Statistical Computing with R Masters in Data Science 503 (S4) First Batch, SMS, TU, 2021

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Review Preview

- Histogram
 - Correct use of central tendency and dispersion parameter
- Boxplot
 - Summary of skewed variable/s; comparison by groups
- Scatterplot
 - Linear or non-linear correlation and regression analysis
- Vector in R
 - Vector operations in R
- Function in R
 - Built-in function
 - User defined function

Vectorized operations in R:

 A key difference between R and many other languages is a topic known as vectorization

 We can write "total" function in R using "loop" but it will be much faster to use built-in "sum" function of R to get the "total"

 "Sum" is already coded in C to work with vector of numbers in R and the loop written in C is hidden from us in R

Many other built-in functions also works in this way in R!

Vector operation in R:

- a <- 1:10 #seq(1,10) is equivalent to this
- b <- 1:10 #c(1,2,3,4,5,6,7,8,9,10) is same as this
- How to add a and b vectors (variables)?
- Do we need to write functions for this?
- What will happen if we type: a + b in R console?
- [1] 2 4 6 8 10 12 14 16 18 20 #Matrix addition?

Quick Think!

 What will happen when you type these two codes in R/R studio and run:

• y <- seq(1, 10, length.out = 5)

• (y < - seq(1, 10, length.out = 5))

Vector operation in R: Continued ...

- a <- 1:10 #Vector length = 10
- b <- 5 #Vector length = 1
- What will happen if we type: a * b in R console?
- [1] 5 10 15 20 25 30 35 40 45 50
- Why? # R does not have scalars!

Exercise: Do this in R or R Studio!

• Define:

- a = [1, 2, 3, 4, 5, 6, 7, 8, 9, 10]
- b = [5]
- c = [2, 3, 4, 5, 6, 7, 8, 9, 10, 11]

- #Check class of 'a'
- #Check class of 'b'
- #Check class of 'c'

• Define:

- ac = cbind(a,c)
- acb = ac * b

#Check class of 'ac'

#Check class of 'acb'

• Get:

- d = arithmetic mean by cases (row means)
- #Hint: Use "apply" function
- acbd = add new variable d in the acb matrix
- #Check class of 'acbd' object
- Summarize all the variables of 'acbd' object and interpret the results carefully!

Vector operation in R: Continued ...

- a <- 1:10 #Vector length = 10
 b <- 1:5 #Vector length = 5
- What will happen if we type: a + b in R console?
- [1] 2 4 6 8 10 7 9 11 13 15 #Matrix addition!
- Why? #1+1, 2+2, 3+3, 4+4, 5+5, 6+1, 7+2, 8+3, 9+4, 10+5
- #Vector recycling is done when we add vectors with unequal lengths, take a note of this!

Vector operation in R: Continued ...

- a <- 1:10 #Vector length = 10
- b <- 1:7 #Vector length = 7

- What will happen if we type: a + b in R console?
- What does the "error" message tells you?

What is the key message?

Function in R: Continued ...

```
best_practice <- c("Let", "the", "computer", "do", "the", "work")</pre>
print_words <- function(sentence) {</pre>
        print(sentence[1])
                                                    What is wrong with this approach?
        print(sentence[2])
        print(sentence[3])
        print(sentence[4])
        print(sentence[5])
        print(sentence[6])
print_words(best_practice)
                                   # [1] "Let" [1] "the" [1] "computer" [1] "do" [1] "the" [1] "work"
                                   # [1] "Let" [1] "the" [1] "computer" [1] "do" [1] "the" [1] "NA"
print_words(best_practice[-6])
                                             #[1] "Let" "the" "computer" "do" "the"
best_practice[-6]
```

Can we improve it in R? We can use functions with "for" loop in R!

```
print words <- function(sentence) {</pre>
                                                        for (variable in collection) {
   for (word in sentence) {
                                                           do things with variable
   print(word)
                                         "for" loop
print words(best_practice)
[1] "Let" [1] "the" [1] "computer" [1] "do" [1] "the" [1] "work"
print words(best practice[-6])
[1] "Let" [1] "the" [1] "computure" [1] "do" [1] "the"
```

"for" and "while" loops can be very slow in R!

What to do?

R:

Loops in R will not be slow if we:

Don't use a loop when a vectorized alternative exists

 Don't grow objects (via c, cbind, etc) during the loop – R has to create new object and copy across the information just to add new element or row/column

Allocate an object to hold the result and fill it during the loop

Can we do even better in R? Alternative to "loop" in R??

While working with data.frame in R:

- It is better to use:
 - apply
 - lapply
 - sapply
 - vapply

We will discuss this in detail while doing breakdown analysis session in R later!

- functions instead of "for loop" to run the script much faster in R!
- Same applies to the "while loop" too!

R script:

• It is a collection of codes and can be saved with an .R extension

It contains #comments

• It can be used to create "markdown" file using knitr package

 R markdown files are ready to deploy files as an online or offline reports using bookdown package e.g. https://bookdown.org/ Use of R Markdown (*.rmd): Sharing!

What is R script? How to use it??

```
p wp_websrapping_in_R.R ×
      Run Source - =
      library(rvest)
   1
       library(dplyr)
   2
   3
   4
       wiki_link = "https://en.wikipedia.org/wiki/COVID-19_pandemic_in_Nepal"
   5
       wiki_page = read_html(wiki_link)
   6
   7
       covid_table = wiki_page %>% html_nodes("table") %>% .[14] %>%
   8
         html_table() %>% . [[1]]
   9
  10
       names(covid_table) = paste(names(covid_table), covid_table[1, ], sep = "_")
  11
       covid_table = covid_table[-1, ]
  12
  13
       summary(covid_table)
  14
  15
       colnames(covid_table)
  16
       names(covid_table) [names(covid_table) == "Date_Date"] = "Date"
  17
       names(covid_table) \[ \text{rames(covid_table)} == "Confirmed cases_Total" \] = "Confirmed_Cases_Total" \]
       names(covid_table) \[ \text{rames(covid_table}) == "Confirmed cases_New" \] = "Confirmed_Cases_New" \]
  18
       names(covid_table) \[ \text{rames(covid_table)} == "Confirmed cases_Active" \] = "Confirmed_Cases_\]
  19
       names(covid_table)[names(covid_table) == "RT-PCR tests_Total"] = "PCR_Total"
  20
  21
       names(covid_table) \[ \text{rames(covid_table)} == "RT-PCR tests_New" \] = "PCR_New" \]
  22
       names(covid_table) \[ \text{names(covid_table)} == "TPR_TPR" \] = "TPR" \]
  23
       names(covid_table)[names(covid_table) == "RR_RR"] = "RR"
       names(covid_table)[names(covid_table) == "CFR_CFR"] = "CFR"
  24
       names(covid_table)[names(covid_table) == "Ref._Ref."] = "Ref"
  25
       colnames(covid_table)
  26
 1:1
       (Top Level) $
                                                                                         R Script $
```

R script: Why %>% was used?

 This is called "pipe" operator and described in chapter 14 of the course book provided to you

 Pipes are a powerful tool for clearly expressing a sequence of multiple operations

• The pipe, %>%, comes from magrittr package

 The pipe helps you to write code in a way that is easier to read and understand! Example: Understand the run in R/R Studio:

```
rnorm(100) %>%
matrix(ncol = 2) \%>\%
plot() %>%
str()
```

Condition: if and else

```
if (condition) {
      #code executed when condition is TRUE
} else {
      #code executed when condition is FALSE
}
```

Can you think of an example?

What will be the output?

```
if (y < 20) {
    x <- "Too low"
} else {
    x <- "Too high"
}</pre>
```

Multiple conditions:

```
if (this) {
      # do that
} else if (that) {
      # do something else
} else {
      # remaining
```

Multiple Conditions: If, else if, else

```
if (temp <= 0) {
"freezing"
} else if (temp <= 10) {
"cold"
} else if (temp <= 20) {
"cool"
} else if (temp <= 30) {
"warm"
} else {
"hot"
```

Question/queries?

Thank you!

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