## Statistical Computing with R Masters in Data Science 503 (S5) Fourth Batch, SMS, TU, 2025

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#### Review Preview

Basics of R

Basics of coding in R

 Chapter from "R for Everyone" book  Chapter from "Hands-on Programming with R" book

We discussed it in the last class

We will discuss this in today's class

#### (Variable) Naming convention is R?

- The article "The State of Naming Conventions in R" suggest to use:
  - allowercase e.g. adjustcolor
  - period.separated e.g. plot.new
  - underscore\_separated e.g. numeric\_version
  - lowerCamelCase e.g. addTaskCallback
  - UpperCamelCase e.g. SignatureMethod

#### This link suggests to use "underscore\_separated"

https://www.r-bloggers.com/2014/07/consistent-naming-conventions-in-r/

- It argues that:
  - allowercase names are **difficult to read**, especially for non-native readers.
  - period.separated names are confusing for users of <u>Python</u> and other languages in which dots are meaningful.
  - UpperCamelBack is ugly and requires excessive use of the shift button.

#### Functions in R: Built-in functions

- round()
  - round(3.1415)
  - 3
- factorial()
  - factorial(3)
  - 6
  - $3! = 3 \times 2 \times 1$
- mean()
  - mean(1:6)
  - =(1+2+3+4+5+6)/6=3.5

```
round()
round(3.1415, digits = 2)
3.14
```

```
factorial()
factorial(2*3)
720
6! = 6 x 5 x 4 x 3 x 2 x 1
```

```
mean()
mean(c(1:30)
15.5
```

# "Sample" function: Random sampling without or with replacement in R

```
die <- 1:6
```

- sample(x = die, size = 1)
- sample(x = die, size = 1)
- sample (x = die, size = 1, replace=TRUE)
- sample(x = die, size = 2)
- sample(x die, size = 2)
- sample(x = die, size = 2, replace=TRUE)

# "Sample" function to split a datafile into train and test datasets

# Make sure to have "iris.csv" datafile in the working directory and use read.csv to import it in R Studio:

- read.csv("iris.csv")
- We can do the 70:30 random split of iris data frame as follow:
  - set.seed(123)
  - tt.sample <- sample(c(TRUE, FALSE), nrow(iris), replace=T, prob=c(0.7,0.3))
  - train <- iris[tt.sample, ]</li>
  - test <- iris[!tt.sample, ]</li>

This is very handy as main data is frequently divided into the Training and Testing datasets in Data Science models! Why?

#### User-defined function in R:

- my\_function <- function() {}</li>
- Where,
- my\_function = name of the function e.g. roll (roll the die)

• function() = telling R that it is a user-defined function

- { = We need to start our code after this braces
- } = We need to close our codes before this braces

#### User-defined function 1: roll()

```
roll <- function() {
         die <- 1:6
         dice <- sample(die, size = 2, replace = TRUE)
         sum(dice)
}</pre>
```

First roll: roll()

Second roll: roll()

Third roll: roll()

#### Function creation in R: HOPR, Chapter 1

- 1. **The name**. A user can run the function by typing the name followed by parentheses, e.g., roll2().
- 3. **The arguments**. A user can supply values for these variables, which appear in the body of the function.

The body. R will run this code whenever a user calls the function. roll2 <- function(bones = 1:6) {
 dice <- sample(bones, size = 2,
 replace = TRUE)
 sum(dice)</pre>

The default values.
 Optional values that R can use for the arguments if a user

does not supply a value.

The last line of code.
 The function will return the result of the last line.

Figure 1-6. Every function in R has the same parts, and you can use function to create these parts.

#### User-defined function 2: roll2()

```
roll2 <- function(dice = 1:6) {
        dice <- sample(dice, size = 2, replace = TRUE)
        sum(dice)
}</pre>
```

First roll: roll2()

Second roll: roll2()

Third roll: roll2()

#### User-defined function 3: roll3(data?)

```
roll3 <- function(dice) {
         dice <- sample(dice, size = 2, replace = TRUE)
         sum(dice)
}</pre>
```

First roll: roll3(dice = 1:6)

Second roll: roll3(dice = 1:12)

Third roll: roll3(dice = 1:24) # Is this possible in two dice?

#### 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])
best practice[-6]
                                             #[1] "Let" "the" "computer" "do" "the"
```

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

```
print_words <- function(sentence) {</pre>
   for (word in sentence) {
                                                        for (variable in collection) {
   print(word)
                                         "for" loop
                                                           do things with variable
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"
```

https://swcarpentry.github.io/r-novice-inflammation/03-loops-R/

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

What to do?

#### 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 family of "apply" functions from base R:

- apply
- lapply
- sapply
- vapply

More here:

https://www.datacamp.com/t utorial/r-tutorial-apply-family

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!

#### 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?

```
#Checking values of y with x:
                                       #Will this work?
                                       check.y <- function(y) {</pre>
if (y < 20) {
 x <- "Too low"
                                       if (y < 20) {
                                       print("Too Low") } else {
} else {
 x <- "Too high"
                                       print("Two high")
#Can you get anything from this?
                                       check.y(10)
                                       check.y(30)
```

#### Creating binary variables with "ifelse"

#Will this work? #Will this work?

y <- 1:40

ifelse(y<20, "Too low", "Too high") ifelse(y<20, 1, 0)

It's a logical as:

ifelse(y<20, TRUE, FALSE) Good to make binary variables with numerical categories!

Good to make binary variables with text categories!

This one is preferred in DS!

#### Multiple conditions:

```
In a function:
if (this) {
        # do that
} else if (that) {
        # do something else
} else if (that) {
        # do something else
} else
# remaining
```

```
check.x <- function(x=1:99){
if (x<20){
print("Less than 20")} else{
if (x<40) {
print("20-39")} else{
if (x<100) {
print("41-99")}
}}}
• check.x(15)
check.x(30)
check.x(45)
Good to make categorical variables!
```

#### Multiple Conditions: combining "ifelse"

#### Will this work too?

```
x <- 1:99
```

x1 <- ifelse(x<20, 1,0) #Binary numbers

x2.1 <- ifelse(x<20, "<20", "20+") #Binary text

x2.2 ? For x between 20 and less than 40

x2.3 ? For x between 40 and less than 100

Now combine them in a single column with <20=1, 20-39=2 and 40-99=3 for x i.e. create categorical variable of x!

#### Will this work?

```
x3 <- ifelse(x<20,1,ifelse(x<40,2,3))
x3
table(x3)
```

```
#This code shows how Petal. Length
categories was created from Petal. Length
variable of iris data frame
iris <- within(iris, {
Petal.cat <- NA
Petal.cat[Petal.Length <1.6] <- "Small"
Petal.cat[Petal.Length >=1.6 &
Petal.Length<5.1] <- "Medium"
Petal.cat[Petal.Length >=5.1] <- "Large"
#The 1.6=Q1 and 5.1=Q3 were obtained
from the "summary" of the Petal.Lenght
variable i.e. summary(iris$Petal.Length)
Iris$Petal.cat
table(iris$Petal.cat)
```

#### Multiple Conditions: If, else if, else if, else if

```
#Make this function work!
if (temp <= 0) {
"freezing"}
else if (temp <= 10) {
"cold"}
else if (temp <= 20) {
"cool"}
else if (temp <= 30) {
"warm"}
else {
"hot"}
```

```
# What is missing?
# How to address it?
```

### Questions/queries?

## Thank you!

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