

Introduction to R

Session 03: Basics of programming

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- Basic idea: you have a vector of values, and you have an "iterator" variable
- ➤ You go through the vector, setting the iterator variable to each value one at a time
- ► Then you run a chunk of that code with the iterator variable set

```
for (iteratorvariable in vector) {
  code chunk
}
```



```
data(mtcars)
abovemed <- mtcars %>% filter(cyl >= median(cyl))
belowmed <- mtcars %>% filter(cyl < median(cyl))
for (i in c('mpg', 'disp', 'hp', 'wt')) {
   print(mean(abovemed[[i]])-mean(belowmed[[i]]))
}</pre>
```



- ► Sometimes, you want R to do something only in some cases
- The basic syntax to do this is an "if" statement

```
if (test_expression) {
statement
}
```

- ► The "test_expression" usually depends on logical operators
 - ▶ '&' is AND
 - '|' is OR
 - ▶ To check equality use '==', not '='
 - > '>=' is greater than OR equal to, similarly for '<='



- Sometimes is useful to "break" (stop) a loop, or skip an iteration
- ▶ Do this with "break" and "next"
- ▶ Often "break" and "next" are nested within an "if" statement

```
#printing odd numbers
m=20
for (k in 1:m){
   if (!k %% 2)
       next
   print(k)
}
```





Example: finding prime numbers



```
for(num in 1:100){
       # Program to check if the input number is prime or not
       flag = 0
       # prime numbers are greater than 1
       if(num > 1) {
6
         # check for factors
         flag = 1
8
         for(i in 2:(num-1)) {
9
            if ((\text{num } \%\% \text{ i}) \stackrel{\checkmark}{=} 0) {
              flag = 0 #if number is divisible, then not prime
10
11
              break #and we can break the loop
12
13
14
                         flag = 1
15
       if (num = 2)
16
       if(flag == 1) {
17
          print(paste(num, "is-a-prime-number"))
18
       } else
          print(paste(num, "is - not - a - prime - number"))
19
20
21
```

Functions



- Functions are used when you have to repeat the same operation multiple times
- Avoids coding mistakes (and if there is one you only need to fix it once)
- ► Let's you break code into simpler parts which become easy to maintain and understand
- It's pretty straightforward to create your own function in R programming.

```
func_name <- function (argument) {
statement
return(whatyouwant)
}</pre>
```

Example: finding prime numbers



```
# Identifying prime number
   prime_num<-function(num){</pre>
2
     # Program to check if the input number is prime or not
     flag = 0
     # prime numbers are greater than 1
     if (num > 1) {
6
       # check for factors
7
       flag = 1
       for(i in 2:(num-1)) {
          if ((num \% i) = 0) {
            flag = 0 #if number is divisible, then not prime
11
            break #and we can break the loop
14
15
      if(num == 2) flag = 1
16
      return (flag)
17
18
```

Example: finding prime numbers



```
prime_num(2)
prime_num(3)
prime_num(4)
prime_num(6131)
prime_num(4684561123)
```

Functions



Functions can have multiple arguments.

► Functions can have default values for arguments.

► Functions can output vectors, list, or any other object.

Example: power function



```
pow <- function(x, y = 2) {
    result1 <- x^y
    return(result1)
}
pow(3)
pow(3,3)</pre>
```

Some basic guiding principles



- ▶ Automate everything that can be automated.
- Design your workflow so that when you add new data or tweak something you don't need to manually recreate anything
- Avoid copying and pasting figures/tables from one software into another
 - e.g., impossible to automate: copying a graph from R into Word
- ▶ General workflow: raw data \rightarrow clean data \rightarrow analysis \rightarrow (pretty) table or figure



Don't forget to look at the data



▶ Use the R studio browser or excel

 Scroll through the variables and accompany yourself with what you've got visually

Look for missing observations



- Check the size of your dataset using dim
- ► Check the structure of your data using str
- Check for missing variables
 - Sometimes the data has -99 for missing
 - Others is truly missing (i.e., NA)
 - Others is an empty string

Always check your merges (combining data sets)



- During a stage of arranging datasets, you will likely merge
- ► Make sure you count before and after you merge so you can figure out what went wrong, if anything
- Also make sure you understand what type of merge you are using/need
 - many to one
 - one to many
 - one to one
 - ▶ many to many (please no!!!)

Good practices



- Beyond checking the data, here are a few things that will prevent errors
 - 1. Organized subdirectories
 - 2. Automation
 - 3. Naming conventions
 - 4. Version control

Naming conventions



- ► Things to keep in mind when naming:
 - 1. variables,
 - 2. datasets
 - 3. scripts
- ► Avoid
 - meaningless words (e.g., 1mb2)
 - dating (e.g., temp05012020), except to save old version in the archive
 - numbering (e.g., outcome25)
 - ▶ any of these will confuse your future self

Naming conventions for variables



- ► Variables should be readable to a stranger
 - Say that you want to create the product of two variables. Use an underscore and mash the two together
 - price_mpg <- price * mpg</p>
- Name the variable exactly what it is when possible
 - bmi <- weight / (height*height * 703)</p>

Headers



- Always start your programming scripts with a header
- ▶ The header is a big comment on the code and says:
 - ► The purpose of the script
 - ► The author(s)
 - The date the script was created
 - ► The last date in which it was updated/modified (and by whom)

Automating Tables and Figures (more on visualization below)

► Goal is to make "beautiful tables/figures" that are never edited post-production

► Tables/figures should be readable on their own (with the table/figure notes)

► Large fixed costs learning commands like "stargazer" in R or "estout" in R

Eliminate junk



- Maximize the data-to-ink ratio by using as little ink as possible to show your data
- ► Remove non data ink, e.g., extra gridlines
- ► Remove redundant data
- Remove indicators you don't need
- Display only as many decimal places as are relevant

Choose the type of visualization based on the information you want to convey

- Use a table only if you can't use a figure. Usually when
 - You need to show exact numerical values
 - You want to allow for multiple localized comparisons
 - You have relatively few numbers to show
- Avoid pie charts
 - ▶ It is difficult for the audience to visually distinguish the size of each wedge
 - A good alternative for showing shares of a whole is a stacked bar chart



Other random tips



- ► Make the font larger than the Stata/R preset
- Label your axis
- ► The Y-axis tickmarks ideally should be horizontal (R puts them sideways)
- Use legends if you have multiple lines
- Avoid having two-axis
- Avoid pie-charts

Example



- Add x-axis tick marks
- ▶ We only want people aged 5-18
- Make the y-axis in percent terms
- Use a more appealing color scheme
- ► The y-axis is misleading, lets make it go to 100
- Add a legend to explain what the colors mean
- Make the numbers in the y-axis easier to read and remove legend box