Chp5&6 HW

Wendy Liang 2/13/2021

6.2

- a. iii. Less flexible and hence will give improved prediction accu racy when its increase in bias is less than its decrease in variance.
- b. iii. Less flexible and hence will give improved prediction accu racy when its increase in bias is less than its decrease in variance.
- c. ii. More flexible and hence will give improved prediction accu racy when its increase in variance is less than its decrease in bias.

5.8

(a)

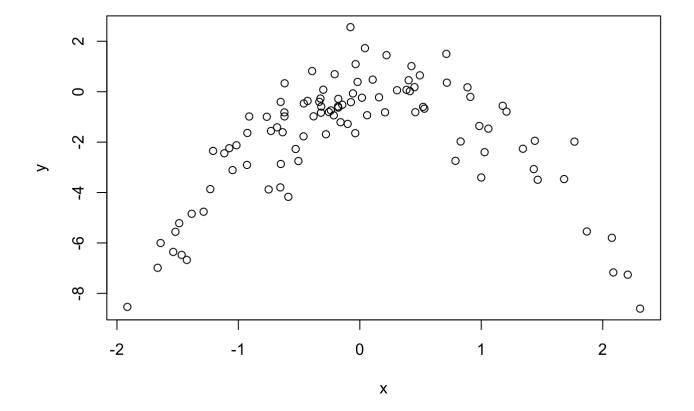
```
set.seed(1)
y <- rnorm(100)
x <- rnorm(100)
y <- x - 2 * x^2 + rnorm(100)</pre>
```

n=100 and p=2

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

(b)

```
plot(x, y)
```



The data obviously suggests a curved relationship

(c)

[1] 1.086596

```
library(boot)
set.seed(1)
Data <- data.frame(x, y)

#1
fit.glm.1 <- glm(y ~ x)
cv.glm(Data, fit.glm.1)$delta[1]

## [1] 5.890979

#2
fit.glm.2 <- glm(y ~ poly(x, 2))
cv.glm(Data, fit.glm.2)$delta[1]</pre>
```

```
#3
fit.glm.3 <- glm(y ~ poly(x, 3))
Logading [MathJax]/jax/output/H1ML-Css/jax.js]
```

```
## [1] 1.102585
```

```
#4
fit.glm.4 <- glm(y ~ poly(x, 4))
cv.glm(Data, fit.glm.4)$delta[1]</pre>
```

```
## [1] 1.114772
```

(d)

```
set.seed(10)
#1
fit.glm.1 <- glm(y ~ x)
cv.glm(Data, fit.glm.1)$delta[1]</pre>
```

```
## [1] 5.890979
```

```
#2
fit.glm.2 <- glm(y ~ poly(x, 2))
cv.glm(Data, fit.glm.2)$delta[1]</pre>
```

```
## [1] 1.086596
```

```
#3
fit.glm.3 <- glm(y ~ poly(x, 3))
cv.glm(Data, fit.glm.3)$delta[1]</pre>
```

```
## [1] 1.102585
```

```
#4
fit.glm.4 <- glm(y ~ poly(x, 4))
cv.glm(Data, fit.glm.4)$delta[1]</pre>
```

```
## [1] 1.114772
```

(e)

The LOOCV estimate for the test MSE is minimum for "fit.glm.2", we can also see clearly in (b) plot that the relation between "x" and "y" is quadratic.

(f)

```
summary(fit.glm.4)
```

```
##
## Call:
## glm(formula = y \sim poly(x, 4))
## Deviance Residuals:
##
      Min
                1Q Median
                                  3Q
                                          Max
## -2.8914 -0.5244 0.0749
                              0.5932
                                       2.7796
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.8277 0.1041 -17.549
                                            <2e-16 ***
## poly(x, 4)1
              2.3164
                           1.0415
                                    2.224
                                            0.0285 *
## poly(x, 4)2 -21.0586
                           1.0415 -20.220
                                            <2e-16 ***
## poly(x, 4)3 -0.3048
                           1.0415 -0.293
                                            0.7704
## poly(x, 4)4 -0.4926 1.0415 -0.473
                                           0.6373
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 1.084654)
##
##
      Null deviance: 552.21 on 99
                                   degrees of freedom
## Residual deviance: 103.04 on 95 degrees of freedom
## AIC: 298.78
##
## Number of Fisher Scoring iterations: 2
```

The p-values show that the linear and quadratic terms are statistically significant and that the cubic and 4th degree terms are not statistically significant.

6.10

(a)

```
set.seed(1)
x <- matrix(rnorm(1000 * 20), 1000, 20)
b <- rnorm(20)
b[3] <- 0
b[4] <- 0
b[9] <- 0
b[19] <- 0
eps <- rnorm(1000)
y <- x %*% b + eps</pre>
```

(b)

```
train <- sample(seq(1000), 100, replace = FALSE)

test <- -train

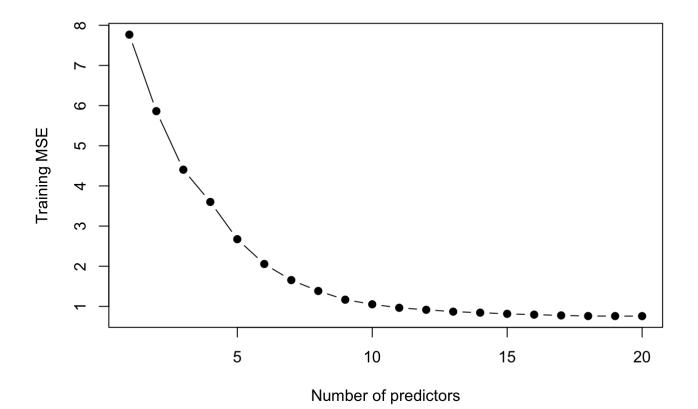
x.train <- x[train, ]

x.test <- x[test, ]

y.train <- y[train]

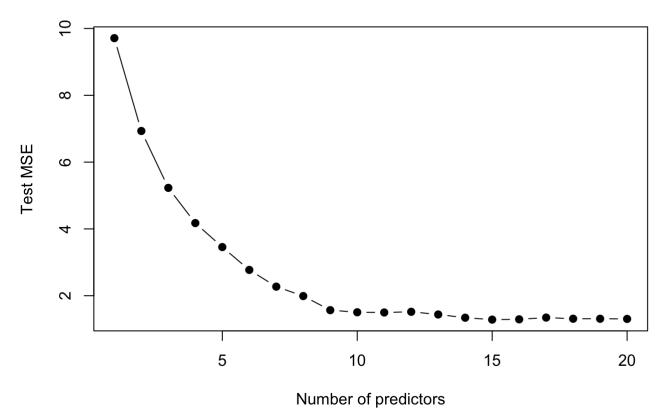
Loading [MathJax] (jax/output/HTML-CSS/jax.js
```

```
library(leaps)
data.train <- data.frame(y = y.train, x = x.train)
regfit.full <- regsubsets(y ~ ., data = data.train, nvmax = 20)
train.mat <- model.matrix(y ~ ., data = data.train, nvmax = 20)
val.errors <- rep(NA, 20)
for (i in 1:20) {
    coefi <- coef(regfit.full, id = i)
        pred <- train.mat[, names(coefi)] %*% coefi
    val.errors[i] <- mean((pred - y.train)^2)
}
plot(val.errors, xlab = "Number of predictors", ylab = "Training MSE", pch = 19, type = "b")</pre>
```



(d)

```
data.test <- data.frame(y = y.test, x = x.test)
test.mat <- model.matrix(y ~ ., data = data.test, nvmax = 20)
val.errors <- rep(NA, 20)
for (i in 1:20) {
    coefi <- coef(regfit.full, id = i)
    pred <- test.mat[, names(coefi)] %*% coefi
    val.errors[i] <- mean((pred - y.test)^2)
}
Loading[MathJax]/jax/output/HTML_CSS/jax.js
plot(val.errors, xlab = "Number of predictors", ylab = "Test MSE", pch = 19, type = "b")</pre>
```



(e)

```
which.min(val.errors)
```

```
## [1] 15
```

(f)

coef(regfit.full, which.min(val.errors))

```
(Intercept)
##
                          x.2
   -0.003933937
##
                  0.359127426
                                0.202707344
                                              1.036265913 -0.253843053 -1.282753293
##
            x.8
                         x.11
                                       x.12
                                                     x.13
                                                                   x.14
                                                                                 x.15
##
    0.691581077
                  0.895769881
                                0.526887865 - 0.207638251 - 0.507929833 - 0.892604795
##
           x.16
                         x.17
                                       x.18
## -0.343062241
                  0.184479252
                                1.646950451 -1.060191640
```

The best model caught all zeroed out coefficients.

(g)

```
val.errors <- rep(NA, 20)
x_cols = colnames(x, do.NULL = FALSE, prefix = "x.")
for (i in 1:20) {
    coefi <- coef(regfit.full, id = i)
      val.errors[i] <- sqrt(sum((b[x_cols %in% names(coefi)] - coefi[names(coefi) %in% x_c ols])^2) + sum(b[!(x_cols %in% names(coefi))])^2)
}
plot(val.errors, xlab = "Number of coefficients", ylab = "Error between estimated and tr ue coefficients", pch = 19, type = "b")</pre>
```

