Homework 6

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2025-06-15

1

- i. nonlinear
- ii. nonlinear
- iii. ridge regression
- iv. lasso

2

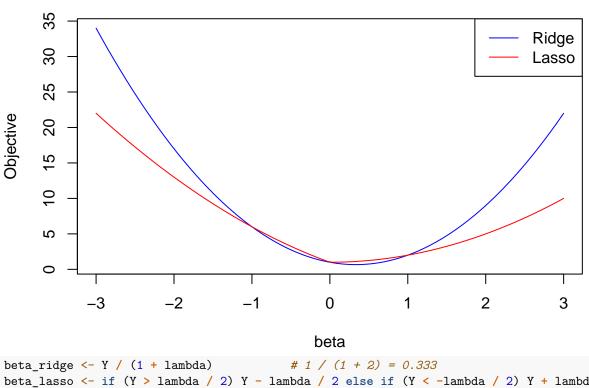
 \mathbf{a}

```
Y <- 1
lambda <- 2
beta <- seq(-3, 3, length = 100)

RSS <- (Y - beta)^2
ridge_obj <- RSS + lambda * beta^2
lasso_obj <- RSS + lambda * abs(beta)

plot(beta, ridge_obj, type = "l", col = "blue", ylab = "Objective", main = "Ridge vs Lasso")
lines(beta, lasso_obj, col = "red")
legend("topright", legend = c("Ridge", "Lasso"), col = c("blue", "red"), lty = 1)</pre>
```

Ridge vs Lasso



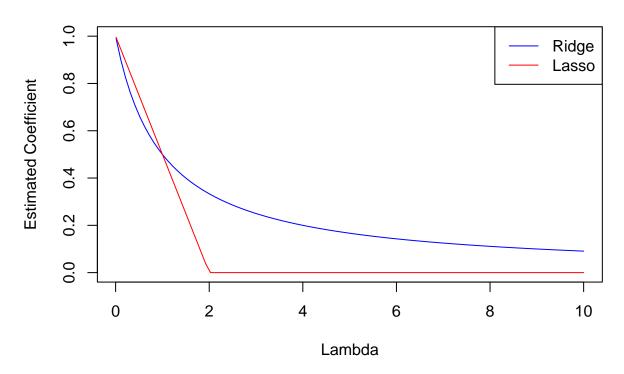
```
beta_ridge <- Y / (1 + lambda)  # 1 / (1 + 2) = 0.333
beta_lasso <- if (Y > lambda / 2) Y - lambda / 2 else if (Y < -lambda / 2) Y + lambda / 2 else 0
beta_ridge</pre>
```

[1] 0.3333333 beta_lasso

[1] 0

b

Ridge vs Lasso Estimates



3

 \mathbf{a}

```
set.seed(1234)
x <- rnorm(100)
error <- rnorm(100)</pre>
```

b

```
beta0 <- 1
beta1 <- 2
beta2 <- 3
beta3 <- 4
Y <- beta0 + beta1*x + beta2*x^2 + beta3*x^3 + error</pre>
```

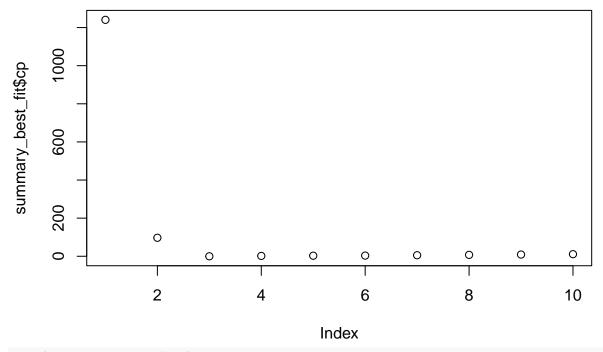
 \mathbf{c}

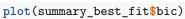
```
library(leaps)

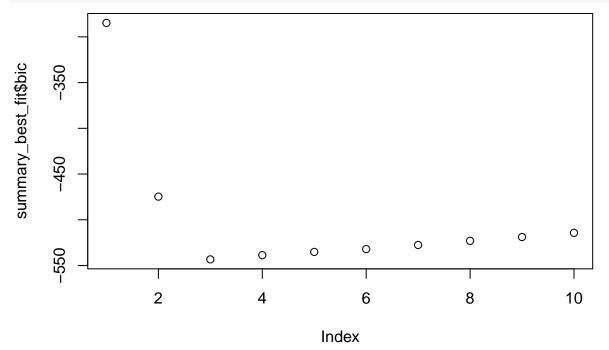
x_poly <- data.frame(poly(x,10, raw = T))
data <- data.frame(Y, x_poly)
best_fit <- regsubsets(Y ~., data = data, nvmax = 10)

summary_best_fit <- summary(best_fit)

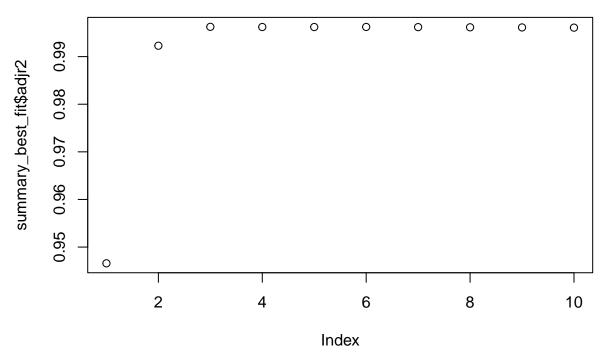
plot(summary_best_fit$cp)</pre>
```







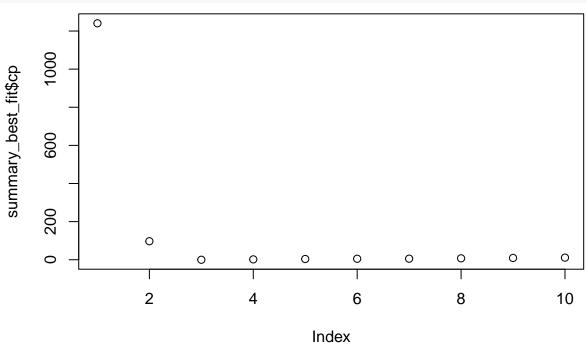
plot(summary_best_fit\$adjr2)

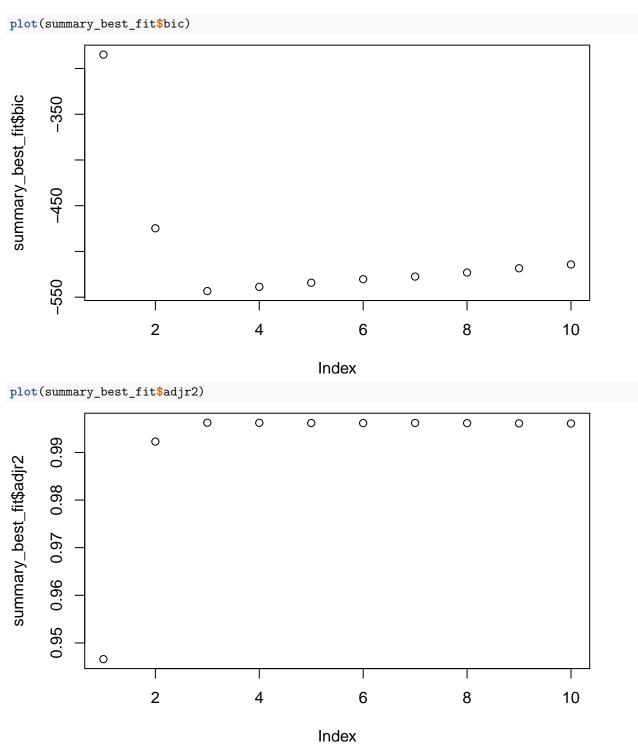


using CP, it looks like 3 predictors is the best. same for BIC. 3 predictors looks best. ADJR2 also suggest 3 predictors is the best. so the best model would be $Y = beta0 + beta1 \times 1 + beta2 \times 1^2 + beta3 \times 1^3$

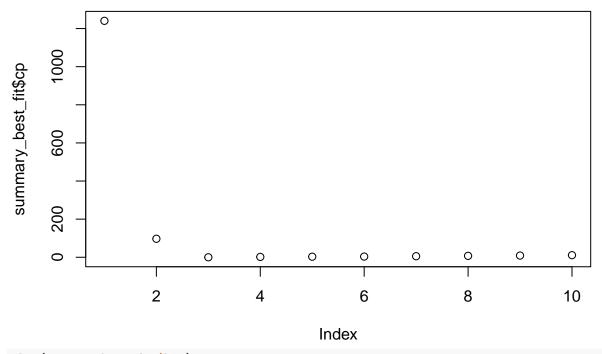
```
\mathbf{d}
```

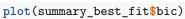
```
best_fit <- regsubsets(Y ~., data=data, nvmax = 10, method = "forward")
summary_best_fit <- summary(best_fit)
plot(summary_best_fit$cp)</pre>
```

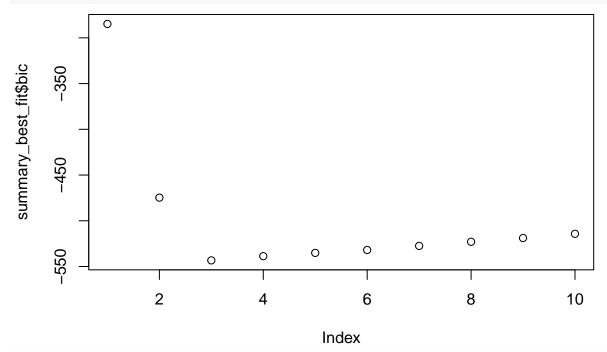




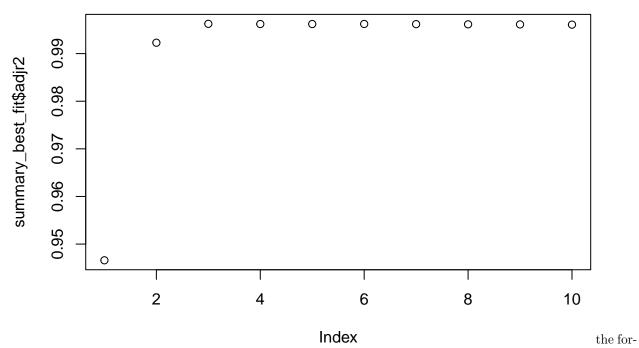
```
best_fit <- regsubsets(Y ~., data=data, nvmax = 10, method = "backward")
summary_best_fit <- summary(best_fit)
plot(summary_best_fit$cp)</pre>
```







plot(summary_best_fit\$adjr2)



ward and backward models suggest 3 predictors as well

 \mathbf{e}

```
library(glmnet)

## Loading required package: Matrix

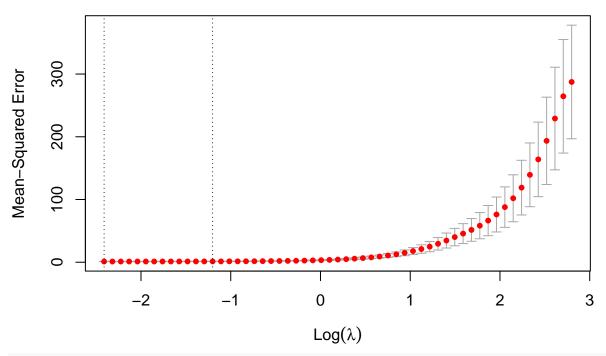
## Loaded glmnet 4.1-9

X_matrix <- model.matrix(Y ~ ., data = data)[, -1]

cv_lasso <- cv.glmnet(X_matrix, Y, alpha = 1)

plot(cv_lasso)</pre>
```

3 3 3 3 3 3 4 4 4 4 4 4 4 3 3 1 1 1 1



```
coef(cv_lasso, s = "lambda.min")
```

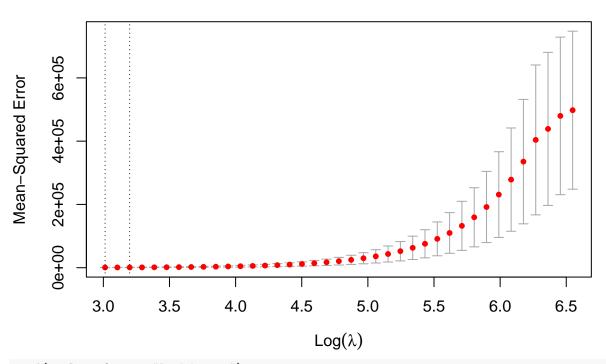
```
## 11 x 1 sparse Matrix of class "dgCMatrix"
## (Intercept) 1.178705
                1.847133
## X1
## X2
                2.838895
                4.028284
## X3
## X4
## X5
## X6
## X7
## X8
## X9
## X10
x4 and above got eliminated
```

 \mathbf{f}

```
Y <- beta0 + 7 * x^7 + error

data2 <- data.frame(Y, poly(x, 10, raw = TRUE))
best_fit2 <- regsubsets(Y ~ ., data = data2, nvmax = 10)
summary_best2 <- summary(best_fit2)

X_matrix2 <- model.matrix(Y ~ ., data = data2)[, -1]
cv_lasso2 <- cv.glmnet(X_matrix2, Y, alpha = 1)
plot(cv_lasso2)</pre>
```



```
coef(cv_lasso2, s = "lambda.min")

## 11 x 1 sparse Matrix of class "dgCMatrix"

## s0

## (Intercept) 2.952989

## X1 .
```

X7 6.795857 ## X8 .

X9 . ## X10 .

4

X2

 \mathbf{a}

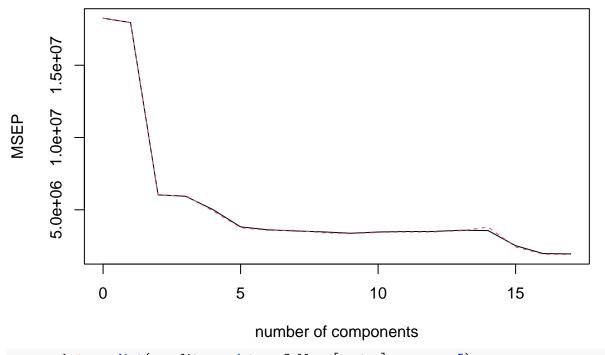
v purrr

1.0.2

```
load("../data/College.rda")
library(ISLR)
library(tidyverse)
## -- Attaching core tidyverse packages ---
                                                       ----- tidyverse 2.0.0 --
## v dplyr
               1.1.4
                                     2.1.5
                         v readr
## v forcats
               1.0.0
                         v stringr
                                     1.5.1
## v ggplot2
               3.5.1
                         v tibble
                                     3.2.1
## v lubridate 1.9.4
                         v tidyr
                                     1.3.1
```

```
## -- Conflicts -----
                                               ## x tidyr::expand() masks Matrix::expand()
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
## x tidyr::pack() masks Matrix::pack()
## x tidyr::unpack() masks Matrix::unpack()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
set.seed(1234)
train <- sample(1:nrow(College), nrow(College)/2)</pre>
test <- setdiff(1:nrow(College), train)</pre>
train_x <- model.matrix(Apps ~ ., data = College)[train, -1]</pre>
test_x <- model.matrix(Apps ~ ., data = College)[test, -1]</pre>
train_y <- College$Apps[train]</pre>
test_y <- College$Apps[test]</pre>
lm fit <- lm(Apps ~ ., data = College, subset = train)</pre>
pred_lm <- predict(lm_fit, newdata = College[test, ])</pre>
mean((pred_lm - test_y)^2)
## [1] 998713.2
b
ridge_fit <- cv.glmnet(train_x, train_y, alpha = 0)</pre>
ridge_pred <- predict(ridge_fit, s = ridge_fit$lambda.min, newx = test_x)</pre>
mean((ridge_pred - test_y)^2)
## [1] 963730.7
\mathbf{c}
lasso_fit <- cv.glmnet(train_x, train_y, alpha = 1)</pre>
lasso_pred <- predict(lasso_fit, s = lasso_fit$lambda.min, newx = test_x)</pre>
mean((lasso_pred - test_y)^2)
## [1] 965527.3
d
library(pls)
## Attaching package: 'pls'
## The following object is masked from 'package:stats':
##
##
       loadings
pcr_fit <- pcr(Apps ~ ., data = College, subset = train, scale = TRUE, validation = "CV")</pre>
validationplot(pcr_fit, val.type = "MSEP")
```

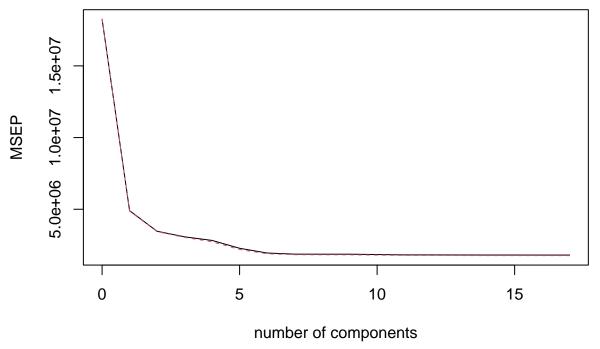
Apps



```
pcr_pred <- predict(pcr_fit, newdata = College[test, ], ncomp = 5)
mean((pcr_pred - test_y)^2)
## [1] 1734251
pls fit <- plsr(Apps ~ ... data = College, subset = train, scale = TRUE, validation = "CV")</pre>
```

pls_fit <- plsr(Apps ~ ., data = College, subset = train, scale = TRUE, validation = "CV")
validationplot(pls_fit, val.type = "MSEP")</pre>

Apps



```
pls_pred <- predict(pls_fit, newdata = College[test, ], ncomp = 5)
mean((pls_pred - test_y)^2)</pre>
```

[1] 1061050