Functions in R

Day 1 - Introduction to Data Analysis with R

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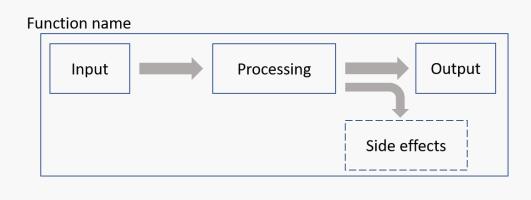
March 2, 2025

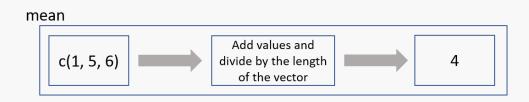
Functions in R

Functions make multiple operations available under one command.

Functions:

- Have a name
- Have (0, 1, or any number of)
 arguments as input
- They calculate something using the arguments
- They have a return value (the output)
- Can have side effects (like plotting)





General structure of a function call: function_name (argument = value)

The mean function

```
function_name ( argument = value )
```

```
mean (x = c(1,5,6))
```

```
mean(x = c(1,5,6)) # or short: mean(c(1,5,6)) #> [1] 4
```

- Arguments can also be variables
- The output of a function can be stored in a variable

```
values <- c(1,5,6)
result <- mean(x = values)
result
#> [1] 4
```

The mean function

But what does the mean function do? What are the arguments that I can use?

→ Call the function help using ?

?mean

```
mean {base}
                                                                                                         R Documentation
Arithmetic Mean
Description
Generic function for the (trimmed) arithmetic mean.
Usage
mean(x, ...)
## Default 83 method:
mean(x, trim = 0, na.rm = FALSE, ...)
Arguments
         An R object. Currently there are methods for numeric/logical vectors and date, date-time and time interval objects.
         Complex vectors are allowed for trim = 0, only.
        the fraction (0 to 0.5) of observations to be trimmed from each end of x before the mean is computed. Values of trim
         outside that range are taken as the nearest endpoint.
        a logical value indicating whether NA values should be stripped before the computation proceeds.
         further arguments passed to or from other methods.
Value
If trim is zero (the default), the arithmetic mean of the values in x is computed, as a numeric or complex vector of length one. If
x is not logical (coerced to numeric), numeric (including integer) or complex, NA real is returned, with a warning.
If trim is non-zero, a symmetrically trimmed mean is computed with a fraction of trim observations deleted from each end
before the mean is computed.
```

- Arguments are the input to a function
- Functions can provide default values for some arguments
- Default values for arguments are indicated in the function help

```
Arguments

An R object. Currently there are methods for numeric/logical vectors and date, date-time and time interval objects. Complex vectors are allowed for trim = 0, only.

the fraction (0 to 0.5) of observations to be trimmed from each end of x before the mean is computed. Values of trim outside that range are taken as the nearest endpoint.

a logical evaluating to TRUE or FALSE indicating whether NA values should na.rm be stripped before the computation proceeds.

further arguments passed to or from other methods.
```

What happened here?

```
# NA is a missing value
values <- c(1, 5, 6, NA)
mean(x = values)
#> [1] NA
```

→ na.rm argument is FALSE by default.

Set it to TRUE if you want to calculate the mean despite missing values:

```
mean(x = values, na.rm = TRUE)
#> [1] 4
```

Arguments with default values are optional, arguments without default values are not!

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

Argument matching can be achieved by position or by name

```
mean(x, trim = 0, na.rm = FALSE, ...)
values <- c(1, 5, 6, NA)
```

These calls to mean all are the same:

```
mean(values, , TRUE) # by position
mean(x = values, na.rm = TRUE) # by name
mean(values, na.rm = TRUE) # a mix of both
```

Argument matching can be achieved by position or by name

Named arguments are (generally) preferred

- Easier to remember
- Easier to read
- Some functions have a lot of arguments

```
mean(x = values, na.rm = TRUE) # by name
```

However, it is common to match the first argument by position (especially when the first argument is the data)

```
mean(values, na.rm = TRUE) # a mix of both
```

Where do functions come from?

Base R functions: built into R

mean() # calculate mean seq() # generate a sequence of values

From additional packages

- Packages must be installed first
- Call a function from a package using packageName::functionName()
- Load the package with
 library(packageName) and then use
 the function

Custom functions:

Write your own functions and then use them in the code

```
# use read_csv function from readr package
readr::read_csv()

# or use library()
library(readr)
read_csv()
```

```
# custom function that prints input in a sentence
my_function <- function(x) {
   return(x + 5)
}
my_function(5)</pre>
```

Summary

R basics - Functions

Summary

- Functions take input in the form of (named) arguments, calculate something and return a result
- Functions are called by their name, followed by parentheses:
 functionName(argument1 = value, argument2 = value, ...)
- Functions from additional packages can be called in two ways:
 - packageName::functionName()
 - first load the package with library(packageName) then call the function anywhere in the script with functionName()
- Call ?functionName to open the help of a function