STA 235H - Model Selection II: Stepwise and Shrinkage

Fall 2022

McCombs School of Business, UT Austin

Last class



- Started with our prediction chapter
 - o Bias vs. Variance
 - Validation set approach and Crossvalidation
 - How to choose a model for a continuous outcome (RMSE)

Today: Continuing our journey

- Some ways to select models:
 - Stepwise selection
 - Shrinkage methods: Ridge and Lasso.



One step at a time

Stepwise selection

- We have seen how to choose between some given models. But what if we want to test all possible models?
- Stepwise selection: Computationally-efficient algorithm to select a model based on the data we have (subset selection).

Algorithm for forward stepwise selection:

- 1. Start with the *null model*, M_0 (no predictors)
- 2. For $k = 0, \ldots, p 1$:
 - a) Consider all p-k models that augment M_k with one additional predictor.
 - b) Choose the *best* among these p-k models and call it M_{k+1} .
- 3. Select the single best model from M_0, \ldots, M_p using CV.

Backwards stepwise follows the same procedure, but starts with the full model.

Q1: Will forward stepwise subsetting yield the same results as backwards stepwise selection?

a) Yes

b) No

How do we do stepwise selection in R?

```
## nvmax RMSE Rsquared MAE RMSESD RsquaredSD MAESD
## 1 1 2.269469 0.6101788 1.850376 0.04630907 0.01985045 0.04266950
## 2 2.087184 0.6702660 1.639885 0.04260047 0.01784601 0.04623508
## 3 3 2.087347 0.6702094 1.640405 0.04258030 0.01804773 0.04605074
## 4 4 2.088230 0.6699245 1.641402 0.04270561 0.01808685 0.04620206
## 5 5 2.088426 0.6698623 1.641528 0.04276883 0.01810569 0.04624618
```

• Which one would you choose out of the 5 models? Why?

How do we do stepwise selection in R?

```
# We can see the number of covariates that is optimal to choose:
lm.fwd$bestTune
    nvmax
## 2
# And how does that model looks like:
summary(lm.fwd$finalModel)
## Subset selection object
## 5 Variables (and intercept)
         Forced in Forced out
## id
                       FALSE
             FALSE
## female
           FALSE
                       FALSE
## citv
        FALSE
                   FALSE
                   FALSE
       FALSE
## age
        FALSE
                       FALSE
## got
## 1 subsets of each size up to 2
## Selection Algorithm: forward
           id female city age got
# If we want to recover the coefficient names, we can use the coef() function:
coef(lm.fwd$finalModel, lm.fwd$bestTune$nvmax)
## (Intercept)
                    city
                                 got
                           -6.306371
     7.035888
                 2.570454
```

Honey, I shrunk the coefficients!

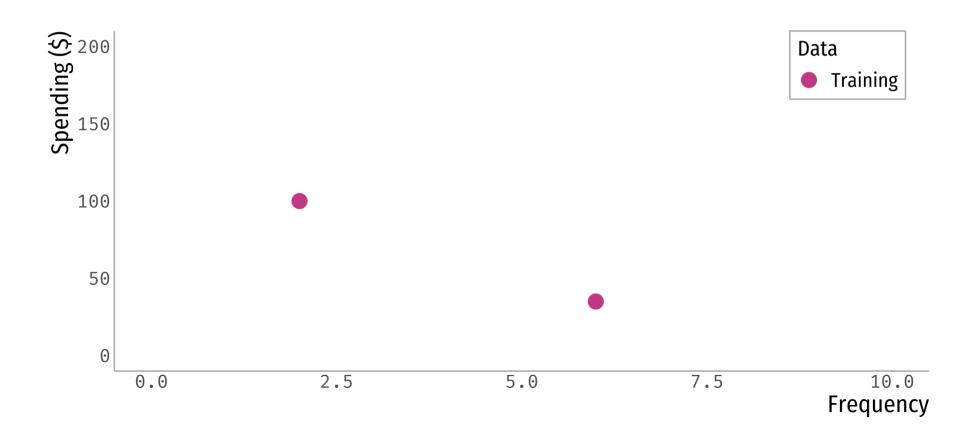
What is shrinkage?

- We just reviewed the **stepwise procedure**: Subsetting model selection approach.
 - Select k out of p total predictors
- Shrinkage (a.k.a Regularization): Fitting a model with all p predictors, but introducing bias (i.e. shrinking coefficients towards 0) for improvement in variance.
 - Ridge regression
 - Lasso regression

Let's build a ridge.

Ridge Regression: An example

• Predict spending based on frequency of visits to a website

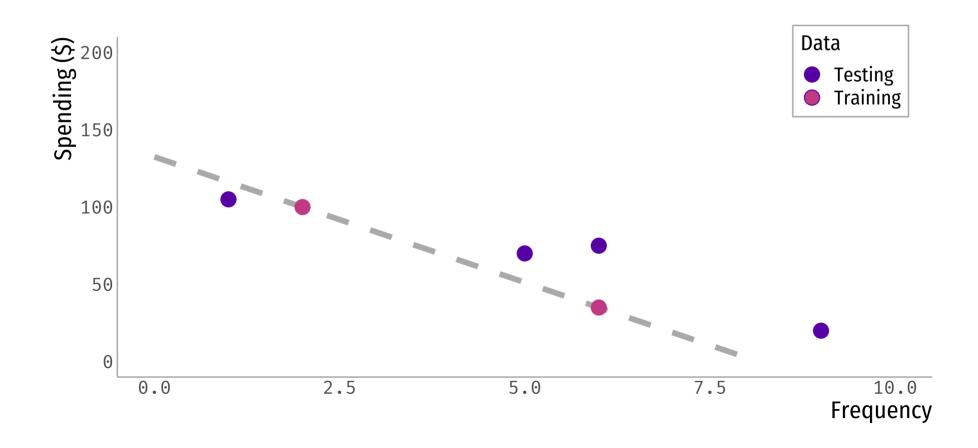


Ordinary Least Squares

• In an OLS: Minimize sum of squared-errors, i.e. $\min_{\beta} \sum_{i=1}^{n} (\operatorname{spend}_{i} - (\beta_{0} + \beta_{1} \operatorname{freq}_{i}))^{2}$

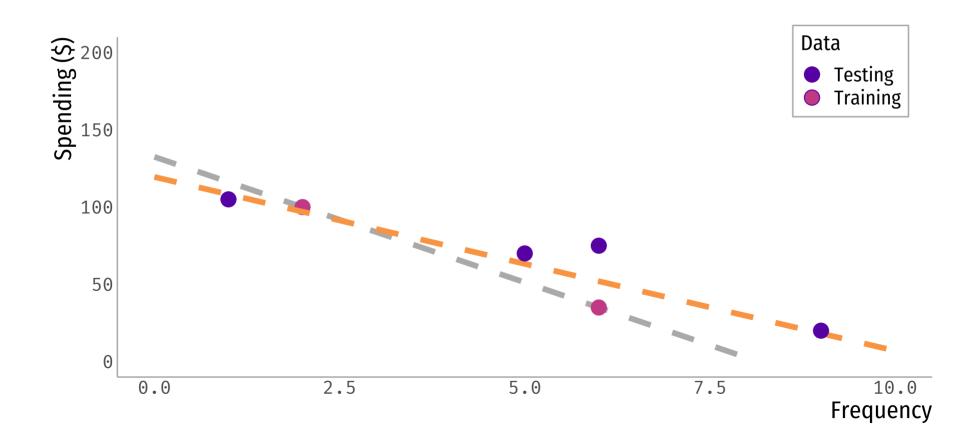
What about fit?

• Does the OLS fit the testing data well?



Ridge Regression

• Let's shrink the coefficients!: Ridge Regression



Ridge Regression: What does it do?

- Ridge regression introduces bias to reduce variance in the testing data set.
- In a simple regression (i.e. one regressor/covariate):

$$\min_{eta} \sum_{i=1}^n \underbrace{(y_i - eta_0 - x_i eta_1)^2}_{OLS}$$

Ridge Regression: What does it do?

- Ridge regression introduces bias to reduce variance in the testing data set.
- In a simple regression (i.e. one regressor/covariate):

$$\min_{eta} \sum_{i=1}^n \underbrace{(y_i - eta_0 - x_i eta_1)^2}_{OLS} + \underbrace{oldsymbol{\lambda} \cdot eta_1^2}_{RidgePenalty}$$

• λ is the penalty factor \rightarrow indicates how much we want to shrink the coefficients.

Q2: In general, which model will have smaller β coefficients?

a) A model with a larger λ

b) A model with a smaller λ

Remember... we care about accuracy in the testing dataset!

RMSE on the testing dataset: OLS

$$RMSE = \sqrt{rac{1}{4}\sum_{i=1}^{4}(\mathrm{spend}_i - (132.5 - 16.25 \cdot \mathrm{freq}_i))^2} = 28.36$$

RMSE on the testing dataset: Ridge Regression

$$RMSE = \sqrt{rac{1}{4}\sum_{i=1}^{4}(\mathrm{spend}_i - (119.5 - 11.25 \cdot \mathrm{freq}_i))^2} = 12.13$$

Ridge Regression in general

• For regressions that include more than one regressor:

$$\min_{eta} \sum_{i=1}^n \underbrace{(y_i - \sum_{k=0}^p x_i eta_k)^2}_{OLS} + \underbrace{\lambda \cdot \sum_{k=1}^p eta_k^2}_{RidgePenalty}$$

• In our previous example, if we had two regressors, female and freq:

$$\min_{eta} \sum_{i=1}^n (\operatorname{spend}_i - eta_0 - eta_1 \operatorname{female}_i - eta_2 \operatorname{freq}_i)^2 + \lambda \cdot (eta_1^2 + eta_2^2)$$

- Because the ridge penalty includes the β 's coefficients, scale matters:
 - Standardize variables (you will do that as an option in your code)

How do we choose λ ?

Cross-validation!

- 1) Choose a grid of λ values
 - The grid you choose will be context dependent (play around with it!)
- 2) Compute cross-validation error (e.g. RMSE) for each
- 3) Choose the smallest one.

λ vs RMSE?

λ vs RMSE? A zoom

```
library(caret)
set.seed(100)
hbo <- read.csv("https://raw.githubusercontent</pre>
lambda seq <- c(0,10^s) = (-3, 1, length = 100)
ridge <- train(logins ~ . - unsubscribe - id,</pre>
            data = train.data,
            method = "glmnet",
            preProcess = "scale",
            trControl = trainControl("cv", num
            tuneGrid = expand.grid(alpha = 0,
                          lambda = lambda seq)
cv lambda <- data.frame(lambda = ridge$results)</pre>
                         rmse = ridge$results$RI
```

• We will be using the caret package

```
library(caret)
set.seed(100)
hbo <- read.csv("https://raw.githubusercontent</pre>
lambda seq <- c(0,10^s) = (-3, 1, length = 100)
ridge <- train(logins ~ . - unsubscribe - id,</pre>
            data = train.data.
            method = "glmnet",
            preProcess = "scale",
            trControl = trainControl("cv", num
            tuneGrid = expand.grid(alpha = 0,
                          lambda = lambda seq)
cv lambda <- data.frame(lambda = ridge$results)</pre>
                         rmse = ridge$results$RI
```

- We will be using the caret package
- We are doing cross-validation, so remember to set a seed!

```
library(caret)
set.seed(100)
hbo <- read.csv("https://raw.githubusercontent</pre>
lambda_seq <- c(0,10^seq(-3, 1, length = 100))
ridge <- train(logins ~ . - unsubscribe - id,
            data = train.data.
            method = "glmnet",
            preProcess = "scale",
            trControl = trainControl("cv", num
            tuneGrid = expand.grid(alpha = 0,
                          lambda = lambda seq)
cv lambda <- data.frame(lambda = ridge$results)</pre>
                         rmse = ridge$results$RI
```

- We will be using the caret package
- We are doing cross-validation, so remember to set a seed!
- You need to create a grid for the λ 's that will be tested

```
library(caret)
set.seed(100)
hbo <- read.csv("https://raw.githubusercontent</pre>
lambda_seq <- c(0,10^seq(-3, 1, length = 100))
ridge <- train(logins ~ . - unsubscribe - id,
            data = train.data.
            method = "glmnet",
            preProcess = "scale",
            trControl = trainControl("cv", num
            tuneGrid = expand.grid(alpha = 0,
                          lambda = lambda seq)
cv lambda <- data.frame(lambda = ridge$results)</pre>
                         rmse = ridge$results$RI
```

- We will be using the caret package
- We are doing cross-validation, so remember to set a seed!
- You need to create a grid for the λ 's that will be tested
- The function we will use is train: Same as before
 - method="glmnet" means that it will run an elastic net.
 - alpha=0 means is a ridge regression
 - o lambda = lambda_seq is not necessary (you can provide your own grid)

```
library(caret)
set.seed(100)
hbo <- read.csv("https://raw.githubusercontent</pre>
lambda_seq <- c(0,10^seq(-3, 1, length = 100))
ridge <- train(logins ~ . - unsubscribe - id,
            data = train.data.
            method = "glmnet",
            preProcess = "scale",
            trControl = trainControl("cv", num
            tuneGrid = expand.grid(alpha = 0,
                          lambda = lambda seq)
cv lambda <- data.frame(lambda = ridge$results)</pre>
                         rmse = ridge$results$R
```

- We will be using the caret package
- We are doing cross-validation, so remember to set a seed!
- You need to create a grid for the λ 's that will be tested
- The function we will use is train: Same as before
- Important objects in CV:
 - \circ results\$lambda: Vector of λ that was tested
 - \circ results\$RMSE: RMSE for each λ
 - \circ bestTune\$lambda: λ that minimizes the error term.

OLS regression:

Ridge regression:

```
coef(ridge$finalModel, ridge$bestTune$lambda)

## 5 x 1 sparse Matrix of class "dgCMatrix"
## s1
## (Intercept) 6.564243424
## female 0.002726465
## city 0.824387472
## age 0.046468790
## got -2.639308962

rmse(ridge, test.data)
```

```
## [1] 2.097452
```

Throwing a lasso

Lasso regression

• Very similar to ridge regression, except it changes the penalty term:

$$\min_{eta} \sum_{i=1}^n \underbrace{(y_i - \sum_{k=0}^p x_i eta_k)^2 + \lambda \cdot \sum_{k=1}^p |eta_k|}_{CLS}$$

• In our previous example:

$$\min_{eta} \sum_{i=1}^n (\operatorname{spend}_i - eta_0 - eta_1 \operatorname{female}_i - eta_2 \operatorname{freq}_i)^2 + \lambda \cdot (|eta_1| + |eta_2|)$$

• Lasso regression is also called l_1 regularization:

$$\left|\left|\beta\right|\right|_1 = \sum_{k=1}^p \left|\beta\right|$$

Ridge vs Lasso

Ridge

Final model will have p coefficients

Usually better with multicollinearity

Lasso

Can set coefficients = 0

Improves interpretability of model

Can be used for model selection

And how do we do Lasso in R?

```
library(caret)
set.seed(100)
hbo <- read.csv("https://raw.githubusercontent</pre>
lambda seq <- c(0,10^s) = (-3, 1, length = 100)
lasso <- train(logins ~ . - unsubscribe - id,</pre>
            method = "glmnet",
            preProcess = "scale",
            trControl = trainControl("cv", num
            tuneGrid = expand.grid(alpha = 1,
                          lambda = lambda seq)
cv lambda <- data.frame(lambda = lasso$results:</pre>
                         rmse = lasso$results$RI
```

Exactly the same!

• ... But change alpha=1!!

And how do we do Lasso in R?

Ridge regression:

```
coef(ridge$finalModel, ridge$bestTune$lambda)

## 5 x 1 sparse Matrix of class "dgCMatrix"
## s1
## (Intercept) 6.564243424
## female 0.002726465
## city 0.824387472
## age 0.046468790
## got -2.639308962

rmse(ridge, test.data)
```

```
## [1] 2.097452
```

Lasso regression:

[1] 2.09153

Your Turn

In-class Activity

- Go to https://sta235h.rocks/Week11 and read the exercise.
- Answer the questions in the activity and write them in your sheet.
- Submit your sheet at the end of the class.
- Important note: We will be doing a classification task (binary outcome). In this case, we <u>do not</u> use RMSE as a measure to compare outcomes, but <u>accuracy</u>.
 - Accuracy: How many observations (%) can I classify correctly?

Main takeway points

- You can shrink coefficients to introduce bias and decrease variance.
- Ridge and Lasso regression are similar:
 - Lasso can be used for model selection.
- Importance of understanding how to estimate the penalty coefficient.



References

- James, G. et al. (2021). "Introduction to Statistical Learning with Applications in R". Springer. Chapter 6.
- STDHA. (2018). "Penalized Regression Essentials: Ridge, Lasso & Elastic Net"