

Developing an R package: a tutorial

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Note for the reader

Many **hyperlinks**[↗] are directly embedded in the slide contents.

Requirements (1)

- **R** [↗] (the latest version if possible¹, **4.1.0** since 2021-05-18)
- You can use the R command line combined with any text editor, but we recommend to use an R-oriented IDE² like **Rstudio** [↗] or **RKward** [↗]
- All content presented here have been tested on a Linux environment but should work on **any OS**
- **Note for Windows users:** you can update R from within R with the **installr** [↗] package and you will need to install Rtools [↗] to enable all R development functionality

¹Keep your software up-to-date! If you need an older version of R for a specific project, use appropriate tools like containers, it should be an exception not a habit.

²Integrated Development Environment [↗]

Requirements (2)

- See the script `install_requirements.R`[↗] to install the packages that will be used in the tutorial
- To (re)generate the slides, see the scripts³ `.setup.R`[↗] to install the requirement, `.build.R`[↗] to build the `pdf` slides

³or the attached `Makefile`[↗] if you are comfortable with using `make`

References

- **Official R documentation:** *Writing R Extensions*[↗]
- Karl Broman[↗] **tutorial:** *R package primer* (web version[↗] and sources[↗])
- Hadley Wickham[↗] and Jenny Bryan[↗] **book:** *R packages* (web version[↗] and sources[↗])
- Hilary Parker[↗] **tutorial** on writing R packages[↗]
- R studio **cheatsheets**[↗] on package development[↗] and Rstudio IDE[↗]

What is an R package?

- a library containing a set of R functions (and possibly more) implementing functionality not available in default R functions⁴
- a standardized way to distribute R codes (for other users)

⁴or reimplementing existing functionality in a different way

Where can I find R packages?

- the [CRAN](#)[↗] (Comprehensive R Archive Network): official repository for R packages

```
install.packages("devtools")
```

- [bioconductor](#)[↗]: bioinformatics-oriented package repository
- any git forge: github, gitlab
- on your colleagues' computers⁵

⁵if they develop in R

Why R packages?

- The **best way** to write and distribute **R code** with **documentation**, **examples**, **tests**, etc.
- A good practice⁶ when coding in R:
 - your project is structured (code, data, doc), easier to use and re-use
 - documentation is essential (including for your future self)
 - your code is standardized, you can check it and test your functions
 - easy management of dependencies
 - etc.

⁶even for codes you don't plan to publish/distribute

How to write an R package?

A wide variety of tools to help you:

- Rstudio IDE built-in development features
- R base built-in tools: `build` (`R CMD build`), `check` (`R CMD check`)
- Some packages to develop packages:
 - `usethis`: to automate package and project setup
 - `devtools`: complete collection of development tools
 - `roxygen2`: to document your code and generate help pages
 - `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)

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