Developing an R package: a tutorial

Going further with your R package development

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Getting started

Additional R packages to help you create R packages

- testthat [™]: to implement automatic tests of your functions
- remotes . to install package from anywhere (integrated in devtools)
- rmarkdown[®] and knitr[®]: to create detailed documentation materials and notebooks (code showcase)
- pkgdown [™] to create a website for your package

Setup your environment

• install additional R packages providing development tools: testthat, remotes¹

install.packages(c("testthat"))

¹not necessary if you already installed devtools

Additional references regarding R programming

• Hadley Wickham book: Advanced R (web version and sources)

Digression: Good practice for software development and programming (not just in R)

Good practice (1)

- The code should be human readable² and easily understandable (use comments, code presentation and formatting)
 - Experiment: read your (5 weeks/months/years) old codes, are you sure that you will understand it? (worst with code written by others)
- Use a versioning system (e.g. git[™]) to manage your code evolution/version and for collaborative development

²being machine readable is necessary for the code to work but not sufficient

Good practice (2)

- Implement automatic tests (e.g. unit tests) for each new function/module/etc.
 (and not afterward) to verify your implementation and results and avoid breaking your code³
- Use continuous integration⁴: to automatically run build, check, tests as your package development progresses (e.g. commit after commit if you are using a versioning system like git)

³never trust yourself, you will implement bugs

⁴software forge offers such service like gitlab CI/CD ^C or github actions ^C

Good practice (3)

- Write a documentation for your code/package/library, including explained code showcases/demos
- Publish your source codes (preferably on a software forge), so that other can continue your work, especially when you move on to other projects, carreer path
- Archive your source codes (because your software forge or webpage can disappear)

Software forge (1)

An online server and/or website offering code/software development and management functionality

- versioning
- collaborative work and planning
- · issue, feedback, bug reports, feature requests
- software release/publication
- · continuous integration
- possibility to get a publication identification like a DOI⁵
- · etc.

⁵eventually externally with Zenodo [☑], c.f. later

Software forge (2)

Examples of software forge

- gitlab: free and open-source git forge hosting software (different hosts are available: in the academic world⁶ or abroad⁷)
- github : very popular git forge with gratis and commercial solutions to host development projects (maybe more simple to reach outside the french academic community)
- · other: bitbucket ☑

Discontinued forges: gitorious, Google code, Inria Gforge (It happens!)

 $^{^6 {\}rm e.g.}\ https://plmlab.math.cnrs.fr, https://gitlab.inria.fr, {\rm etc.}$

⁷e.g. https://gitlab.com

⁸but owned by Microsoft

Archive your code (publication \neq archiving)

- What happens if your software forge (or the webpage where you host your code) disappear?
- The Software Heritage initiative [™]
 - "Our ambition is to collect, preserve, and share all software that is publicly available
 in source code form. On this foundation, a wealth of applications can be built,
 ranging from cultural heritage to industry and research."
 - Simple deposit procedure from a software forge⁹

⁹See https://archive.softwareheritage.org/save/

Get a DOI for your code with Zenodo

- a DOI¹⁰ to facilitate your software identification and citation (e.g. in publication using it)
- Upload your codes to Zenodo[™] and get a unique DOI for the current version (possible integration with github to directly generate identification for the different versions of your code)
- · Possible to identify codes, datasets, creative contents
- · More at https://help.zenodo.org/features/ and in the FAQ [™]

¹⁰ Digital Object Identifier ¹⁷

Test your functions

Implement automatic tests with testthat (1)

Good practice:

- write tests when you code functions (and not after)
- as soon as you create/modify a function, verify if tests are passing

Implement automatic tests with testthat (2)

```
To enable testthat in your package: usethis::use_testthat()<sup>C</sup>
To create a test case: usethis::use_test()<sup>C</sup>
(e.g. usethis::use_test("feature1"))
```

· Test file example:

```
test_that("multiplication works", {
    res <- my_multiplication(2, 2)
    expect_equal(res, 4)
})</pre>
```

tests sub-directory structure after initialization

```
tests
```

- +-- testthat
- +-- testthat.R

tests sub-directory structure after creating a test

tests

- +-- testthat
- +-- test-feature1.R
- +-- testthat.R

Implement automatic tests with testthat (3)

- Tests use expect_XX() functions to verify conditions of any type on any R expression
- Unit tests (i.e. test regarding a single function/functionality) can be grouped into test_that("id", {}) chunks
- You have to enumerate and write yourself all test cases
- More details at https://r-pkgs.org/tests.html and testthat exhaustive tour

test

Verify that the test you write are passing (e.g. your code is doing what you want)

- · devtools::test() ♂
- in Rstudio interface (Build panel More Test package¹¹)
- Note: tests will be run during package check

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¹¹keyboard shortcut: CTRL + SHIFT + T

Sharing (your code) is caring

Publish and distribute your package

- Others can use your work, collaborate with you to improve it (collaborative development)
- Many repositories: the CRAN[™] (official), bioconductor[™] (bioinformatics-oriented package repository)
- the remotes package (exported by devtools) can be used to install packages stored almost anywhere on the Internet (CRAN, bioconductor, git forges, etc.) or locally

CRAN

- Strict policy to accept a package [™] (READ IT!)
- Pipeline
 - 1. devtools::build() [♂] (or R CMD build)
 - 2. $devtools::check()^{\columnwidth{\columnwidth{\mathcal{C}}}}$ (or R CMD check --as-cran)
 - 3. upload it 12 to https://cran.r-project.org/submit.html
- devtools::release()[™] can help you to prepare the release (i.e. the version of your package that will be publish)

¹²in bundle state

Reverse dependencies

- Important: if you are releasing a new version of existing package, it is your responsibility to check that it does not break downstream dependencies¹³ (i.e. all packages that list your package in the Depends, Imports, Suggests or LinkingTo fields)
- · usethis::use_revdep() $^{\mbox{\tiny CP}}$ to enable the revdepcheck package $^{\mbox{\tiny CP}}$ that can help you in that task

¹³called "reverse dependencies"

- · versioning system: see the official website d and the book d
 - manage evolution of your code
 - branch-base system for production/development code cohabitation
 - · decentralized system: if you lose your remote, you do not lose the project history
 - easy to distribute (with git clone) and to move from remote to remote
- Command line tool or possible to manage everything from R/Rstudio:
 - · usethis::use_git() [™] to initialize a repository in your project
 - Git panel in Rstudio to manage your local repository and interact with remote (ssh key generation, etc.)
- More detail at https://r-pkgs.org/git.html

Distribute your package on a git repository

To install packages hosted on:

```
· any git forge: remotes::install_git() ♂
```

Possibility to specify the branch, the sub-directory where to find the package, etc.

```
remotes::install_github("RcppCore/Rcpp")

remotes::install_git(
    "https://github.com/getkeops/keops",
    subdir = "rkeops", branch = "dev", args="--recursive"
)
```

Organize your package project

- Package root directory = Rstudio project/git repository root directory (default behavior when using usethis::create_package() or Rstudio new project package)
- The package root directory is a sub-directory of the Rstudio project/git repository
 - you can specify the path to your package directory to devtools functions
 - Rstudio project setup: Tools Project Options Build tools Package directory

Advanced documentation

Writing a "vignette"

- A document¹⁴ presenting/detailing your package (or a functionality in your package), included in the package (and visible on CRAN)
- Written in a markup language: Rmarkdown[™] to integrate R code chunks, or LaTeX or Markdown
- To create a vignette: usethis::use_vignette("my-vignette")
- \cdot Rendering (in pdf/html/etc.) with the package knitr $^{\c C}$

¹⁴See https://r-pkgs.org/vignettes.html

¹⁵See also this cheat sheet ☑

Create a website

- Create and build a standardized website for your package with pkgdown 2716
- · Hostable on Github or Gitlab pages, or on your own webpage
- To create the website template: usethis::use_pkgdown()
- To build the website 17 (e.g. generate the HTML source): pkgdown::build_site()
- More details in the pkgdown vignette [™]

¹⁶See also https://github.com/r-lib/pkgdown

¹⁷README.md become the homepage, man documentation are used to generate function references, and vignettes are rendered into articles

Continuous Integration

- · Automate package testing and checking when you modify it
- Generally associated with a software forge
- · See usethis::use_gitlab_ci() or usethis::use_github_actions() or use_github_actions() or use_github_actions(
- You define a set of actions (e.g. tests and checks) that are run after each commit, or before any pull/merge request (configurable)

Non R code

interfacing language

Rcpp: Seamless R and C++ Integration

- · See the Rcpp webpage [™] and the introduction vignette [™]
- C++ API to use R types and R like functions¹⁸ in C++
- Automatic export of C++ functions to R¹⁹ in particular when creating/building a package
- Expose C++ functions and classes to R²⁰
- Conversion from C++ to R and back²¹

¹⁸See the "Rcpp-sugar" vignette [☑]

¹⁹See the "Rcpp-attributes" vignette ¹⁹

²⁰See the "Rcpp-modules" vignette [☑]

²¹See the "Rcpp-extending" vignette ^C

Rcpp: compilation on the fly

In convolve.cpp file:

```
#include <Rcpp.h>
using namespace Rcpp:
// [[Rcpp::export]]
NumericVector convolveCpp(
    NumericVector a. NumericVector b
    int na = a.size(), nb = b.size();
    int nab = na + nb - 1;
    NumericVector xab(nab);
    for (int i = 0: i < na: i++)</pre>
        for (int j = 0; j < nb; j++)
            xab[i + j] += a[i] * b[j];
    return xab;
```

Compilation on the fly in R:

```
sourceCpp("convolve.cpp")
convolveCpp(x, y)
```

Rcpp in a package (1)

· Create a Rcpp-based package template:

```
Rcpp::Rcpp.package.skeleton("NewPackage", attributes = TRUE)
```

- All C++ codes should be in the src sub-directory
- Add the comment // [[Rcpp::export]] before every C++ functions that should be exported to R
- · Add LinkingTo: Rcpp in DESCRIPTION file

Rcpp in a package (2)

- To generate the C++ to R wrappers: devtools::load_all()²² or devtools::build() will call Rcpp::compileAttributes()²³
- The files src/RcppExports.cpp and R/RcppExports.R are automatically created (or updated) and contain the code necessary to expose your C++ functions in R
- You C++ code will be compiled during your package installation

²²Reminder: CTRL + SHIFT + L

²³or you can call it yourself

Rcpp in a package (3)

- Compatible with roxygen2 doc generation
- Rcpp::compileAttributes()
 converts //' C++ doc comment
 chunks to #' roxygen2 doc comment
 chunks in the R/RcppExports.R file

```
#include <Rcpp.h>
using namespace Rcpp;

//' Do something
//' @author someone
//' @description
//' This function does something
//'
//' @param x An integer vector
//' @export
// [[Rcpp::export]]
void my_fun(IntegerVector a) {
    // do something...
}
```

The Rcpp ecosystem (1)

• RcppEigen[©]: 'Rcpp' Integration for the Eigen[©] Templated Linear Algebra Library

- RcppArmadillo[©]: 'Rcpp' Integration for the Armadillo[©] Templated Linear Algebra Library
- RcppGSL . Rcpp Integration for GNU GSL Vectors and Matrices
- BH[©]: Boost C++ Header Files ("a set of libraries providing support for tasks and structures such as linear algebra, pseudo-random number generation, multi-threading, image processing, regular expressions, and unit testing")
- · and more...

The Rcpp ecosystem (2)

How to use the previous C++ libraries in your package?

- Install the corresponding R package (with install.packages("<pkg>"))
- 2. Add LinkingTo: <pkg> in your DESCRIPTION file
- 3. Add the comment // Rcpp::depends(<pkg>)]] when including the corresponding library in your C++ code, e.g.:

```
#include <RcppArmadillo.h>
// Rcpp::depends(RcppArmadillo)]]
```

4. Use the C++ corresponding library in a standard way in your C++ code

reticulate: R Interface to Python

CRAN page[♂] and webpage ♂

Calling Python from R (dedicated vignette[™])

```
library(reticulate)
scipy <- import("scipy")
scipy$amin(c(1,3,5,7))</pre>
```

- · Conversion from R to Python matrix/array (dedicated vignette ♂)
- Python code chunks in Rmarkdown (dedicated vignette ^[2]])

Managing Python from R

- Python Version Configuration (dedicated vignette [□] and help page [□])
- Use virtual environment with reticulate::use_virtualenv() and reticulate::use_condaenv()

Using Python code in an R package

- Using reticulate in a R package (dedicated vignette [™])
- Configuring Python dependencies of your R package (dedicated vignette [™])

Control your R environment

renv

https://rstudio.github.io/renv/articles/renv.html

packrat

```
https://github.com/rstudio/packrat/
https://rstudio.github.io/packrat/
```

Configuring R

- References: here and here
- Configure where you install packages and from where you load packages (i.e. in which directory on your system)
- · Setup a default CRAN mirror for package installation
- Define default R objects, functions that will be available without additional file sourcing
- Modify R global options (see the functions options() and getOption() to check R global options)

.Renviron: configure the environment where R is run (1)

.Renviron = a file defining environment variables (as in bash) with the following syntax (!!not R code!!):

```
Key1=value1
Key2=value2
```

To edit your .Renviron file, you can use usethis::edit_r_environ().

.Renviron: configure the environment where R is run (2)

- To modify the directory where packages are installed²⁴ and loaded from²⁵: you can set²⁶ R_LIBS_USER=/path/to/my/lib/dir (useful to have project-specific package installation²⁷)
- Define environment variables (e.g. MYVAR=5) that will be available in R (with Sys.getenv("MYVAR")) or have an effect an your R code behavior

²⁴by install.package(), devtools::install(), remotes::install_from_xxx()

²⁵ by library() or require()

²⁶default value is 'R_LIBS_USER=~/R/%p/%v

²⁷to avoid package version conflict between project

Where storing the .Renviron file

R tries to use an .Renviron file in the following order:

- 1. in the working directory where R is started (if existing), e.g. in your RStudio project root directory
- 2. in your home directory (if existing)

Note: You can modify this behavior by setting (outside of R/RStudio²⁸) the following environment variable: R_ENVIRON_USER=/path/to/my/.Renviron

Anyway: R has a global Renviron.site file that is read first. Using your own .Renviron file allows you to modify the default environment defined in this file.

²⁸as in your bash environment

.Rprofile: configure and modify your R session

- .Rprofile = an R source file that will be run at R startup (after .Renviron was read)
- · What for?
 - define your own default R objects/functions
 - · write a startup message
 - · modify R global options
 - etc.

To edit your . Rprofile file, you can use usethis::edit $_r$ profile().

.Rprofile: an example

```
# setup a default CRAN repository
options(repos = c(CRAN = "https://cran.rstudio.org"))

# modify an option only in interactive mode
if(interactive()) {
    options(width = 120)
}
```

Note: interactive mode = as in R console²⁹ (in RStudio or in a terminal)

²⁹versus script mode (like scripts run by Rscript)

Where storing the .Rprofile file

R tries to use an .profile file in the following order:

- 1. in the working directory where R is started (if existing), e.g. in your RStudio project root directory
- 2. in your home directory (if existing)

Note: You can modify this behavior by setting (outside of R/RStudio³⁰) the following environment variable: R_PROFILE_USER=/path/to/my/.Renviron

Anyway: R has a global Rprofile.site file that is read first and using your own .Rprofile file allows you to modify the default R session defined in this file.

³⁰as in your bash environment

.Renviron/.Rprofile and reproducibility

Attention: you should be careful that your code is usable without your .Renviron and .Rprofile files

- .Renviron and .Rprofile files are personal files, another user may configure its environment differently
- Example: charging packages or modifying (global or packages) options that have an impact on output values³¹ in your .Rprofile file may affect the reproducibility of your code (i.e. the results can be different or you code can be broken without your .Rprofile file)

³¹e.g. options(stringsAsFactors = FALSE)

The end

