

An introduction to R – Part 2

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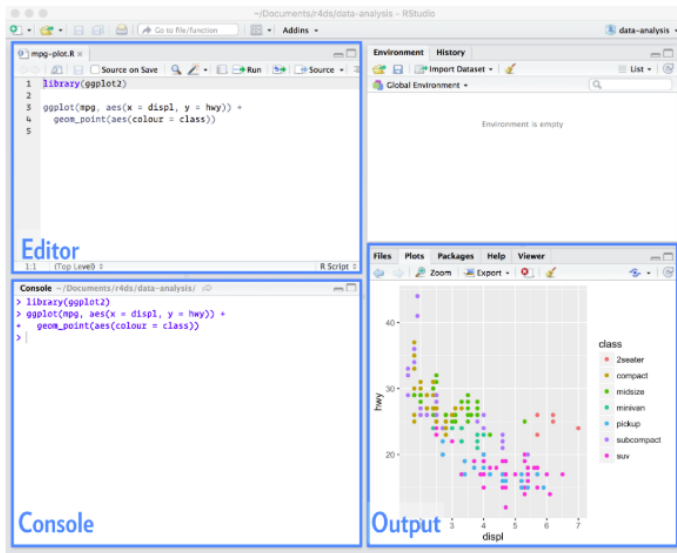
November 21, 2018

- 1 Programming with R
 - R Script
 - Control structures
 - User-defined Functions
 - Loop functions

R Script

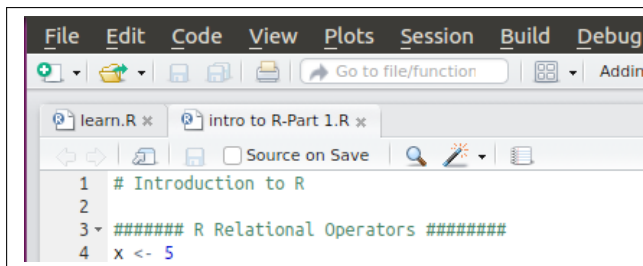
- Everything we have shown so far can be done using R console
- A script is a good way to keep track of what you're doing
- If you have a long analysis, and you want to be able to recreate it later, a good idea is to type it into a script

R Script



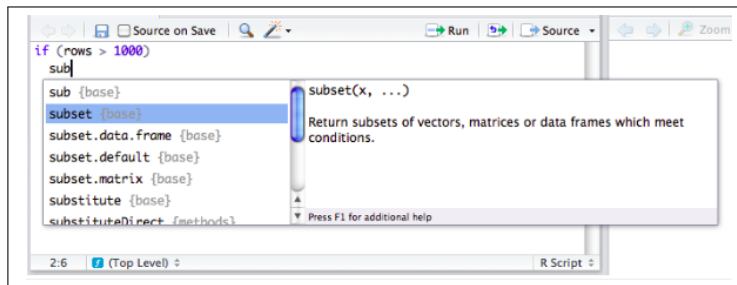
R Script file

- To create a new file you use the: **File -> New File -> R Script (Ctrl + Shift + N)**
- To open an existing file you use the **File -> Open File**
- If you open several files within RStudio they are all available as tabs



Code Completion

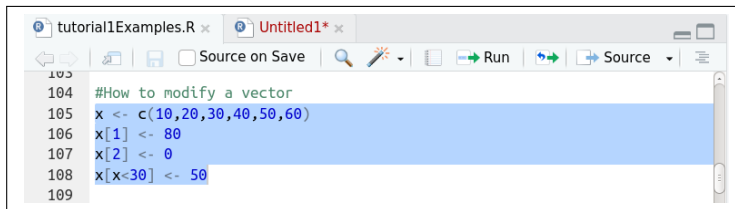
- RStudio supports the automatic completion of code using the Tab key
- If you have an function named subset, you can type sub and then Tab and RStudio will automatically complete the full name



Executing Code

Executing a single line

- To execute the line of source code where the cursor currently resides you press the **Ctrl + Enter** key (or use the Run toolbar button)
- After executing the line of code, RStudio automatically advances the cursor to the next line

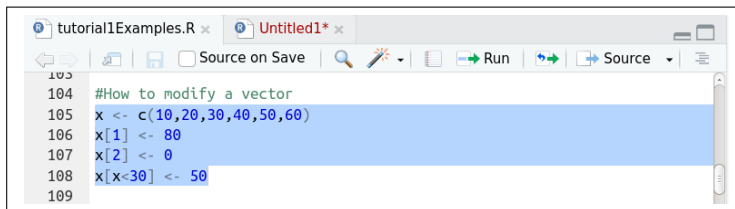


```
103
104 #How to modify a vector
105 x <- c(10,20,30,40,50,60)
106 x[1] <- 80
107 x[2] <- 0
108 x[x<30] <- 50
109
```


Executing Code

Executing multiple lines

- Select the lines and press the **Ctrl + Enter** key (or use the Run toolbar button)
- To run the entire document press the **Ctrl + Shift + Enter** key



The screenshot shows the RStudio IDE interface. The top toolbar includes buttons for 'Run' (a green arrow) and 'Source' (a blue arrow). Below the toolbar, a code editor window titled 'tutorial1Examples.R' contains the following R code:

```
103  
104 #How to modify a vector  
105 x <- c(10,20,30,40,50,60)  
106 x[1] <- 80  
107 x[2] <- 0  
108 x[x<30] <- 50  
109
```

The code lines 105 through 108 are highlighted in blue, indicating they are selected. The 'Run' button in the toolbar is also visible.

Keyboard Shortcuts

- Ctrl + Shift + N: New document
- Ctrl + O: Open document
- Ctrl + S: Save active document
- **Ctrl + 1**: Move focus to the Source Editor
- **Ctrl + 2**: Move focus to the Console

Control structures

Control structures

- Allow to control the flow of execution of a series of statements
- There are different types of control flow statements:
 - executing a set of statements **only if some condition is met**
 - executing a set of statements a given number of times
 - executing a set of statements **until** some condition is met
 - stop executing one thing or quit entirely
- Control structures allow to put some “logic” into the code

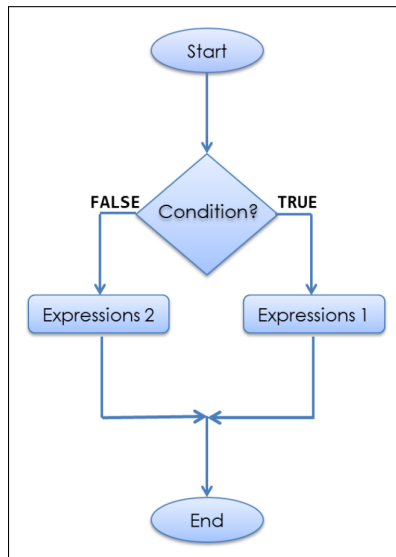
Conditional execution

If statements

- if statements allow to do different things based on the value of some **condition**.
- Conditional expressions in R.

Simple condition

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



If statements

```
> x <- 10
> if ( x > 0 )
{
  print("This is Positive number")
}
[1] "This is Positive number"
```

If-else statements

if-else

```
if (this) {  
  # Do that  
} else if (that) {  
  # Do something else  
} else {  
  #  
}
```

Example

```
> x <- -10  
> if(x >= 0)  
  {  
    print("Non-negative number")  
  } else {  
    print("Negative number")  
  }
```


If-else statements

```
> x <- c(2, 1, 3)
> if(sqrt(9) > 2)
{
  mean(x)
} else {
  sum(x)
}
[1] 2
```

```
> if(sqrt(9) > 4) { mean(x) } else { sum(x) }
[1] 6
```

ifelse

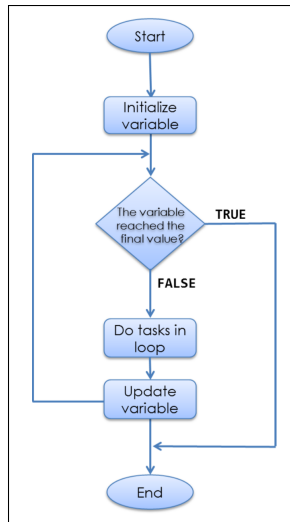
`ifelse` (test, TrueValue, FalseValue)

```
> x <- c(2, 1, 3, 6, 8, 1)
> y <- ifelse(x > 4, x, sum(x))
> y
[1] 21 21 21 6 8 21
```

Repetitive execution: for loops, repeat and while

Repetitive execution

- Loop is used to repeatedly carry out some computation.



Loop “For”

```
> for (variable in vector) {  
  commands  
}
```

```
> for (i in 1:5) {  
  print(i)  
}  
[1] 1  
[1] 2  
[1] 3  
[1] 4  
[1] 5
```

Loop “For”

Note that the variable that is set in a for loop is changed in the calling environment.

```
> i <- 1
> for (i in seq(5, 10, by = 2)) {
  print(i)
}

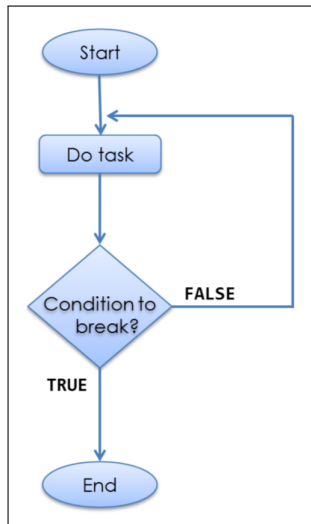
[1] 5
[1] 7
[1] 9

> i
[1] 9
```

Loop “Repeat”

- “Repeat”: repeats the same expression.
- The syntax is:

```
> repeat {  
  expression  
}
```



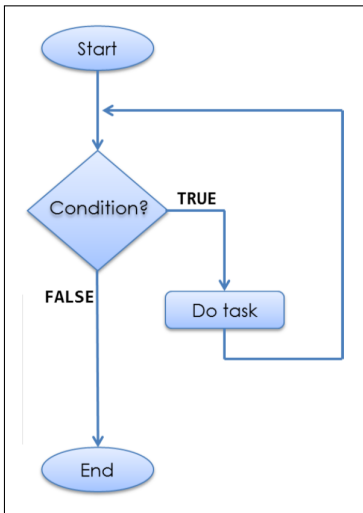
Loop “Repeat”

```
> x <- 7
> repeat {
  print(x);
  x <- x+ 2;
  if(x > 10) { break }
}
[1] 7
[1] 9
```


Loop “While”

- The while structure evaluates a expression as long as a stated condition is TRUE.
- The syntax is:

```
> while (condition) {  
    expression  
}
```



Loop “While”

```
> x <- 0
> while (x < 10) {
  print(x);
  x <- x + 5
}
[1] 0
[1] 5
```

User-defined Functions

Summary measures functions

- R offers functions to compute summary measures for quantitative variables
- These functions operate on a vector x

R function	Description
<code>mean(x)</code>	arithmetic mean of x
<code>median(x)</code>	median of x
<code>var(x)</code>	variance of x
<code>sd(x)</code>	standard deviation of x
<code>range(x)</code>	minimum and maximum values of x

User-defined Functions

- One of the great strengths of R is the user's ability to add functions
- The structure of a function is given below:

```
myfunction <- function(arg1, arg2, ... ){  
  statements  
  return(object)  
}
```

- Function elements:
 - Function name
 - Function arguments
 - Statements
 - Return

User-defined Functions

```
f <- function(num) {  
  for(i in seq_len(num)) {  
    print("Hello, world!")  
  }  
}  
f(3)|
```

- Function elements:
 - Function name: `f`
 - Function arguments: `num`
 - Statements: function's body
 - Return: It doesn't return anything
- **Tip: In general, if you find yourself doing a lot of cutting and pasting, that's usually a good sign that you might need to write a function**

User-defined Functions

```
f <- function(num) {  
  hello <- "Hello, world!"  
  for(i in seq_len(num)) {  
    print(hello)  
  }  
  chars <- nchar(hello) * num  
  chars  
}  
a = f(3)  
a
```

- Function elements:
 - Function name: `f`
 - Function arguments: `num`
 - Statements: function's body
 - Return: `chars`. In R, the return value of a function is always the very last expression that is evaluated.

User-defined Functions

```
f <- function(num) {  
  hello <- "Hello, world!"  
  for(i in seq_len(num)) {  
    print(hello)  
  }  
  chars <- nchar(hello) * num  
  chars  
}  
a = f(3)  
a
```

- The user must specify the value of the argument `num`. If it is not specified by the user, R will throw an error:

```
f()
```

```
Error in f(): argument "num" is missing, with no  
default
```


User-defined Functions

```
f <- function(num = 1) {  
  hello <- "Hello, world!"  
  for(i in seq_len(num)) {  
    print(hello)  
  }  
  chars <- nchar(hello) * num  
  chars  
}  
a = f()  
a
```

- We can modify this behavior by setting a default value for the argument `num`

User-defined Functions

Functions have named arguments which can optionally have default values. Because all function arguments have names, they can be specified using their name.

```
> f(2)
[1] "Hello, world!"
[1] "Hello, world!"
[1] 26
> |
```

Argument matching

R functions arguments can be matched *positionally* or by name. Given the following function:

```
g <- function(n, mean = 0, sd = 1) {...}
```

• `mydata <- g(100, 2, 1)`

- 100 is assigned to `n`
- 2 is assigned to `mean`
- 1 is assigned to `sd`

Argument matching

Given a function which sums all elements of the vector:

```
mysum <- function(vec, neg = TRUE) {...}
```

it has two arguments:

- `vec`: the vector of numbers
- `neg`: is a logical indicating whether negative values should be summed or not

Argument matching

Given a function which sums all elements of the vector:

```
mysum <- function(vec, neg = TRUE) {...}
```

- `mysum(v)`
 - Positional match first argument, and TRUE for `neg`
- `mysum(vec = v)`
 - Specify `vec` argument by name, and TRUE for `neg`
- `mysum(vec = v, neg = FALSE)`
 - Specify both arguments by name
- When specifying the function arguments by name, it doesn't matter in what order you specify them: `mysum(neg = FALSE, vec = v)`

The ... Argument

- Functions can have a special argument ...
- The ... argument is necessary when the number of arguments passed to the function cannot be known in advance

```
i01 <- function(...){  
  x <- list(...)  
  x  
}  
y <- i01(1,2,3)  
x <- i01(1,2,3,4,5)
```

Loop functions

Looping on the command line

- Writing “for” and “while” loops is useful but not particularly easy
- using loops in R is slow
- R has important family of functions which implement looping in a compact form

Functions

- **lapply()**: Loop over a [list](#) and evaluate a function on each element
- **sapply()**: same as “lapply” but try to simplify the result
- **apply()**: apply a function over the margins of an [array](#)

lapply()

- 1 `lapply()` loops over a list, iterating over each element in that list
 - 2 it applies a function to each element of the list (function specified by you)
 - 3 returns a list
- “`lapply()`” takes the following arguments: (1) a list, (2) a function, (3) other (`?lapply`).
 - The output will be coerced to a list using `as.list()`

lapply()

To apply the `mean()` function to all elements of a list.

```
x <- list(a = 1:5, b = 10:15)
lapply(x, mean)
```

Note that here we passed the `mean` function as an argument to the `lapply()` function.

sapply()

- The `sapply()` function behaves similarly to `lapply()`.
 - The only difference is in the return value
 - `sapply()` will try to simplify the result of `lapply()` (if possible).
- If the result is a list where every element is length 1, then a vector is returned.
 - If the result is a list where every element is a vector of the same length (> 1), a matrix is returned.
 - If it can't figure things out, a list is returned

supply()

```
x <- list(a = 1:4, b = 10:15, c = 20:25, d = 30:35)
> lapply(x, mean)
$a
[1] 2.5

$b
[1] 12.5

$c
[1] 22.5

$d
[1] 32.5
```

```
> sapply(x, mean)
a      b      c      d
2.5    12.5   22.5   32.5
```

apply()

The function `apply()` can be used to apply a function to the rows (second argument equal to 1) or columns (second argument equal to 2) of a matrix, arrays (> 2) dimensions or data.frames.

```
# matrix a  
> a  
> apply(a, 1, sum) # Apply sum to rows  
> apply(a, 2, mean) # Apply mean to columns
```

The result is equivalent to:

```
rowSums(a) # form row sums  
colMeans(a) # Apply mean to columns
```

References



Hadley Wickham & Garrett Golemund (2017)

R for data science: import, tidy, transform, visualize, and model data
O'Reilly.



Roger D. Peng (2015)

R Programming for Data Science



Venables W.N. & Smith D. M. (2018)

An introduction to R: Notes on R - A programming for Data Analysis and Graphics
Version 3.5.1.



Emmanuel Paradis (2005)

R for beginners



Link:

Advanced R

<https://adv-r.hadley.nz/>

The End

