Statistics 452: Statistical Learning and Prediction

Chapter 5: Resampling Methods

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Resampling Methods

- Cross-validation and bootstrap.
- Grouped together because they involve random sampling of subsets of the data.
- ▶ But purpose is different: CV estimates the test error, bootstrap is used to estimate the variance of estimators.

Test Error

- ▶ Illustrate with squared error for quantitative *Y*.
- ► Training observations $\{(x_1, y_1), \dots, (x_n, y_n)\}$ are used to produce \hat{f} .
- ▶ If we had a large number of test observations (x_0, y_0) , the test MSE

$$Ave((y_0 - \hat{f}(x_0))^2)$$

reflects how well \hat{f} predicts new observations.

▶ With just a finite test set we get an *estimate* of the test error.

Validation

- ▶ This is what we have already been doing.
- ▶ We split our data into two parts, a training set and a validation, or hold-out set.
 - Use the training set for fitting and the validation set for estimating the test error.

Validation on the Auto Data

Split the Auto data in half.

```
library(tidyverse)
library(ISLR)
data(Auto)
Auto <- dplyr::select(Auto,mpg,horsepower)</pre>
n <- nrow(Auto)
set.seed(42)
# Split in half
train <- sort(sample(1:n,size=n/2)) # sorting not necessary</pre>
head(train)
## [1] 1 2 3 7 11 14
validn <- setdiff(1:n,train)</pre>
head(validn)
## [1] 4 5 6 8 9 10
Auto.train <- Auto[train.]
Auto.validn <- Auto[validn,] # same as Auto[-train,]
```

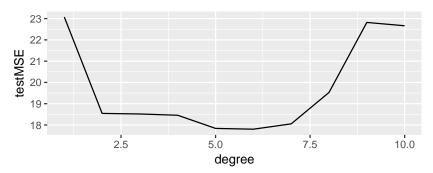
- Use the train half to train a polynomial model in horsepower and then estimate the test MSE on the validn half.
 - Software note: poly() returns polynomials and is useful in a model formula to save typing

```
afit <- lm(mpg ~ poly(horsepower,2),data=Auto.train)
yhat.v <- predict(afit,newdata=Auto.validn)
tMSE <- with(Auto.validn,mean((mpg - yhat.v)^2))
tMSE</pre>
```

```
## [1] 18.54359
```

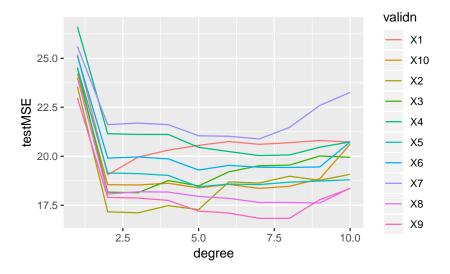
Validation to Select the Degree of Polynomial

```
testMSE <- function(dd,train,validn) {
   afit <- lm(mpg ~ poly(horsepower,dd),data=train)
   yhat.v <- predict(afit,newdata=validn)
   return(with(validn,mean((mpg - yhat.v)^2)))
}
nd <- 10; dd <- (1:nd); tm <- rep(NA,nd)
for(i in dd) {
   tm[i] <- testMSE(i,Auto.train,Auto.validn)
}
dMSE <- data.frame(degree=dd,testMSE= tm)
ggplot(dMSE,aes(x=degree,y=testMSE)) + geom_line()</pre>
```



Validation with Different Validation Sets

```
nValid <- 10
valid <- function() {</pre>
  n <- nrow(Auto)
  train <- sample(1:n,size=n/2)</pre>
  Auto.train <- Auto[train,]</pre>
  Auto.validn <- Auto[-train,]
  tm <- rep(NA,nd)
  for(i in dd) {
    tm[i] <- testMSE(i,Auto.train,Auto.validn)</pre>
  tm
tMSE <- replicate(nValid, valid())</pre>
tMSE <- data.frame(degree=dd,tMSE)</pre>
tMSE <- gather(tMSE, validn, testMSE, X1:X10, -degree)</pre>
```



▶ Note the variability in the estimated test MSE.

Cross-Validation (CV)

- Rather than a single data split, do multiple splits into "folds" of approximately equal size.
 - ▶ Common numbers of folds are k = n, 10 and 5.
- ► Train on all but one hold-out fold, and test on the hold-out to get MSE_i; i = 1,..., k.
- ▶ Repeat for each fold and average the estimated test MSEs:

$$CV_{(k)} = \frac{1}{k} \sum_{i=1}^{k} MSE_i.$$

Test Error vs Expected Test Error

 A quantity related to test error is the expected test error, obtained by averaging the test error over repeated training sets,

$$Ave[Ave((y_0 - \hat{f}(x_0))^2)]$$

where the outer Ave is over the training sets that give us \hat{f} .

- Picture this as repeating the following:
 - 1. Sample training and test data
 - 2. Train the model, and use on the test data to obtain the average squared error

and averaging the average from step 2.

- Cross validation estimates the expected test error.
 - A procedure with good expected test error tends to have good test error.

Leave-Out-One CV (LOOCV)

- ▶ Break the data into *n* folds, with one observation in each fold.
- Computational trick for a linear model fit by least squares:

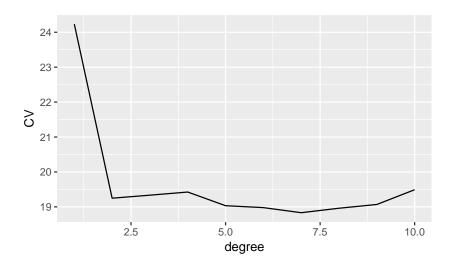
$$CV_{(n)} = \frac{1}{n} \sum_{i=1}^{n} \left(\frac{y_i - \hat{y}_i}{1 - h_i} \right)^2,$$

where \hat{y}_i is the fitted value from the least squares fit and h_i is the leverage of the *i*th observation.

LOOCV on Auto Data

```
loocv <- function(dd) {
   CVn <- rep(NA,length(dd))
   for(i in dd) {
      fit <- lm(mpg ~ poly(horsepower,i),data=Auto)
      hh <- hatvalues(fit)
      ff <- fitted.values(fit)
       CVn[i] <- with(Auto,mean(((mpg-ff)/(1-hh))^2))
   }
   CVn
}
cv.err <- loocv(dd)
cv.err <- data.frame(degree=dd,CV=cv.err)</pre>
```

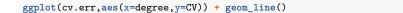


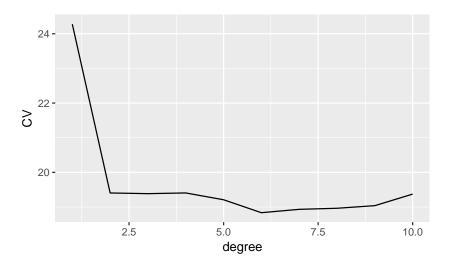


10-Fold CV on Auto Data

- ► Can use a function cv.glm() from the boot package.
 - ▶ Uses output from glm().
 - ▶ glm default is normal errors; i.e., lm().

```
library(boot)
cv.err <- rep(NA,nd)
set.seed(123)
for(i in dd) {
   fit <- glm(mpg ~ poly(horsepower,i),data=Auto)
    cc <- cv.glm(Auto,fit,K=10)
    cv.err[i] <- cc$delta[1]
}
cv.err <- data.frame(degree=dd,CV=cv.err)</pre>
```





Bias-Variance Trade-Off for k-Fold CV

- ▶ In general, computation of *k*-fold CV increases with *k*.
- ▶ But more important is the accuracy of the CV estimator as a function of *k*.
- ▶ There are two components to accuracy, bias and variance.
 - ► It can be shown that the bias of the CV estimator of the test error *decreases* as *k* increases.
 - It can be shown that the variance of the CV estimator increases with k.

Bias

- If data splitting results in a training set that is small, the error of the statistical learning method will be larger than if we fit to all data.
- ▶ Implies an upward bias in the estimate of the test error.
- ➤ On the other extreme, LOOCV uses almost all the data to train, and so will have almost no bias.

Variance

- ▶ This is harder to reason through.
- ► The LOOCV estimate is an average of many squared errors that are (i) highly variable, and (ii) positively correlated.
 - Averaging many things is good.
 - The positive correlation arrises from using mostly the same data to fit the model each time.
- For k-fold CV with smaller k, we average fewer MSEs that are
 (i) less variable and (ii) less correlated.
- ▶ Which "wins"? Values k = 5 or 10 have been shown to work well empirically and are recommended.

Simulation Example

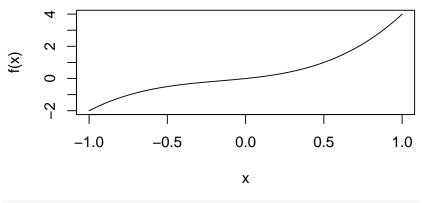
- We can illustrate the bias and variance of the CV estimator of the test error with one similation model.
- Simulate from the model

$$Y = X + X^2 + 2X^3 + \epsilon$$

for $\epsilon \sim N(0,1)$ and a fixed grid of X-values.

- Fit a linear regression.
- ▶ View the contributions to $CV_{(n)}$ and $CV_{(10)}$ and the variance of these two estimators.

```
n <- 100
x <- seq(-1,1,length=n)
plot(x,x+x^2+2*x^3,ylab="f(x)",type="l")</pre>
```

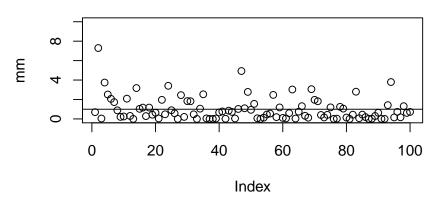


```
simdat <- function() {
  y<-rnorm(n,mean=x+x^2+2*x*3,sd=1)
  data.frame(y=y,x=x)
}</pre>
```

```
# LOOCV contributions for data set of size n.
# LOOCV estimate of test error is their average.
loocv.contrib <- function() {
    MSE <- rep(NA,n)
    dat <- simdat()
    for(i in 1:n) {
        fit <- lm(y-x,data=dat,subset=(-i))
            newdat <- data.frame(x=dat$x[i])
        pp <- predict(fit,newdata=newdat)
        MSE[i] <- (dat$y[i] - pp)^2
    }
    MSE
}</pre>
```

▶ Repeat the following a few times

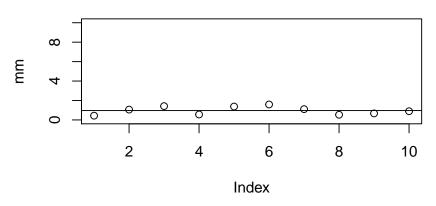
```
mm <- loocv.contrib()
plot(mm,ylim=c(0,10))
abline(h=mean(mm)) # LOOCV estimate of test error</pre>
```



```
# 10-fold CV contributions for data set of size n
cv.contrib <- function(n=100,k=10) {
    MSE <- rep(NA,k)
    dat <- simdat()
    inds <- sample(1:n)
    for(i in 1:(n/k)) {
        ss <- inds[(i-1)*k + (1:(n/k))]
        fit <- lm(y-x,data=dat,subset=(-ss))
        newdat <- data.frame(x=dat$x[ss])
        pp <- predict(fit,newdata=newdat)
        MSE[i] <- mean((dat$y[ss] - pp)^2)
    }
    MSE
}</pre>
```

Repeat the following a few times

```
mm <- cv.contrib()
plot(mm,ylim=c(0,10))
abline(h=mean(mm)) # Estimate of test error</pre>
```



Hard to see how the CV estimates (horzontal lines) vary over samples, so record them over many simulations.

```
set.seed(123)
NREPS <- 1000
LOOCVres <- CVres <- rep(NA, NREPS)
for(i in 1:NREPS) {
  LOOCVres[i] <- mean(loocv.contrib())
  CVres[i] <- mean(cv.contrib())</pre>
mean(LOOCVres): var(LOOCVres):mean(CVres): var(CVres)
## [1] 1.122344
## [1] 0.02634004
## [1] 1.124939
## [1] 0.02689808
```

 Can see suggestion of lower bias from LOOCV, but not lower variance of 10-fold CV for this scenario.

CV on Classification Problems

- We have illustrated the idea behind CV when the response is quantitative.
- ► We then use the MSE (mean squared error) to quantify test error.
- ► For classification problems we measure the error of a procedure by the misclassification error.
- For example,

$$CV_{(n)} = \frac{1}{n} \sum_{i=1}^{n} I(y_i \neq \hat{y}_i)$$

See the text for examples.

The Bootstrap

► The bootstrap uses resampling to quantify uncertainty in an estimator.

Assumptions and sampling distributions

- Under model assumptions, the sampling distribution of the statistics used for inference are known.
 - Sampling distribution: Distribution of a statistic over repeated samples of data from the population.
 - For regression coefficients, the sampling distribution leads to t-tests and Cls
- ► The bootstrap is a data-driven approach to approximating the sampling distribution of inferential statistics.
 - ► Find the distribution of a statistic over repeated samples of data **from the original sample**.
 - ▶ Reasonable if original sample is representative of the population.
 - ▶ Base inference on the bootstrap approximate distribution.

Advantages

- ▶ Bootstrap may give reasonable uncertainty estimates when assumptions for traditional inference don't hold.
- ▶ We can expand the definition of the procedure used to obtain the estimates to include variable selection and/or other smoothing (more on this later).

Resampling

- Resampling means drawing samples, with replacement, from the original sample.
 - ▶ E.G., drawing cars, with replacement.

```
set.seed(42)
n <- nrow(Auto)
Autos <- data.frame(index=1:n,Auto)
resamAuto <- sample_n(Auto,size=n,replace=TRUE)
head(resamAuto)</pre>
```

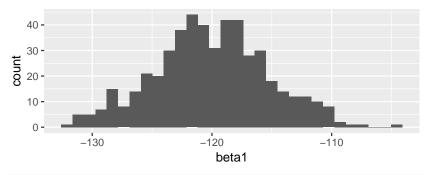
```
## mpg horsepower
## 364 22.4 110
## 373 27.0 90
## 114 21.0 107
## 328 36.4 67
## 254 20.5 95
## 206 28.0 75
```

Bootstrap Standard Errors

- Resample some number B times.
- For each resample compute the estimates.
- ► Take the sample SD of the bootstrap estimates.

```
B <- 500; beta1Boot <- rep(NA,B)
for(i in 1:B) {
    rAuto <- sample_n(Auto, size=n, replace=TRUE)
    fit <- lm(mpg-poly(horsepower,2), data=rAuto)
    beta1Boot[i] <- coefficients(fit)[2]
}
beta1Boot <- data.frame(beta1=beta1Boot)</pre>
```

```
ggplot(beta1Boot,aes(x=beta1)) + geom_histogram()
```



```
with(beta1Boot,sd(beta1))
```

```
## [1] 4.72912
```

```
fit <- lm(mpg~poly(horsepower,2),data=Auto)
round(summary(fit)$coefficients[2,],4)</pre>
```

```
## Estimate Std. Error t value Pr(>|t|)
## -120.1377 4.3739 -27.4668 0.0000
```