

Functions - part 1

STAT 133

Gaston Sanchez

Department of Statistics, UC–Berkeley

`gastonsanchez.com`

`github.com/gastonstat/stat133`

Course web: `gastonsanchez.com/teaching/stat133`

Functions

R comes with many functions and packages that let us perform a wide variety of tasks. Sometimes, however, there's no function to do what we want to achieve. In these cases we need to create our own functions.

Anatomy of a Function

Anatomy of a function

`function()` allows us to create a function. It has the following structure:

```
function_name <- function(arg1, arg2, etc)
{
  expression_1
  expression_2
  ...
  expression_n
}
```

Anatomy of a function

- ▶ Generally we will give a name to a function
- ▶ A function takes one or more inputs (or none), known as *arguments*
- ▶ The expressions forming the operations comprise the body of the function
- ▶ Simple expression doesn't require braces
- ▶ Compound expressions are surrounded by braces
- ▶ Functions return a single *value*

Function example

A function that squares its argument:

```
square <- function(x) {  
  x * x  
}
```

Function example

A function that squares its argument:

```
square <- function(x) {  
  x * x  
}
```

- ▶ the function's name is "square"
- ▶ it has one argument `x`
- ▶ the function's body consists of one simple expression
- ▶ it returns the value `x * x`

Function example

It works like any other function in R:

```
square(10)
```

```
## [1] 100
```

In this case, `square()` is also vectorized

```
square(1:5)
```

```
## [1] 1 4 9 16 25
```

Why is `square()` vectorized?

Function example

Functions with a body consisting of a simple expression can be written with no braces (in one single line!):

```
square <- function(x) x * x
```

```
square(10)
```

```
## [1] 100
```

If the body of a function is a compound expression we use braces:

```
sum_sqr <- function(x, y) {  
  xy_sum <- x + y  
  xy_ssqr <- (xy_sum)^2  
  list(sum = xy_sum,  
        sumsqr = xy_ssqr)  
}
```

```
sum_sqr(3, 5)
```

```
## $sum  
## [1] 8  
##  
## $sumsqr  
## [1] 64
```

Function example

Once defined, functions can be used in other function definitions:

```
sum_of_squares <- function(x) sum(square(x))
```

```
sum_of_squares(1:5)
```

```
## [1] 55
```

Area of a Rectangle

A function which, given the values l (length) and w (width) computes the value $l \times w$

```
area_rect <- function(l, w) l * w
```

- ▶ The formal arguments of the function are `l` and `w`
- ▶ The body of the function consists of the simple expression `l * w`
- ▶ The function has been assigned the name "area_rect"

```
area_rect(5, 3)
```

```
## [1] 15
```

Evaluation of Functions

Function evaluation involves:

- ▶ A set of variables associated to the arguments is temporarily created
- ▶ The variable definitions are used to evaluate the body function
- ▶ Temporary variables are removed at the end
- ▶ The computed values are returned

Evaluation Example

Evaluating the function call `area_rect(5, 3)` takes place as follows:

- ▶ Temporarily create a variable `l` with value 5, and `w` with value 3
- ▶ Use those values to compute `l * w`
- ▶ Remove the temporary variable definition
- ▶ Return the value `15`

Test yourself

```
ft2 <- function(from, to) from:to^2
```

What does `ft2(1, 3)` return?

- A) 1 2 3 4 5 6 7 8 9
- B) 1 4 9
- C) 1 4 9 16 25 36 49 64 81
- D) 1 2 3
- E) 1 2 3 1 2 3

Nested Functions

We can also define a function inside another function:

```
getmax <- function(a) {  
  maxpos <- function(u) which.max(u)  
  list(position = maxpos(a),  
        value = max(a))  
}
```

```
getmax(c(2, -4, 6, 10, pi))
```

```
## $position  
## [1] 4  
##  
## $value  
## [1] 10
```


Function names

Different ways to name functions

- ▶ `squareroot()`
- ▶ `SquareRoot()`
- ▶ `squareRoot()`
- ▶ `square.root()`
- ▶ `square_root()`

Function names

Invalid names

- ▶ `5squareroot()`: cannot begin with a number
- ▶ `_sqrt()`: cannot begin with an underscore
- ▶ `square-root()`: cannot use hyphenated names

In addition, avoid using an already existing name, e.g. `sqrt()`

Function Output

Function output

- ▶ The body of a function is an expression
- ▶ Remember that every expression has a value
- ▶ Hence every function has a value

Function output

The value of a function can be established in two ways:

- ▶ As the last evaluated simple expression (in the body)
- ▶ An explicitly **returned** value via `return()`

The return() command

Sometimes the `return()` command is included to explicitly indicate the output of a function:

```
add <- function(x, y) {  
  z <- x + y  
  return(z)  
}
```

```
add(2, 3)
```

```
## [1] 5
```

The return() command

If no return() is present, then R returns the last evaluated expression:

```
# output with return()  
add <- function(x, y) {  
  z <- x + y  
  return(z)  
}  
  
add(2, 3)  
  
## [1] 5
```

```
# output without return()  
add <- function(x, y) {  
  x + y  
}  
  
add(2, 3)  
  
## [1] 5
```

The return() command

Depending on what's returned or what's the last evaluated expression, just calling a function might not print anything:

```
# nothing is printed  
add <- function(x, y) {  
  z <- x + y  
}  
  
add(2, 3)
```

```
# output printed  
add <- function(x, y) {  
  z <- x + y  
  return(z)  
}  
  
add(2, 3)  
  
## [1] 5
```


The return() command

Here we call the function and assign it to an object. The last evaluated expression has the same value in both cases:

```
# nothing is printed
add <- function(x, y) {
  z <- x + y
}

a1 <- add(2, 3)
a1

## [1] 5
```

```
# output printed
add <- function(x, y) {
  z <- x + y
  return(z)
}

a2 <- add(2, 3)
a2

## [1] 5
```

The return() command

If no return() is present, then R returns the last evaluated expression:

```
add1 <- function(x, y) {  
  x + y  
}
```

```
add2 <- function(x, y) {  
  z <- x + y  
  z  
}
```

```
add3 <- function(x, y) {  
  z <- x + y  
}
```

```
add4 <- function(x, y) {  
  z <- x + y  
  return(z)  
}
```

The return() command

return() can be useful when the output may be obtained in the middle of the function's body

```
f <- function(x, y, add = TRUE) {  
  if (add) {  
    return(x + y)  
  } else {  
    return(x - y)  
  }  
}
```

```
f(2, 3, add = TRUE)
```

```
## [1] 5
```

```
f(2, 3, add = FALSE)
```

```
## [1] -1
```

Function Arguments

Function arguments

Functions can have any number of arguments (even zero arguments)

```
# function with 2 arguments
add <- function(x, y) x + y

# function with no arguments
hi <- function() print("Hi there!")

hi()

## [1] "Hi there!"
```

Arguments

Arguments can have default values

```
hey <- function(x = "") {  
  cat("Hey", x, "\nHow is it going?" )  
}
```

```
hey()
```

```
## Hey  
## How is it going?
```

```
hey("Gaston")
```

```
## Hey Gaston  
## How is it going?
```

Arguments with no default values

If you specify an argument with no default value, you must give it a value everytime you call the function, otherwise you'll get an error:

```
sqr <- function(x) {  
  x^2  
}
```

```
sqr()
```

```
## Error in sqr(): argument "x" is missing, with no  
default
```

Arguments with no default values

Sometimes we don't want to give default values, but we also don't want to cause an error. We can use `missing()` to see if an argument is missing:

```
abc <- function(a, b, c = 3) {  
  if (missing(b)) {  
    result <- a * 2 + c  
  } else {  
    result <- a * b + c  
  }  
  result  
}
```

```
abc(1)
```

```
## [1] 5
```

```
abc(1, 4)
```

```
## [1] 7
```

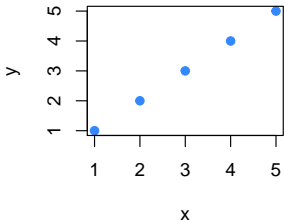

Arguments with no default values

You can also set an argument value to NULL if you don't want to specify a default value:

```
abcd <- function(a, b = 2, c = 3, d = NULL) {  
  if (is.null(d)) {  
    result <- a * b + c  
  } else {  
    result <- a * b + c * d  
  }  
  result  
}
```

More arguments

```
# arguments with and without default values  
myplot <- function(x, y, col = "#3488ff", pch = 19) {  
  plot(x, y, col = col, pch = pch)  
}  
  
myplot(1:5, 1:5)
```



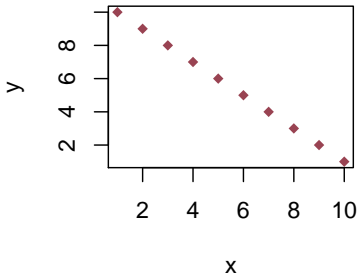
More arguments

```
# arguments with and without default values  
myplot <- function(x, y, col = "#3488ff", pch = 19) {  
  plot(x, y, col = col, pch = pch)  
}
```

- ▶ x and y have no default values
- ▶ col and pch have default values (but they can be changed)

More arguments

```
# changing default arguments  
myplot(1:10, 10:1, col = "#994352", pch = 18)
```



Positional and Named Arguments

```
output <- some_function(pos1, pos2, name1 = 1, name2 = "yes", name3 = 3)
```

- ▶ `pos1` positional argument
- ▶ `pos2` positional argument
- ▶ `name1` named argument
- ▶ `name2` named argument
- ▶ `name3` named argument

Argument Matching

- ▶ Arguments with default values are known as **named** arguments
- ▶ Arguments with no default values are referred to as **positional** arguments
- ▶ Arguments can be matched positionally or by name

Argument Matching

```
values <- seq(-2, 1, length.out = 20)
```

```
# equivalent calls
```

```
mean(values)
```

```
mean(x = values)
```

```
mean(x = values, na.rm = FALSE)
```

```
mean(na.rm = FALSE, x = values)
```

```
mean(na.rm = FALSE, values)
```

Partial Matching

Named arguments can also be partially matched:

```
# equivalent calls  
seq(from = 1, to = 2, length.out = 5)  
seq(from = 1, to = 2, length = 5)  
seq(from = 1, to = 2, len = 5)
```

`length.out` is partially matched with `length` and `len`

Matching Order

Order of argument matching operations:

- ▶ Check for exact match for a named argument
- ▶ Check for a partial match
- ▶ Check for a positional match

Example

Write a function that checks if a number is positive (output TRUE) or negative (output FALSE)

Example

Write a function that checks if a number is positive (output TRUE) or negative (output FALSE)

```
is_positive <- function(x) {  
  if (x > 0) TRUE else FALSE  
}
```

```
is_positive(2)
```

```
## [1] TRUE
```

```
is_positive(-1)
```

```
## [1] FALSE
```

Example

Write a function that checks if a number is positive (output TRUE) or negative (output FALSE)

```
# a simpler way  
is_positive <- function(x) {  
  x > 0  
}
```

Example

Write a function that checks if a number is positive (output TRUE) or negative (output FALSE)

```
# no need to do this  
is_positive <- function(x) {  
  if (x > 0) print('TRUE') else print('FALSE')  
}
```

Remember that every function has a value: the last statement that is evaluated (or an output from return())

Example

What happens in these cases?

```
is_positive(0)
```

```
is_positive(NA)
```

```
is_positive(TRUE)
```

```
is_positive("positive")
```

```
is_positive(1:5)
```

Using arguments for other functions

There are various functions that include the argument `na.rm` to indicate if missing values should be removed. One of them is `mean()`:

```
# default na.rm = FALSE
mean(c(1, 2, 3, NA, 5))

## [1] NA

# default na.rm = TRUE
mean(c(1, 2, 3, NA, 5), na.rm = TRUE)

## [1] 2.75
```

Using arguments for other functions

If we create a function that uses other functions containing `na.rm`, it is wise to include that argument:

```
meansd <- function(x, na.rm = FALSE) {  
  c(mean = mean(x, na.rm = na.rm),  
    sd = sd(x, na.rm = na.rm))  
}
```

```
meansd(c(1, 2, 3, NA, 5), na.rm = TRUE)
```

```
##      mean      sd  
## 2.750000 1.707825
```


Dots argument

If you check functions like `c()`, `paste()`, `plot()`, you'll notice the use of a special argument: `...`

- ▶ it matches zero, one or more actual arguments
- ▶ it allows us to pass arguments to other functions inside the function
- ▶ `...` allows us to “cascade” arguments to other functions without including them in the definition

Dots argument

Using ...

```
fplot <- function(y, ...) {  
  x <- 1:length(y)  
  plot(x, y, type = 'n', ylim = c(0, 1), ...)  
  points(x, y, col = "#93a8f2", pch = 19)  
  lines(x, y, col = "#dd93f2", lwd = 3)  
}  
  
fplot(runif(10), bty = 'n')  
fplot(runif(10), bty = 'n', main = "some title")  
fplot(runif(10), bty = 'n', xlab = '')
```