## Resampling and regularization Homework

Jiachen Feng

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5.8

(a)

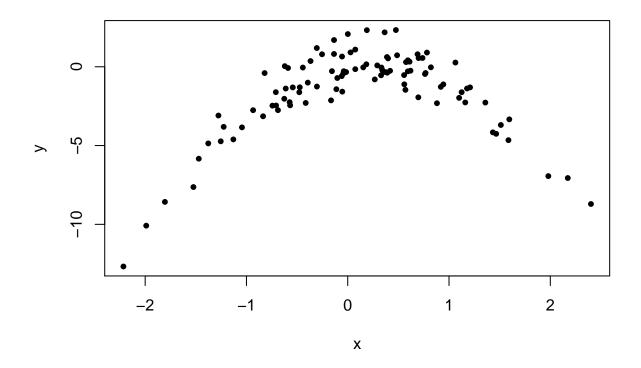
```
set.seed(1)
x <- rnorm(100)
y <- x-2*x^2+rnorm(100)</pre>
```

n is 100, p is 2.

$$y = x - x^2 + \epsilon$$

(b)

plot(x,y,pch=20)



The scatterplot is like a quadratic function.

(c)

```
library(boot)
set.seed(100)
data <- data.frame(x,y)</pre>
mod1 \leftarrow glm(y~x)
cv.glm(data,mod1)$delta[1]
## [1] 7.288162
mod2 \leftarrow glm(y poly(x, 2))
cv.glm(data,mod2)$delta[1]
## [1] 0.9374236
mod3 \leftarrow glm(y poly(x,3))
cv.glm(data,mod3)$delta[1]
## [1] 0.9566218
mod4 \leftarrow glm(y poly(x,4))
cv.glm(data,mod4)$delta[1]
## [1] 0.9539049
 (d)
set.seed(200)
mod1 \leftarrow glm(y~x)
cv.glm(data,mod1)$delta[1]
## [1] 7.288162
mod2 \leftarrow glm(y poly(x, 2))
cv.glm(data,mod2)$delta[1]
## [1] 0.9374236
mod3 \leftarrow glm(y poly(x,3))
cv.glm(data,mod3)$delta[1]
## [1] 0.9566218
mod4 \leftarrow glm(y poly(x,4))
cv.glm(data,mod4)$delta[1]
## [1] 0.9539049
Same.
 (e) model 2. The function is quadratic, so that this model fits best.
  (f)
summary(mod1)
##
## Call:
## glm(formula = y ~ x)
##
## Deviance Residuals:
##
       Min
                                       ЗQ
                   1Q
                        Median
                                                Max
## -9.5161 -0.6800
                        0.6812 1.5491
                                             3.8183
##
```

```
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.6254
                        0.2619 -6.205 1.31e-08 ***
                0.6925
                           0.2909 2.380 0.0192 *
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 6.760719)
##
##
      Null deviance: 700.85 on 99 degrees of freedom
## Residual deviance: 662.55 on 98 degrees of freedom
## AIC: 478.88
## Number of Fisher Scoring iterations: 2
summary(mod2)
##
## Call:
## glm(formula = y \sim poly(x, 2))
## Deviance Residuals:
      Min
                10
                    Median
                                 30
## -1.9650 -0.6254 -0.1288 0.5803
                                      2.2700
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.5500 0.0958 -16.18 < 2e-16 ***
## poly(x, 2)1 6.1888
                                  6.46 4.18e-09 ***
                          0.9580
## poly(x, 2)2 -23.9483
                          0.9580 -25.00 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 0.9178258)
##
      Null deviance: 700.852 on 99 degrees of freedom
## Residual deviance: 89.029 on 97 degrees of freedom
## AIC: 280.17
## Number of Fisher Scoring iterations: 2
summary(mod3)
##
## Call:
## glm(formula = y \sim poly(x, 3))
##
## Deviance Residuals:
      Min 1Q Median
                                 3Q
                                         Max
## -1.9765 -0.6302 -0.1227
                            0.5545
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.55002 0.09626 -16.102 < 2e-16 ***
## poly(x, 3)1 6.18883
                           0.96263 6.429 4.97e-09 ***
```

```
## poly(x, 3)2 -23.94830
                            0.96263 -24.878 < 2e-16 ***
                            0.96263
                                      0.274
                                                0.784
## poly(x, 3)3
                 0.26411
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
  (Dispersion parameter for gaussian family taken to be 0.9266599)
##
##
##
       Null deviance: 700.852 on 99 degrees of freedom
## Residual deviance: 88.959
                               on 96 degrees of freedom
## AIC: 282.09
##
## Number of Fisher Scoring iterations: 2
summary(mod4)
##
## Call:
##
  glm(formula = y \sim poly(x, 4))
## Deviance Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
                                           Max
##
  -2.0550
           -0.6212 -0.1567
                               0.5952
                                         2.2267
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
                            0.09591 -16.162 < 2e-16 ***
## (Intercept)
                -1.55002
                 6.18883
                            0.95905
                                      6.453 4.59e-09 ***
## poly(x, 4)1
## poly(x, 4)2 -23.94830
                            0.95905 - 24.971
                                             < 2e-16 ***
## poly(x, 4)3
                 0.26411
                                                0.784
                            0.95905
                                      0.275
## poly(x, 4)4
                 1.25710
                            0.95905
                                      1.311
                                                0.193
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
  (Dispersion parameter for gaussian family taken to be 0.9197797)
##
##
       Null deviance: 700.852
##
                               on 99
                                      degrees of freedom
## Residual deviance: 87.379
                               on 95 degrees of freedom
## AIC: 282.3
##
## Number of Fisher Scoring iterations: 2
All of the four models are statistically significant.
```

## 6.2

- (a)
  - i. Wrong.
- ii. Wrong.
- iii. Right.
- iv. Wrong. It is less flexible because the complexity of lasso is related to  $\lambda$ .
- (b) Ridge regression is less flexible and hence will give improved prediction accuracy when its increase in bias is less than its decrease in variance.
- (c) Non-linear methods are more flexible and hence will give improved prediction accuracy when its increase in variance is less than its decrease in bias.

## 6.10

(a)

```
set.seed(1000)
mat <- matrix(rnorm(1000 * 20), 1000, 20)</pre>
p <- rnorm(20)
p[1] <- 0
p[2] <- 0
p[3] <- 0
epsilon <- rnorm(1000)
y <- mat %*% p + epsilon
 (b)
set.seed(1000)
mat1 <- matrix(rnorm(100 * 20), 100, 20)
p1 <- rnorm(20)
p1[1] <- 0
p1[2] <- 0
p1[3] <- 0
epsilon1 <- rnorm(100)</pre>
train <- mat1 %*% p1 + epsilon1
mat2 <- matrix(rnorm(900 * 20), 900, 20)
p2 <- rnorm(20)
p2[1] <- 0
p2[2] <- 0
p2[3] <- 0
epsilon2 <- rnorm(900)</pre>
test <- mat2 %*% p2 + epsilon2</pre>
```

(c)