



## Do it yourself: R Package

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# Functions

- In R, everything is an object
- Most objects are functions (demo: `t.test`; `+`; parenthesis, ...)
- R Packages mainly consist of functions
- Define own functions → structures code and improves maintainability

## Examples (bad and good)

```
library(ggplot2)
```

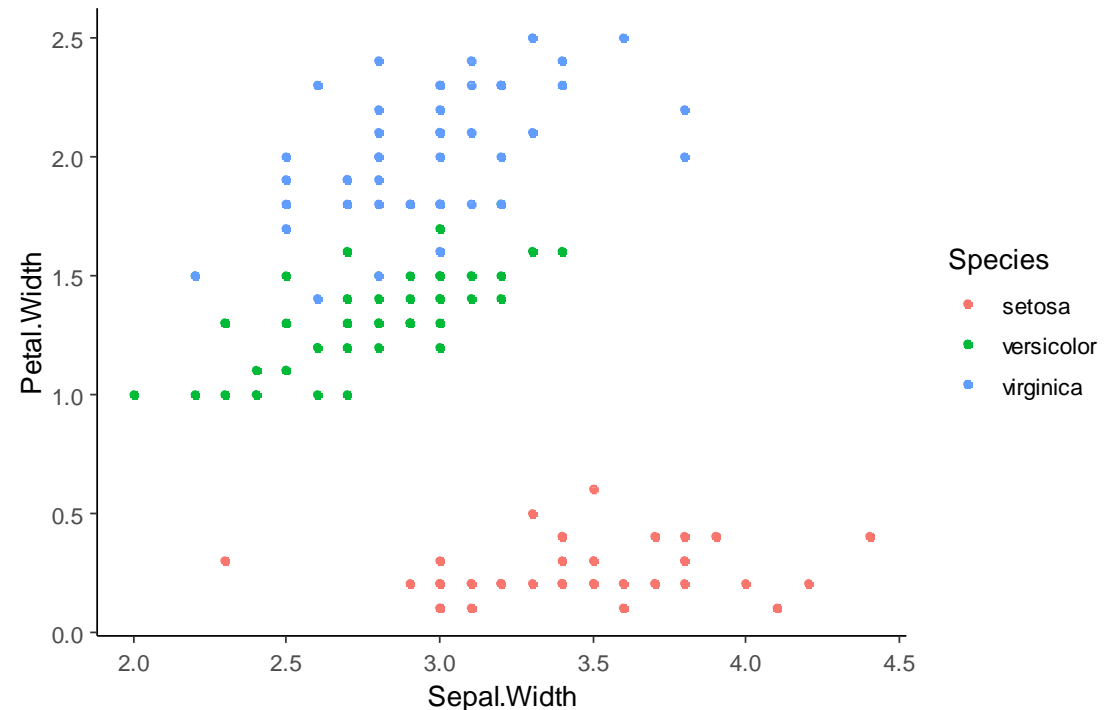
### # Bad

```
ggplot(iris, aes(x = Sepal.Width, y = Petal.Width)) +  
  geom_point(aes(color = Species)) +  
  theme_classic(base_size = 12)
```

```
ggplot(iris, aes(x = Sepal.Length, y = Petal.Width)) +  
  geom_point(aes(color = Species)) +  
  theme_classic(base_size = 12)
```

### # Good

```
scatter <- function(x, y = "Petal.Width") {  
  ggplot(iris, aes_string(x = x, y = y)) +  
    geom_point(aes(color = Species)) +  
    theme_classic(base_size = 12)  
}  
scatter("Sepal.Width")  
scatter("Sepal.Length")
```



# Separate Functions from Analysis

If your analysis uses several self-written functions, it often makes sense to

1. put them in a script "functions.R" and
2. source this in the analysis script by

```
source("functions.R")
```

Example of a "functions.R" file

```
# Scatter of y against x
```

```
scatter <- function(x, y = "Petal.Width") {  
  ggplot(iris, aes_string(x = x, y = y)) +  
    geom_point(aes(color = Species)) +  
    theme_classic(base_size = 12)  
}
```

```
# Does something fancy
```

```
something_fancy <- function(...) {  
  ...  
}
```



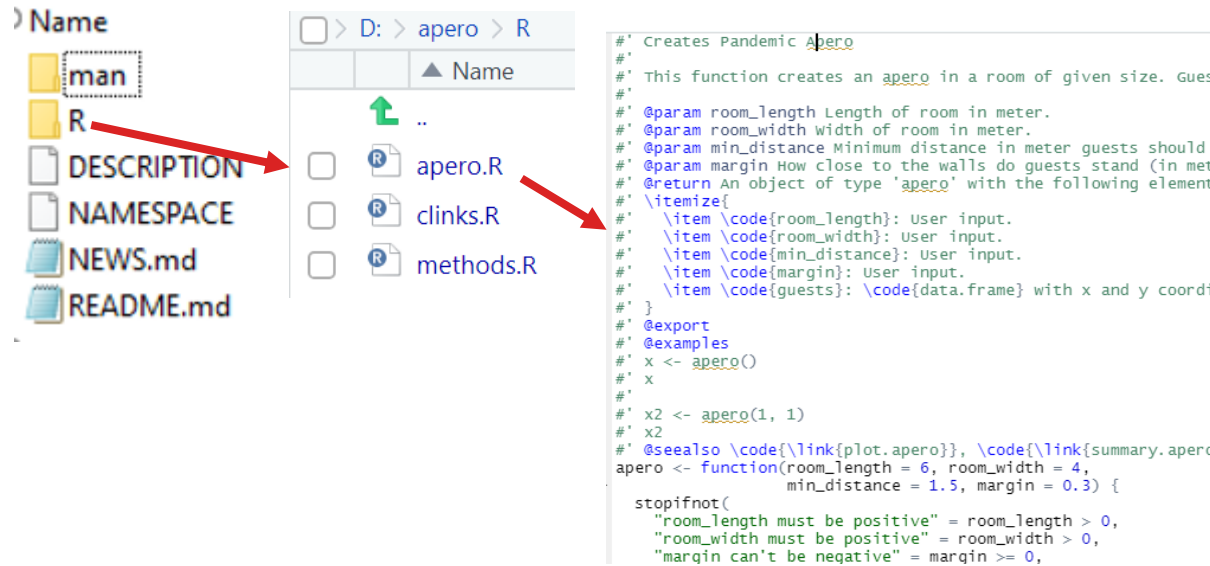


# Well-documented Functions are almost a Package

When is it worth to turn a "functions.R" script into a package?

- If the same functions are used across multiple projects.
- Collaboration with external persons.
- You want to publicly share something useful with others (via Github or CRAN).

Typical content of an R package



The image shows a file explorer view of an R package directory. On the left, a list of files includes 'man', 'R', 'DESCRIPTION', 'NAMESPACE', 'NEWS.md', and 'README.md'. A red arrow points from the 'R' folder to a sub-view of the 'R' directory. This sub-view shows files: '..', 'apero.R', 'clinks.R', and 'methods.R'. Another red arrow points from 'apero.R' to a code editor window. The code editor displays the source code for the 'apero' function, which is a well-documented R function. The code includes comments in French describing the function's purpose and parameters, followed by R code for parameter handling, validation, and plotting.

```
##' Creates Pandemic apero
##'
##' This function creates an apero in a room of given size. Guest
##'
##' @param room_length Length of room in meter.
##' @param room_width width of room in meter.
##' @param min_distance Minimum distance in meter guests should
##' @param margin How close to the walls do guests stand (in meter)
##' @return An object of type 'apero' with the following element
##' \itemize{
##'   \item room_length: User input.
##'   \item room_width: User input.
##'   \item min_distance: User input.
##'   \item margin: User input.
##'   \item guests: \code{data.frame} with x and y coordi
##' }
##'
##' @export
##' @examples
##' x <- apero()
##' x
##'
##' x2 <- apero(1, 1)
##' x2
##' @seealso \link{plot.apero}, \link{summary.apero}
apero <- function(room_length = 6, room_width = 4,
                  min_distance = 1.5, margin = 0.3) {
  stopifnot(
    "room_length must be positive" = room_length > 0,
    "room_width must be positive" = room_width > 0,
    "margin can't be negative" = margin >= 0,
```

Demo of "apero" package.  
Note: In Switzerland, we have "Apéro" all the time...



## Remark: print(), summary(), plot(), ...

R has a very simple system for object-specific functions (S3)

### Example 1

`summary()` on a `data.frame` does something different than on a numeric vector.

### Example 2 (copy to R and short demo)

```
employee <- function(given_name, family_name) {  
  out <- list(  
    given_name = given_name,  
    family_name = family_name  
  )  
  class(out) <- "employee"  
  out  
}
```

```
print.employee <- function(x) {  
  cat("You are", x$given_name, x$family_name)  
}
```

```
me <- employee("Michael", "Mayer")  
me # or print(me)
```

### Example 3: t.test()

# Creating the "apero" Package

Two packages are of key importance to build your own package

- **usethis**: Cares about **content** of package
  - **devtools**: Turns content into **package**
- } By Prof. Jenny Bryan, RStudio et al.

## Preparation

- You need **R**, **RStudio**, and **RTools**.
- Put your R functions into one or multiple R scripts.
- Document them in Roxygen style.

Then continue as in <https://github.com/mayer79/apero> → script "packaging.R"

## Demo

## Some final Words

In R code of package,

- never use `source()`, `setwd()`,  
`library()`, `require()`
- use defensive programming

Everything on <https://github.com/mayer79/apero>  
`devtools::install_github("mayer79/apero")`

## Test with your own code!

Hadley's book free online

<https://r-pkgs.org/>

