Programming

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Introduction to R and RStudio

Recap

Last week

- How to use R, RStudio, R-scripts and R-notebooks
- Data types (elements)
 - character, numeric, integer, logical, factor
- Data structures: composed of data types
 - vector, matrix, list, data.frame
- Subsetting data structures
- Reading files in different formats

This week

How to organize and automate your code:

First half

- Control-flow:
 - Choice: if-else statements
 - Loops: For loops
- Functions
- Environments

Second half

- Principles of tidy data and short comparison of base R and the tidyverse
- Inferential statistics: A primer of linear regression
- Best practices in R

Control-flow

New controls and functions

New control flow constructs

- Choice:
 - We often want to run some code *only if* some *condition* is true.
 - if(cond) {cons.expr} else {alt.expr}
- Loops:
 - We often want to repeat the execution of a piece of code many times.
 - for(var in seq) {expr}

Loops in R happen often under the hood. New functions:

- apply(): apply a function to margins of a matrix
- sapply(): apply a function to elements of a list, vector or matrix return
- lapply(): apply a function to elements of a list, list return

Control-flow (I): Choice

If statement

Operation of a if statement:

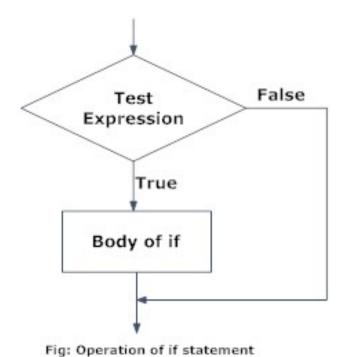
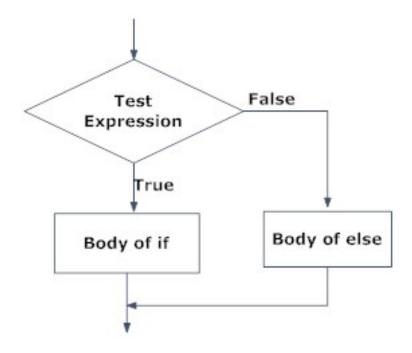


Figure 1: Source: datamentor.io

Code of an if statment:

```
value <- 3
if (value > 3) { #condition
  print("Value greater than 3") #conditional code
}
```

If-else statements



Operation of a if-else statement:

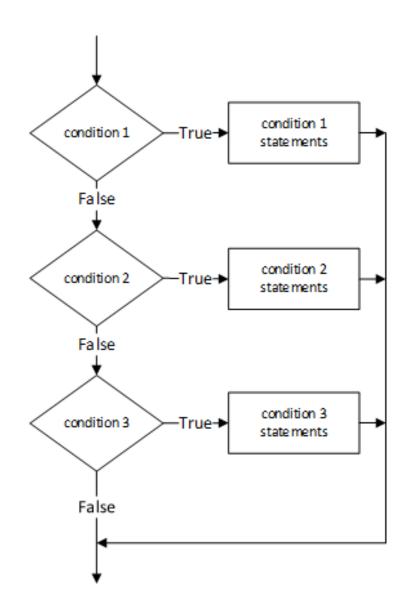
Fig: Operation of if...else statement

Code of an if-else statment:

```
value <- 3
if (value > 3) { #condition true?
  print("Value greater than 3") #code in if block
} else {
    print("Value lower or equal to one") #code in else block
}
```

[1] "Value lower or equal to one"

If-else statements



Operation of a if-else statement:

Code of an if-else statment:

```
value <- 3
if (value > 3) { #condition 1
  print("Value greater than 3") #condition 1 statements
} else if (value > 1) { #condition 2
  print("Value greater than 1") #condition 2 statements
} else if (value > 0) { #condition 3
  print("Value greater than 0") #condition 3 statements
}
```

[1] "Value greater than 1"

Subsetting consists of if-else statements

Remember our example from last time

The computer keeps the value of the element i of example_vector if the element i of the condition (example_vector>3) is true.

```
example_vector = c(1,2,3,4,5,6,7,8,9)
example_vector>3

## [1] FALSE FALSE TRUE TRUE TRUE TRUE TRUE TRUE
example_vector[example_vector>3]

## [1] 4 5 6 7 8 9
```

Control-flow (II): Loops

For loops

For loops are used when we want to perform some repetitive calculations.

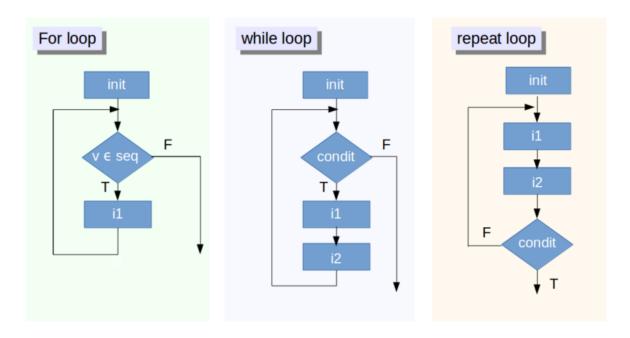


Figure 2: Source: datacamp.com

For-loops

```
# Let's print the numbers 1 to 6 one by one.
print(1)
## [1] 1
print(2)
## [1] 2
print(3)
## [1] 3
print(4)
## [1] 4
print(5)
## [1] 5
```

```
print(6)
## [1] 6
```

For-loops

For-loops allow us to automate this!

For each element of 1:6, print the element:

```
for (i in 1:6){
   print(i)
}

## [1] 1
## [1] 2
## [1] 3
## [1] 4
## [1] 5
## [1] 6
```

For-loops

You can use any variable name, i is a convention for counting/index.

```
for (some_var_name in 1:6){
   print(some_var_name)
}

## [1] 1
## [1] 2
## [1] 3
## [1] 4
## [1] 5
## [1] 6
```

Subsetting consists of for-loops and if-else statements

For each element i, keep the value of the element i of example_vector if the element i of the condition (example_vector>3) is true.

```
example_vector = c(1,2,3,4,5,6,7,8,9)
example_vector>3

## [1] FALSE FALSE TRUE TRUE TRUE TRUE TRUE TRUE
example_vector[example_vector>3]
```

```
## [1] 4 5 6 7 8 9
```

For-loops

Often you don't want to iterate over a range, but over an object

```
for (element in c("Amsterdam","Rotterdam","Eindhoven")){
  if (element == "Amsterdam"){
    print(paste(element,"Terrible football team.", sep=": "))
} else {
    print(paste(element,"Not the prettiest city, but at least their football team is okay.", sep=": "))
```

```
}
}
## [1] "Amsterdam: Terrible football team."
## [1] "Rotterdam: Not the prettiest city, but at least their football team is okay."
## [1] "Eindhoven: Not the prettiest city, but at least their football team is okay."
For-loops
Something a bit more useful
df <- data.frame("V1" = rnorm(5),</pre>
                 "V2" = rnorm(5, mean = 5, sd = 2),
                 "V3" = rnorm(5, mean = 6, sd = 1))
head(df)
##
              ۷1
                       ٧2
## 1 -1.59177940 4.316317 7.368464
## 2 0.76324704 6.760209 6.956894
```

```
## [1] "V1"
## [1] "V2"
```

print(col)

For-loops

[1] "V3"

Doing an operation on each column

3 -0.22899353 4.715755 5.895623 ## 4 -0.04584141 2.125445 5.913738 ## 5 0.90252283 4.809158 5.467448

for (col in names(df)) {

For-loops

Doing an operation on each row

```
for (row in 1:nrow(df)) {
  row_values = df[row, ]

  print(paste("Row ", row, ": ", sum(row_values>5), " values over 2", sep=""))
}
```

```
## [1] "Row 1: 1 values over 2"
## [1] "Row 2: 0 values over 2"
## [1] "Row 3: 1 values over 2"
## [1] "Row 4: 1 values over 2"
## [1] "Row 5: 2 values over 2"
```

While loops

Do something forever until a condition is (not) met

```
i = 0
while (i < 10) {
    i = i + 1
    print(i)
}

## [1] 1
## [1] 2
## [1] 3
## [1] 4
## [1] 5
## [1] 6
## [1] 7
## [1] 8
## [1] 9
## [1] 10</pre>
```

More info on loops: https://www.datamentor.io/r-programming/break-next/

The apply() family

apply()

The apply family is a group of very useful functions that allow you to easily execute a function of your choice over a list of objects, such as a list, a data.frame, or matrix.

We will look at three examples:

- apply
- sapply
- lapply

apply()

apply is used for matrices/dataframes. It applies a function to each *row* or *column*. It returns a vector or a matrix.

```
head(df, 1)

## V1 V2 V3

## 1 0.4826716 4.034959 5.709234

Apply it by row (MARGIN = 1):

apply(df, MARGIN = 1, mean)

## [1] 3.408955 3.142947 4.236908 1.738974 3.368189
```

Apply it by column (MARGIN = 2):

```
apply(df, MARGIN = 2, mean) #Identical to colMeans(df)

## V1 V2 V3

## -0.2649956 3.6483485 6.1542312
```

sapply()

sapply() is used on list-objects. It returns a vector or a matrix.

```
my.list \leftarrow list(A = c(4, 2, 1:3), B = "Hello.", C = TRUE) sapply(my.list, class)
```

```
## A B C
## "numeric" "character" "logical"
my.list <- list(A = c(4, 2, 1:3), B = c("hello", "Hello.", "Aa", "aa"), C = c(FALSE, TRUE))
sapply(my.list, range)</pre>
```

```
## A B C
## [1,] "1" "aa" "0"
## [2,] "4" "Hello." "1"
```

Why is each element a character string?

sapply()

Any data.frame is also a list, where each column is one list-element.

This means we can use sapply on data frames as well, which is often useful.

```
sapply(df, mean)
```

```
## V1 V2 V3
## -0.2649956 3.6483485 6.1542312
```

lapply()

lapply() is exactly the same as sapply(), but it returns a list instead of a vector.

```
lapply(df, class)
```

```
## $V1
## [1] "numeric"
##
## $V2
## [1] "numeric"
##
## $V3
## [1] "numeric"
```

Writing your own functions

What are functions?

Functions are reusable pieces of code that

- 1. take some standard input (e.g. a vector of numbers)
- 2. do some computation (e.g. calculate the mean)
- 3. return some standard output (e.g. one number with the mean)

We have been using a lot of functions: code of the form something() is usually a function.

```
mean(1:6)
## [1] 3.5
```

Our own function

We can make our own functions as follows:

```
squared <- function (x){
  x.square <- x * x
  return(x.square)
}</pre>
```

[1] 16

x, the input, is called the (formal) argument of the function. x.square is called the return value.

Our own function

If there is no return(), the last line is automatically returned, so we can also just write:

I do not recommend this.

```
squared <- function(x){
  x * x
}
squared(-2)</pre>
```

[1] 4

Our own function

We can also combine this with apply()

```
df
```

```
## V1 V2 V3

## 1 0.4826716 4.034959 5.709234

## 2 0.6331722 4.127596 4.668073

## 3 0.4748632 3.950741 8.285120

## 4 -2.0082158 1.110719 6.114420

## 5 -0.9074691 5.017728 5.994309

apply(df, 2, squared)
```

```
## V1 V2 V3

## [1,] 0.2329718 16.280896 32.59536

## [2,] 0.4009070 17.037048 21.79090

## [3,] 0.2254950 15.608353 68.64322

## [4,] 4.0329308 1.233696 37.38613

## [5,] 0.8235002 25.177597 35.93174
```

Default options in functions

• Default options for some arguments are provided in many functions.

• They allow us to provide an additional option, but if no choice is provided, we can choose for the user of the function.

```
is_contained <- function(str_1, str_2, print_input = TRUE){
   if (print_input){
      cat("Testing if", str_1, "contained in", str_2, "\n")
   }
   return(str_1 %in% str_2)
}

is_contained("R", "rstudio")

## Testing if R contained in rstudio
## [1] FALSE
is_contained("R", "rstudio", print_input = TRUE)
## Testing if R contained in rstudio
## [1] FALSE
is_contained("R", "rstudio", print_input = FALSE)
## [1] FALSE
is_contained("R", "rstudio", print_input = FALSE)
## [1] FALSE</pre>
```

Troubleshooting

- Your first self-written for-loop, or function, will probably not work.
- Don't panic! Just go line-by-line, keeping track of what is currently inside each variable.
- Stackoverflow is your friend.

Scoping rules in R

Scoping rules in R: Global environment (workspace)

When you write the name of a variable, R needs to find the value.

In the interactive computation (outside of functions, e.g., your console), this happens in the following order: - First, search the global environment (i.e., your workspace) - If it cannot be found, search each of the loaded packages

```
search()
```

```
[1] ".GlobalEnv"
                             "package:forcats"
                                                 "package:stringr"
##
    [4] "package:dplyr"
                             "package:purrr"
                                                 "package:readr"
##
  [7] "package:tidyr"
                             "package:tibble"
                                                 "package:ggplot2"
## [10] "package:tidyverse" "package:stats"
                                                 "package:graphics"
## [13] "package:grDevices"
                             "package:utils"
                                                 "package:datasets"
## [16] "package:methods"
                             "Autoloads"
                                                 "package:base"
```

The order of packages is important.

Scoping rules in R: Functions

Inside a function, this happens in the following order: - First, search within the function. - If it cannot be found, search in the global environment (i.e., your workspace) - If it cannot be found, search each of the loaded packages

```
y <- 3
```

```
test_t <- function() {
    print(y)
}

test_t()

## [1] 3

y <- 3

test_t <- function() {
    y <- 3
    print(y)
}

test_t()

## [1] 3</pre>
```

Scoping rules in R: Functions

What happens inside a function, stays within a function (unless you specify differently)

```
y <- 3

test_t <- function() {
    y <- 2
    print(y)
}

test_t()

## [1] 2

y</pre>
## [1] 3
```

Scoping rules in R: Packages

Packages are neatly contained/isolated, so they are not affected by your code.

They do so through namespaces. - Allow the package developer to hide functions and data. - Objects in the global environment that match objects in the function's namespace are ignored when running functions from packages (prevent clashes) - Functions are executed within the namespace of the package and have access to the global environment - They provide a way to refer to an object, with the double colon ::

```
dplyr::n_distinct(c(1,2,3,4,2))
## [1] 4
```

Practical