

Lecture 11

Variable selection

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Fall 2025

RStudio setup for this lecture

- Log into RStudio on your Amazon EC2 instance
 - Use AMI `FIN550-RStudio` with IAM role `BigDataEC2Role`

Enter this command via RStudio Terminal

```
aws s3 cp --recursive s3://bigdata-fin550-reif/lecture-11 ~/fin550/lecture-11
```

Selecting variables as predictors

- Including more variables in a model can improve predictive power
- If variables are uninformative, including them can cause overfitting
 - In this case, adding variables actually **reduces** predictive power
- Today: learn how to include variables in a model without overfitting

Selecting from p possible predictors

- Suppose you have p possible predictors: X_1, X_2, \dots, X_p
- Possible models:
 - $Y = \beta_0$
 - $Y = \beta_0 + \beta_1 X_1$
 - $Y = \beta_0 + \beta_5 X_5$
 - $Y = \beta_0 + \beta_3 X_3 + \beta_4 X_4 + \beta_{10} X_{10}$
 - ...
- In total, there are 2^p possible models

Methods for selecting variables

- Estimate all possible models and compare them
 - If p is large, however, then computationally impossible
- Alternative: employ "intelligent search methods"
 - Reduces computation time
 - But, it is possible that these will miss the best possible model
- Today, we learn about three popular methods:
 1. Best subset selection
 2. Forward stepwise selection
 3. Backward stepwise selection

Best subset selection

Try it: load seatbelt data

```
library(tidyverse)
library(boot)

# Load the data into a data frame
seatbelts <- as_tibble(datasets::Seatbelts)

# We will consider 4 possible predictors of DriversKilled:
#   X1 = law
#   X2 = PetrolPrice
#   X3 = kms
#   X4 = kms^2
seatbelts <- seatbelts %>%
  select(DriversKilled, law, PetrolPrice, kms) %>%
  mutate(kms2 = kms^2)
```

Monthly road casualties in Great Britain, 1969-1984

Variable name	Definition
DriversKilled	Car drivers killed (monthly)
law	0/1: was a seatbelt law in effect that month?
PetrolPrice	Price of petrol (gasoline)
kms	Distance driven
kms2	Distance driven squared

Try it: inspect the 1969-1984 monthly road casualties data

How many observations are there? Does that number make sense?

What is the range of the outcome variable, 'DriversKilled'?

Inspect the 1969-1984 monthly road casualties data

```
# How many observations are there? Does that make sense?
```

```
nrow(seatbelts)
```

```
# What is the range of the outcome variable, 'DriversKilled'?
```

```
summary(seatbelts$DriversKilled)
```

```
# [1] 192
```

```
#   Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
#   60.0   104.8   118.5   122.8   138.0   198.0
```

Best subset selection

- Consider all possible models
 - We have four predictors, so we must evaluate $2^4 = 16$ models
- Select the model with the lowest **cross-validated** mean-squared error
- Why not use (non-cross-validated) mean-squared error?
 - Because then we would always select models with more variables
 - That can lead to overfitting

Try it: create a formula object

```
# M0: constant only model (i.e. no predictors)  
f0_1 <- formula(DriversKilled ~ 1)  
f0_1  
  
# Following code is same as: `lm(DriversKilled ~ 1, data=seatbelts)`  
lm(f0_1, data = seatbelts)
```

```
# DriversKilled ~ 1  
#  
# Call:  
# lm(formula = f0_1, data = seatbelts)  
#  
# Coefficients:  
# (Intercept)  
#      122.8
```

Try it: define all possible formulas

M1: all models with 1 predictor

```
f1_1 <- formula(DriversKilled ~ law)
```

```
f1_2 <- formula(DriversKilled ~ PetrolPrice)
```

```
f1_3 <- formula(DriversKilled ~ kms)
```

```
f1_4 <- formula(DriversKilled ~ kms2)
```

M2: all models with 2 predictors

```
f2_1 <- formula(DriversKilled ~ law + PetrolPrice)
```

```
f2_2 <- formula(DriversKilled ~ law + kms)
```

```
f2_3 <- formula(DriversKilled ~ law + kms2)
```

```
f2_4 <- formula(DriversKilled ~ PetrolPrice + kms)
```

```
f2_5 <- formula(DriversKilled ~ PetrolPrice + kms2)
```

```
f2_6 <- formula(DriversKilled ~ kms + kms2)
```

Try it: define all possible formulas

M3: all models with 3 predictors

```
f3_1 <- formula(DriversKilled ~ PetrolPrice + kms + kms2)
```

```
f3_2 <- formula(DriversKilled ~ law + kms + kms2)
```

```
f3_3 <- formula(DriversKilled ~ law + PetrolPrice + kms2)
```

```
f3_4 <- formula(DriversKilled ~ law + PetrolPrice + kms)
```

M4: all models with 4 predictors

```
f4_1 <- formula(DriversKilled ~ law + PetrolPrice + kms + kms2)
```

Try it: create function to calculate cross-validated error

```
# Use glm() to estimate linear regression of formula f on the seatbelts dataset
cv_fun <- function(f) {
  glmfit <-
  cv.glm(data = seatbelts, glmfit)$delta[1]
}

# Run the function
cv_fun(f0_1)

cv_fun(f3_4)
```

Create function to calculate cross-validated error

Use glm() to estimate linear regression of formula f on the seatbelts dataset

```
cv_fun <- function(f) {  
  glmfit <- glm(f, data = seatbelts)  
  cv.glm(data = seatbelts, glmfit)$delta[1]  
}
```

Run the function

```
cv_fun(f0_1)
```

```
cv_fun(f3_4)
```

```
# [1] 647.5111
```

```
# [1] 533.2905
```


Try it: calculate MSE for all 16 formulas

Create a list of formulas

```
formulas <- list(f0_1,  
                f1_1, f1_2, f1_3, f1_4,  
                f2_1, f2_2, f2_3, f2_4, f2_5, f2_6,  
                f3_1, f3_2, f3_3, f3_4,  
                f4_1)
```

Create a vector for storing the LOOCV MSE for formulas

```
formulas_cv <- vector("numeric", length(formulas))
```

Use the function we created to calculate the LOOCV MSE for each formula

```
for (i in 1:length(formulas)) {  
  formulas_cv[[i]] <-  
}
```

Calculate MSE for all 16 formulas

Create a list of formulas

```
formulas <- list(f0_1,  
                f1_1, f1_2, f1_3, f1_4,  
                f2_1, f2_2, f2_3, f2_4, f2_5, f2_6,  
                f3_1, f3_2, f3_3, f3_4,  
                f4_1)
```

Create a vector for storing the LOOCV MSE for formulas

```
formulas_cv <- vector("numeric", length(formulas))
```

Use the function we created to calculate the LOOCV MSE for each formula

```
for (i in 1:length(formulas)) {  
  formulas_cv[[i]] <- cv_fun(formulas[[i]])  
}
```

Try it: identify model with the smallest MSE

```
best_model <- which.min(formulas_cv)
```

```
# Formula
```

```
formulas[[best_model]]
```

```
# MSE
```

```
formulas_cv[[best_model]]
```

```
# DriversKilled ~ law + PetrolPrice + kms2
```

```
# [1] 531.6267
```

lapply() is useful when applying a function to a list/vector

```
# Instead of writing this for loop:  
#   for (i in 1:length(formulas)) {  
#     formulas_cv[[i]] <- cv_fun(formulas[[i]])  
#   }  
# We can use lapply():  
formulas_cv2 <- lapply(formulas, cv_fun)  
  
best_model <- which.min(formulas_cv2)  
formulas[[best_model]]
```

```
# DriversKilled ~ law + PetrolPrice + kms2
```

apply() is equivalent to simplify(lapply())

```
# lapply() returns a list. Recall: formulas_cv2 <- lapply(formulas, cv_fun)  
typeof(formulas_cv2)
```

```
# [1] "list"
```

```
# sapply() works like lapply(), but it "simplifies" the output  
formulas_cv3 <- sapply(formulas, cv_fun)  
typeof(formulas_cv3)
```

```
# [1] "double"
```

```
all.equal(simplify(formulas_cv2), formulas_cv3)
```

```
# [1] TRUE
```

Technical note: parallel computing

- Parallel processing uses multiple CPU cores to run tasks simultaneously
- Great for tasks that repeat the same operation many times (e.g., model fitting)
- Not all code can be parallelized
 - If B depends on output from A, then can't run A and B simultaneously
- `lapply()` and `sapply()` run the same function on multiple inputs
 - R can run these in parallel with minimal edits to the code:

```
library(parallel)
formulas_cv2 <- mclapply(formulas, cv_fun, mc.cores = 4)
```

Problems with best subset selection

- Works well when the number of possible predictors is very small
- But, not feasible with large number of predictors

```
# Our model had 4 possible predictors:  $2^4 = 16$  models  
length(formulas)
```

```
# 40 predictors:  $2^{40} = \text{over 1 trillion models}$   
prettyNum(240, big.mark = ",")
```

```
# 300 predictors:  $2.04 \times 10^{90}$  models!  
format(2300, scientific=TRUE)
```

```
# [1] 16
```

```
# [1] "1,099,511,627,776"
```

```
# [1] "2.037036e+90"
```

Forward stepwise selection

Forward stepwise selection searches intelligently

- Start with all possible one-variable models
 - Find best predictor and keep it, e.g., $X = X_2$
- Now consider all two-variable models that include X_2
 - Find best predictor and keep it, e.g., $X = \{X_2, X_5\}$
- Repeat until you reach a model that includes all possible predictors
- Choose the model (1 variable, 2 variables, ... p variables) with lowest cross-validated MSE

Algorithm for forward stepwise selection

1. Let M_0 denote the null model with no predictors
2. Estimate all models with a single predictor
 - Let M_1 denote the 1-variable model with the lowest mean-squared error
3. Estimate all the models that add another predictor to the model M_1
 - Let M_2 denote the 2-variable model with the lowest mean-squared error
4. Repeat step 3 until you reach a model with all predictors (M_p)
5. Select the model M with the lowest **cross-validated** mean-squared error

Forward stepwise selection has pros and cons

- Instead of 2^p models, we estimate $1 + p(p + 1)/2$ models
 - When $p = 20$, this reduces number of models from 1,048,576 to 211
- However, it is possible that we may miss the best model
- Example: consider model with $p = 3$ predictors
 - Suppose best possible 1-variable model includes X_1
 - Suppose best possible 2-variable model includes X_2 and X_3
 - Then forward stepwise selection will fail to find this 2-variable model

Seatbelts example: M0 and M1

```
# Initialize vector and list to store MSE and formula for each model M (M0-M4)
```

```
forward_cv <- vector("numeric", 5)
```

```
forward_formulas <- vector(mode = "list", 5)
```

```
# M0: no predictors
```

```
forward_cv[1] <- cv_fun(f0_1)
```

```
forward_formulas[[1]] <- f0_1
```

```
# M1: consider all 1-variable models
```

```
M_formulas <- list(f1_1, f1_2, f1_3, f1_4)
```

```
cv <- sapply(M_formulas, cv_fun)
```

```
forward_cv[2] <- min(cv) # Store smallest MSE for M1
```

```
forward_formulas[[2]] <- M_formulas[[which.min(cv)]] # Store best model for M1
```

```
forward_formulas[[2]] # Display the 1-variable model with the lowest MSE
```

```
# DriversKilled ~ PetrolPrice
```

Seatbelts example: M2

```
# M2: consider all 2-variable models that include PetrolPrice
```

```
M_formulas <- list(f2_1, f2_4, f2_5)
```

```
print(as.character(M_formulas))
```

```
cat("\n")
```

```
cv <- sapply(M_formulas, cv_fun)
```

```
forward_cv[3] <- min(cv)
```

```
forward_formulas[[3]] <- M_formulas[[which.min(cv)]]
```

```
forward_formulas[[3]] # Display the 2-variable model with the lowest MSE
```

```
# [1] "DriversKilled ~ law + PetrolPrice"  "DriversKilled ~ PetrolPrice + kms"
```

```
# [3] "DriversKilled ~ PetrolPrice + kms2"
```

```
#
```

```
# DriversKilled ~ PetrolPrice + kms2
```

Seatbelts example: M3 and M4

```
# M3: consider all 3-variable models that include PetrolPrice and kms2
```

```
M_formulas <- list(f3_1, f3_3)
```

```
print(as.character(M_formulas))
```

```
cat("\n")
```

```
cv <- sapply(M_formulas, cv_fun)
```

```
forward_cv[4] <- min(cv)
```

```
forward_formulas[[4]] <- M_formulas[[which.min(cv)]]
```

```
forward_formulas[[4]] # Display the 3-variable model with the lowest MSE
```

```
# M4: all predictors (4-variable model)
```

```
forward_cv[5] <- cv_fun(f4_1)
```

```
forward_formulas[[5]] <- f4_1
```

```
# [1] "DriversKilled ~ PetrolPrice + kms + kms2"
```

```
# [2] "DriversKilled ~ law + PetrolPrice + kms2"
```

```
#
```

```
# DriversKilled ~ law + PetrolPrice + kms2
```

Select the model with the lowest MSE

```
# Display MSE and formulas for models M0-M4
forward_cv
cat("\n")
print(as.character(forward_formulas))
cat("\n")

# Display model with the lowest cross-validated mean-squared error
forward_formulas[[which.min(forward_cv)]]

# [1] 647.5111 556.7901 536.4993 531.6267 536.0520
#
# [1] "DriversKilled ~ 1"
# [2] "DriversKilled ~ PetrolPrice"
# [3] "DriversKilled ~ PetrolPrice + kms2"
# [4] "DriversKilled ~ law + PetrolPrice + kms2"
# [5] "DriversKilled ~ law + PetrolPrice + kms + kms2"
#
# DriversKilled ~ law + PetrolPrice + kms2
```

Backward stepwise selection

Backward stepwise selection is similar to forward

- Instead of starting with a model with 0 predictors, start with all predictors
- Incrementally remove the least predictive variable, one at a time
- Like forward stepwise selection, this reduces model count from 2^p to $1 + p(p + 1)/2$
- Lab assignment: implement backward stepwise selection

Algorithm for backward stepwise selection

1. Let M_p denote the full model with all predictors
2. Estimate all models that remove a single predictor from M_p
 - Let M_{p-1} denote the $p - 1$ variable model with the lowest mean-squared error
3. Estimate all the models that remove another predictor from the model M_{p-1}
 - Let M_{p-2} denote the $p - 2$ variable model with the lowest mean-squared error
4. Repeat step 3 until you reach a model with no predictors (M_0)
5. Select the model M with the lowest **cross-validated** mean-squared error

Comparing the different selection models

- Best subset selection will always find the best model
- However, these three selection models usually yield similar results
- Special case: number of variables p exceeds number of observations n
 - Can still use forward stepwise selection, but not backward stepwise selection
- Shrinkage methods such as Lasso (next lecture) can also be used for variable selection

Summary

- Three variable selection methods:
 1. Best subset selection
 2. Forward stepwise selection
 3. Backward stepwise selection
- When number of predictors exceeds about 10, need intelligent search methods
- Use cross-validated MSE when comparing models with different numbers of variables
- Lab-11 due Sunday at 11:59pm
- Make sure to **stop your instance** when you are done working