv2$delta)
##
        WithGuardianER NoGuardianER
## [1,]
             0.1615721
                           0.1572052
## [2,]
             0.1615721
                           0.1572005
```

Which model, with or without guadian, performs better based on CV prediction error rate?

## 5 Cross-validation in lda(), qda() and KNN.

Read the help file of lda(), qda() and knn.cv() for more details.

Examples of LOOCV in lda() and qda() can be found in previous R example.

You'll try knn.cv() in homework.