### Developing an R package: a tutorial

The essentials to write your package

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## **Getting started**

#### Note

If you are not working from Rstudio, you will not benefit from all its functionality but it is possible to do everything from the R command line.

#### Setup your environment

 install R packages providing development tools<sup>1</sup>: devtools, usethis, roxygen2, lintr

```
install.packages(c("devtools", "usethis", "roxygen2", "lintr"))
```

for a more complete setup: see https://r-pkgs.org/setup.html

¹devtools may require to install additional system libraries depending on your OS, a quick search on the web will help you if you encounter any error.

#### Create a package

• Initialize a package template:

```
usethis::create_package("mypkg")
```

- Directly from Rstudio (equivalent): File New Project New directory R package
- Attention: if you want to initialize an R package without initializing an Rstudio project<sup>2</sup>, use:

```
usethis::create_package("mypkg1", rstudio = FALSE, open = FALSE)
```

 $<sup>^2</sup>$ e.g. because you want to create your package in an existing Rstudio project, or you don't use Rstudio

#### Rstudio project

- Project specific configuration, workspace, history
- Isolated R environment for the project
- RStudio project management feature (e.g. git management)
- More information regarding Rstudio project at https://r-pkgs.org/workflows101.html#projects

### The "old-fashion" built-in R function to create package

(for more advanced users)

```
## two functions and two "data sets" :
f <- function(x, y) x+y
g <- function(x, y) x-y
d <- data.frame(a = 1, b = 2)
e <- rnorm(1000)
## automatically "fill" the package
package.skeleton(list = c("f", "g", "d", "e"), name = "mypkg2")</pre>
```

**Attention:** using package.skeleton() creates a package that is not ready "out-of-the-box", you will have to edit and fix the help pages (e.g. by using roxygen2, c.f. later).

#### Naming your package

- three formal requirements:
  - "The name can only consist of letters, numbers, and periods, i.e., .."
  - "It must start with a letter."
  - "It cannot end with a period."
- Advice: use a catchy name or acronym with a link to your package functionality
- More details on naming convention at https://r-pkgs.org/workflows101.html#naming

#### Always choose a license!

- It governs the possibility to use, modify or redistribute a software
- It helps to identify clear authorship/copyright<sup>3</sup>
- Without a license: fuzzy and unclear (generally "all rights reserved" but you are never sure<sup>4</sup>)

<sup>&</sup>lt;sup>3</sup>depending on legal consideration, varying from one country to another

<sup>&</sup>lt;sup>4</sup>"Was it forgotten or a deliberate choice?"

#### Different types of license

- Use a software-specific license for software and a content-specific license for data<sup>5</sup>
- Recommandation: favor free<sup>6</sup> and open-source licenses (versus proprietary or closed licenses), either permissive or with copyleft

<sup>&</sup>lt;sup>5</sup>e.g. Creative Commons license are for contents and not for software

<sup>&</sup>lt;sup>6</sup>as in "libre" and not as in "gratis" (proprietary software can be gratis)

#### How to choose a license?

```
See https://r-pkgs.org/license.html (and functions usethis::use_XX_license()<sup>7</sup> from the usethis package<sup>C*</sup>)
```

Additional resources on software license:

- https://choosealicense.com
- https://opensource.org/licenses
- https://www.gnu.org/licenses/license-list.en.html

<sup>7</sup>e.g. use\_mit\_license() or use\_gpl\_license()

# R package structure

### Files and sub-directories (1)

#### Empty package:

```
mypkg
+-- DESCRIPTION
+-- NAMESPACE
+-- R
+-- (empty)
```

#### More complete package:

```
mypkg2
+-- data
   +-- d.rda
 +-- e.rda
+-- DESCRIPTION
+-- man
  +-- d.Rd
  +-- e.Rd
 +-- f.Rd
  +-- g.Rd
   +-- mypkg2-package.Rd
+-- NAMESPACE
+-- R
   +-- f.R
   +-- g.R
```

### Files and sub-directories (2)

- Meta-data files: DESCRIPTION and NAMESPACE (c.f. later)
- R sub-directory: where to store R source files implementing the function included in your package
- man sub-directory: where to store the mandatory help pages
- src sub-directory (optional): where to store code to be compiled (written in other languages, not in R) included in your package
- data sub-directory (optional): where to store data files attached to your package

### R source code (1)

#### The R sub-directory:

- Write your code as functions
- Save your code implementing functions in R source code files<sup>8</sup>
- Group related functions in the same file
- Create and edit source code files manually or with usethis::use\_r("name") ☑
- See R code formatting convention<sup>™</sup>
- Check your code formatting with the lintr package and lintr::lint\_package()

#### R source code (2)

#### Debugging-friendly advice:

- Avoid very long functions (split long functions into several shorter ones)
- Factorize re-used code into specific functions (avoid copying-pasting chunk of codes several time)

#### From R scripts to R functions

#### Scripting:

```
# data
a = 7
b = 3
# intermediate operations
tmp1 = 2 * a
tmp2 = b / 6
# final computations
c = tmp1 + tmp2
```

**Objective:** simplify your code by "hiding" intermediate steps into a function

Implementing functions in your package:

```
myFun <- function(x, y) {
   tmp1 = 2 * x
   tmp2 = y / 6
   return(tmp1 + tmp2)
}</pre>
```

In a script using your package:

```
library(mypkg)
# data
a = 7
b = 3
# computations
c = myFun(a,b)
d = myFun(10, 3)
```

#### Meta-data files

- DESCRIPTION: a structured text file giving information about your package (title, description , authors , license , dependencies , etc.)
- NAMESPACE: a text file indicating<sup>9</sup> names of R objects (functions, datasets)
  that are imported in your package (from other packages), and/or exported
  by your package (to be usable when you install your package)

 $<sup>^{9} \</sup>mathrm{and}/\mathrm{or}$  the name of the dynamic library related to compiled codes to be used in your package if relevant

### **DESCRIPTION** file (1)

- Can be edited manually, or created and modified with usethis::use\_description() and other usethis::use\_XXX()
- Setup your package requirements and dependencies (c.f. later)
- More details at https://r-pkgs.org/description.html
- Important: package versioning<sup>™</sup>

### **DESCRIPTION** file (2)

#### Example:

```
Package: mypkg
Title: What the Package Does (One Line, Title Case)
Version: 0.0.0.9000
Authors@R:
    person(given = "First",
          family = "Last",
          role = c("aut", "cre").
           email = "first.last@example.com",
           comment = c(ORCID = "YOUR-ORCID-ID"))
Description: What the package does (one paragraph).
License: `use_mit_license()`, `use_gpl3_license()` or friends to pick a
    license
Encoding: UTF-8
LazvData: true
Roxygen: list(markdown = TRUE)
RoxygenNote: 7.1.1
```

#### **NAMESPACE** file

Can be created with usethis::use\_namespace() , or edited manually, or (better) automatically updated thanks to roxygen2 inline documentation (c.f. later)

#### Example:

```
# Generated by roxygen2: do not edit by hand
importFrom(stats, runif)
export(my_function)
```

### Data in your package

- Binary R data file (.Rda or .Rdata file) can be stored in the data sub-directory
- Raw data can be stored in the inst sub-directory and found after installation with the system.file() function<sup>10</sup> (important: you do not need to search for the inst sub-directory, just anything in it)

```
system.file(filename, package = "mypkg")
system.file(dirname, package = "mypkg")
system.file(package = "mypkg") # package root directory
```

More at https://r-pkgs.org/data.html

 $<sup>^{10}</sup>$ any other file or sub-directory shipped with a package can be found likewise

#### help/man pages

- Available with the R commands ?function\_name (e.g. ?rnorm) or help(function\_name)
- Content:
  - usage description and functioning details
  - input arguments and return value description
  - function authorship
  - link to related functions
  - bibliographic reference
  - minimum working examples
- Encoded in .Rd files in the man sub-directory: structured text files with a specific syntax

### Document your code and generate help page at once

- Avoid creating and editing .Rd file manually (laborious)
- Good practice: DOCUMENT YOUR CODE (with inline comments) for other and YOUR FUTURE SELF
- Inline code documentation with roxygen2 based on tags identified with @

### Document your code and your package with roxygen2 (1)

# Inline code documentation (identified with #' comment characters):

```
Add together two numbers
# 1
   Oparam x A number
#' @param y A number
#' @return The sum of \code{x} and \code{y}
   Qauthor Anonymous
   @examples
\#' \ add(1, 1)
\#' a.d.d.(10. 1)
#' @export
add <- function(x, v) {
 x + v
```

#### Corresponding . Rd file:

```
% Generated by roxygen2 (3.2.0): do not edit by hand
\name{add}
\alias{add}
\title{Add together two numbers}
\usage{
add(x, y)
\arguments{
 \item{x}{A number}
 \item{v}{A number}
\value{
The sum of \code{x} and \code{v}
\description{
Add together two numbers
\examples{
add(1, 1)
add(10, 1)
```

### Document your code and your package with roxygen2 (2)

- Generate the man pages (and update NAMESPACE file) with devtools::document() or in Rstudio interface (Build panel More Document<sup>11</sup>)
- Identify exported functions<sup>12</sup> with the tag @export (automatically added to the NAMESPACE file)
- Identify imported functions<sup>13</sup> with the tag @importFrom package function (automatically added to the NAMESPACE file)

 $<sup>^{11}</sup>$ keyboard shortcut: CTRL + SHIFT + D

<sup>&</sup>lt;sup>12</sup>your functions that will be available to users

<sup>&</sup>lt;sup>13</sup>functions from other packages that you use

### Document your code and your package with roxygen2 (4)

#### More complete example:

```
#' A function to do some stuff
#' @description
#' Do some stuff
#' @details
#' I do the stuff in a complicated way.
#' @param x A number
#' Oparam y A number
#' Greturn what the function is returning
#' Qauthor Someone
#' @importFrom stats rnorm
#' @seealso [mupka::mu other fun()]
#' @examples
#' add(1, 1)
#' add(10, 1)
#' @export
mv fun <- function(x, v) {
    tmp = rnorm(7)
```

#### Tips:

- add Roxygen: list(markdown = TRUE) to the DESCRIPTION file to use markdown syntax in documentation chunks or run usethis::use\_roxygen\_md() (possible conversion from existing standard roxygen2 syntax with roxygen2md package)
- internal functions (only used by other functions in your package, and not to be available for users) can be tagged with @keywords internal and (and without @export tag)

### Document your code and your package with roxygen2 (5)

#### References:

- More details at https://r-pkgs.org/man.html
- roxygen2 cheat sheet <sup>©</sup>
- Help to format your documentation chunks at https://roxygen2.r-lib.org/articles/rd-formatting.html and https://roxygen2.r-lib.org/articles/rd.html

### Manage your dependencies (1)

Several fields in the DESCRIPTION file:

- Depends: R (>= 3.1.0): the minimal R version required by your package
- Imports: ...: packages (with optional minimal versions) required for your package to work
- Suggests: ... (optional): additional packages (with optional minimal versions) that are not necessary for your package to work but that would improve the user experience with your package
- Additional (optional) fields: LinkingTo (useful if external codes needs to be compiled and linked against external library), OS\_type: unix (to specify which OS are supported<sup>14</sup>), SystemRequirements: C++11 (to specify additional external system requirements<sup>15</sup>)

<sup>&</sup>lt;sup>14</sup>here it means that Windows is not supported

<sup>&</sup>lt;sup>15</sup>here a C++ compiler compatible with C++11 standard

### Manage your dependencies (2)

```
Example:
Depends: R (>= 3.1.0)
LinkingTo:
    Rcpp (>= 1.0.1),
    RcppEigen (>= 0.3.3.5)
Imports:
    Rcpp (>= 1.0.1),
    openssl
Suggests:
    testthat (>= 2.1.0)
SystemRequirements: C++11
```

### Manage your dependencies (3)

- usethis::use\_package() to update Imports or Sugggests fields
- More details at https://r-pkgs.org/description.html#dependencies

### Manage your dependences (4)

Imported objects (functions, dataset) should be declared in the NAMESPACE file

ightarrow automatically manage thanks to roxygen2

### Other (optional) files and sub-directories (1)

- README<sup>16</sup>, LICENSE (depending on the license your choose), COPYRIGHT (to detail authorship, copyright associated to the package content)
- src<sup>©</sup>: source codes to be compiled (c.f. later)
- inst<sup>®</sup> to store additional files (e.g. required for tests, vignettes, etc.), raw data, etc.
- tests  $^{\ \ \ }$  to write automatic tests (c.f. later)

 $<sup>^{16}</sup>possible \ format: .md \ or .Rmd (Rmarkdown), see usethis::use_readme_md() or usethis::use_readme_rmd() <math display="inline">^{\c C}$ 

### Other (optional) files and sub-directories (2)

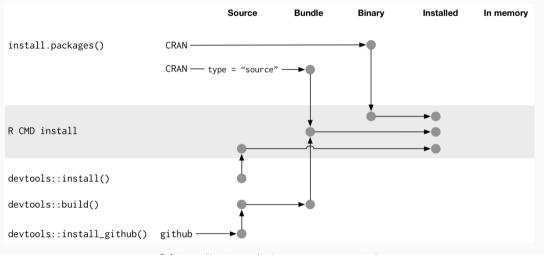
- More details at https://r-pkgs.org/misc.html
- Non-standard files can be present in your project but not shipped in your package: you should create a .Rbuildignore file

### Workflow

#### Package state

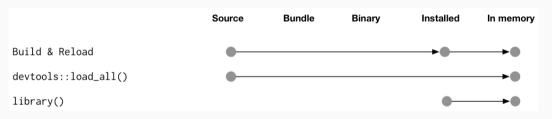
- source: what you are writing
- bundled: a single-file compressed version containing your package source (how package are shipped by the CRAN for installation)
- binary: a single-file binary version of your package containing compiled library (if relevant), mainly used by the CRAN to ship package for installation on Windows
- installed: available as a library on your system, i.e. the package files and sub-directories (along with library files if compilation was needed) have been copied somewhere on your computer
- in-memory: loaded and ready to use (after calling library(mypkg))

### Dev workflow (building and installing a package)



Ref: https://r-pkgs.org/package-structure-state.html

### Dev workflow (loading a package)



Ref: https://r-pkgs.org/workflows101.html#load-all

#### load

Load your package for a test drive (manual test) without building/installing it

- devtools::load\_all() <sup>©</sup>
- in Rstudio interface (Build panel More Load all<sup>17</sup>)

Development cycle: write code, test it, correct your code, test it, etc.

<sup>&</sup>lt;sup>17</sup>keyboard shortcut: CTRL + SHIFT + L

#### document (reminder)

#### Generate the man pages (and update NAMESPACE file)

- devtools::document() <sup>C</sup>
- in Rstudio interface (Build panel More Document<sup>18</sup>)

<sup>&</sup>lt;sup>18</sup>keyboard shortcut: CTRL + SHIFT + D

#### build

#### Prepare your package for installation (and distribution)

- devtools::build() <sup>©</sup>
- in Rstudio interface (Build panel More "Build source package")
- R built-in shell command line tool<sup>19</sup>: R CMD build mypkg
- Create a .tar.gz archive files containing the sources (or a .zip file if you use "Build binary package") ready for installation

<sup>&</sup>lt;sup>19</sup>R.exe on Windows

#### check

Verify that your package is functional and that your package structure is correct

- devtools::check()
- in Rstudio interface (Build panel Check)
- R built-in shell command line tool<sup>20</sup>: R CMD check mypkg\_1.0.0.tag.gz
- Verbose output: often clearly identify problems (and suggest fixes)
- More details at https://r-pkgs.org/r-cmd-check.html

<sup>&</sup>lt;sup>20</sup>R. exe on Windows

#### usethis (exhaustive tour)

See https://usethis.r-lib.org/reference/index.html

Possible to write every files manually for more advanced users.

#### devtools (exhaustive tour)

See https://devtools.r-lib.org/reference/index.html

(devtools exports several functions from other development-oriented packages)