Stat 260, Lecture 12, Iteration

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Load packages and datasets

library(tidyverse)

Reading

- ▶ Iteration chapter of R for Data Science, chapter 17 of printed book and chapter 21 online.
- purrr cheatsheet [https: //github.com/rstudio/cheatsheets/raw/master/purrr.pdf]

Iterating over a vector

- ► For loops allow iteration.
- ➤ A common scenario for iteration is that our data is in a vector (list), and we want to perform the same operation on each element.
- Such iteration is so common that special tools have been developed with the aim of reducing the amount of code (and therefore errors) required for common iterative tasks.
 - ► Tools in base R include the apply() family of functions.
 - A tidyverse package called purrr includes more.

Example data

➤ To illustrate iteration we simulate data and fit four regression models.

```
set.seed(42)
n <- 100
x1 <- rnorm(n); x2<-rnorm(n)
y1 <- x1 + rnorm(n,sd=.5); y2 <- x1+x2+rnorm(n,sd=.5)
y3 <- x2 + rnorm(n,sd=.5); y4 <- rnorm(n,sd=.5)
rr <- list(fit1 = lm(y1 ~ x1+x2),
  fit2 = lm(y2 ~ x1+x2),
  fit3 = lm(y3 ~ x1+x2),
  fit4 = lm(y4 ~ x1+x2))
coef(rr$fit1)</pre>
```

```
## (Intercept) x1 x2
## 0.0008831357 0.9281453769 0.0426465892
```

Exercise

➤ The elements of the list rr from last slide are lm objects. The function coef() is generic. Assign class "Im_vec" to rr and write a coef() method for objects of this class. Hint: Your function could encapsulate a for() loop like the following.

```
for(i in seq_along(rr)) { # safer than 1:length(rr)
  print(coef(rr[[i]]))
}
```

```
##
    (Intercept)
                          x1
                                        x2
   0.0008831357 0.9281453769 0.0426465892
   (Intercept)
                        x1
                                     x2
    0.01572372 1.03114836 1.00306653
##
   (Intercept)
                        x1
                                     x2
## -0.06641184
                0.04316514 0.93035180
##
    (Intercept)
                          x1
                                        x2.
## -0.008394232 -0.018428268 -0.116309416
```

Extracting the regression coefficient for x1

Using a for() loop, we initialize an object to hold the output, loop along a sequence of values for an index variable, and execute the body for each value of the index variable.

```
betahat <- vector("double",length(rr))
for(i in seq_along(rr)) { # safer than 1:length(rr)
  betahat[i] <- coef(rr[[i]])["x1"]
}
betahat</pre>
```

```
## [1] 0.92814538 1.03114836 0.04316514 -0.01842827
```

Looping over elements of a set

-0.01842827

- The index set in the for() loop can be general.
 - We might use this generality to loop over named components of a list

```
fits <- paste0("fit",1:4)
for(ff in fits) {
  print(coef(rr[[ff]])["x1"])
##
          x1
## 0.9281454
##
         v1
## 1.031148
##
           x1
## 0.04316514
##
            x1
```

Looping over a set makes it harder to save the results, though.

Avoid growing vectors incrementally

```
means <- seq.int(1000)
set.seed(123)
system.time({
  output <- double()
  for (i in seq_along(means)) {
    n <- sample(100, 1)
    output <- c(output, rnorm(n, means[[i]]))
}
}
## user system elapsed
## 0.153 0.102 0.305</pre>
```

```
system.time({
  out <- vector("list", length(means))
  for (i in seq_along(means)) {
    n <- sample(100, 1)
    out[[i]] <- rnorm(n, means[[i]])
  }
  out <- unlist(out)
})

## user system elapsed
## 0.024 0.002 0.025</pre>
```

bind_cols() and bind_rows()

```
out <- vector("list", length(means))
n <- 100
for (i in seq_along(means)) {
  out[[i]] <- rnorm(n, means[[i]])
}
out <- bind_cols(out)
out <- vector("list", length(means))
for (i in seq_along(means)) {
  out[[i]] <- tibble(y=rnorm(n, means[[i]]),x=rnorm(n))
}
out <- bind_rows(out)</pre>
```

The body of a loop can be a small part of the code

- ▶ In our examples, most of the code is for setting up the output and looping, with very little to do with the body.
- ➤ To illustrate, consider a small change: instead of the estimated coefficient of x1 we wanted the estimated coefficient of x2:

```
betahat <- vector("double",length(rr))
for(i in seq_along(rr)) { # safter than 1:length(rr)
  betahat[i] <- coef(rr[[i]])["x2"]
}
betahat</pre>
```

```
## [1] 0.04264659 1.00306653 0.93035180 -0.11630942
```

Exercise

Write a for() loop to find the mode() of each column in nycflights13::flights

Using lapply()

► The intent of lapply() is to take care of the output and the loop, allowing us to focus on the body.

```
b1fun <- function(fit) { coef(fit)["x1"] } # body
bfun <- function(fit,cc) { coef(fit)[cc] } # body
lapply(rr,b1fun) # or sapply(rr,b1fun) or unlist(lapply(rr,b1fun))
## $fit1
##
          x1
## 0.9281454
##
## $fit2
##
         x1
## 1.031148
##
## $fit3
##
           x1
## 0.04316514
##
## $fit4
##
            x1
## -0.01842827
lapply(rr,bfun,"x1")
```

\$fit1 15/24

Exercise

▶ Re-write your coef() method for objects of class lm_vec to use lapply().

Iterating with the map() functions from purrr

- ► The purrr package provides a family of functions map(), map_dbl(), etc. that do the same thing as lapply() but work better with other tidyverse functions.
 - map() returns a list, like lapply().
 - map_dbl() returns a double vector, etc.

```
library(purr)
map_dbl(rr,b1fun) # or rr %>% map_dbl(b1fun)

## fit1 fit2 fit3 fit4

## 0.92814538 1.03114836 0.04316514 -0.01842827

# map_dbl(rr,bfun,"x1")
```

Exercises

- Use map_chr() to return the mode() of each column of the nycflights13::flights tibble.
- ▶ Use map() to return the summary() of each column of the nycflights13::flights tibble.

Pipes and map() functions

- Suppose we want to record a model summary returned by the summary() function.
 - summary() applied to an lm() object it computes regression summaries like standard errors and model R².

```
rr %>%
  map(summary) %>%
  map_dbl(function(ss) { ss$r.squared })
##
         fit1
                    fit2
                               fit3
                                          fit4
```

- Notice that we can define a function on-the-fly in the call to a map() function.
- map() functions have a short-cut for function definitions.

```
rr %>%
  map(summary) %>%
  map_dbl(~.$r.squared) # or map_dbl("r.squared")
```

```
## fit1 fit2 fit3 fit4
## 0.78845184 0.91430933 0.73684218 0.04087594
```

▶ In ~. read ~ as "define a function" and . as "argument to the function"

Exercise

Write a call to map_dbl() that does the same thing as map_dbl(rr,b1fun), but define the function on the fly, as in the previous slide. You can use multiple calls to map() functions.

Detour: The apply family of functions in R

► The "original" apply is apply(), which can be used to apply a function to rows or columns of a matrix.

```
mat <- matrix(1:6,ncol=2,nrow=3)</pre>
mat.
##
        [,1] [,2]
## [1.]
## [2,] 2 5
## [3,] 3 6
apply(mat,1,sum) # row-wise sums; rowSums() is faster
## [1] 5 7 9
apply(mat,2,sum) # column-wise; colSums() is faster
## [1] 6 15
```

Detour, cont.

sapply() takes the output of lapply() and simplifies to a vector or matrix.

Detour, cont.

- ▶ Other apply-like functions vapply(), mapply(), tapply(), ...
- ► I don't use these.
 - See their respective help pages for information.