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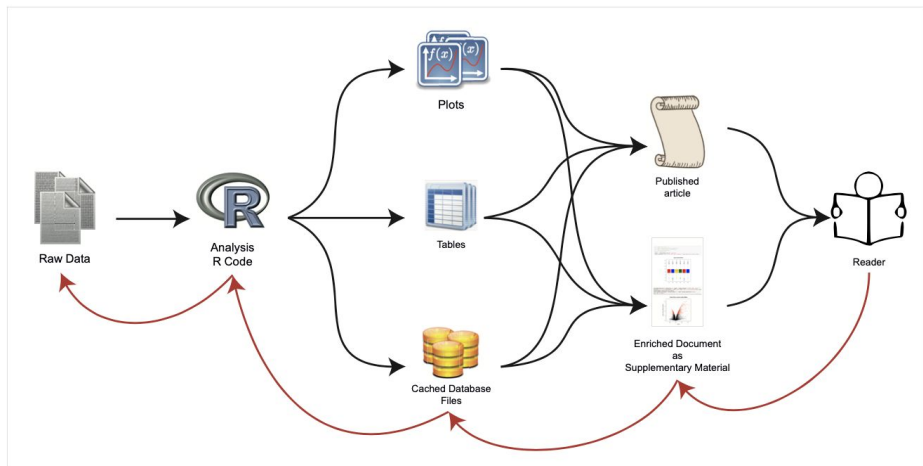
Tools and best practices for R package development

Department of
Statistical Sciences

31/05/2023



Allow people to reproduce your findings



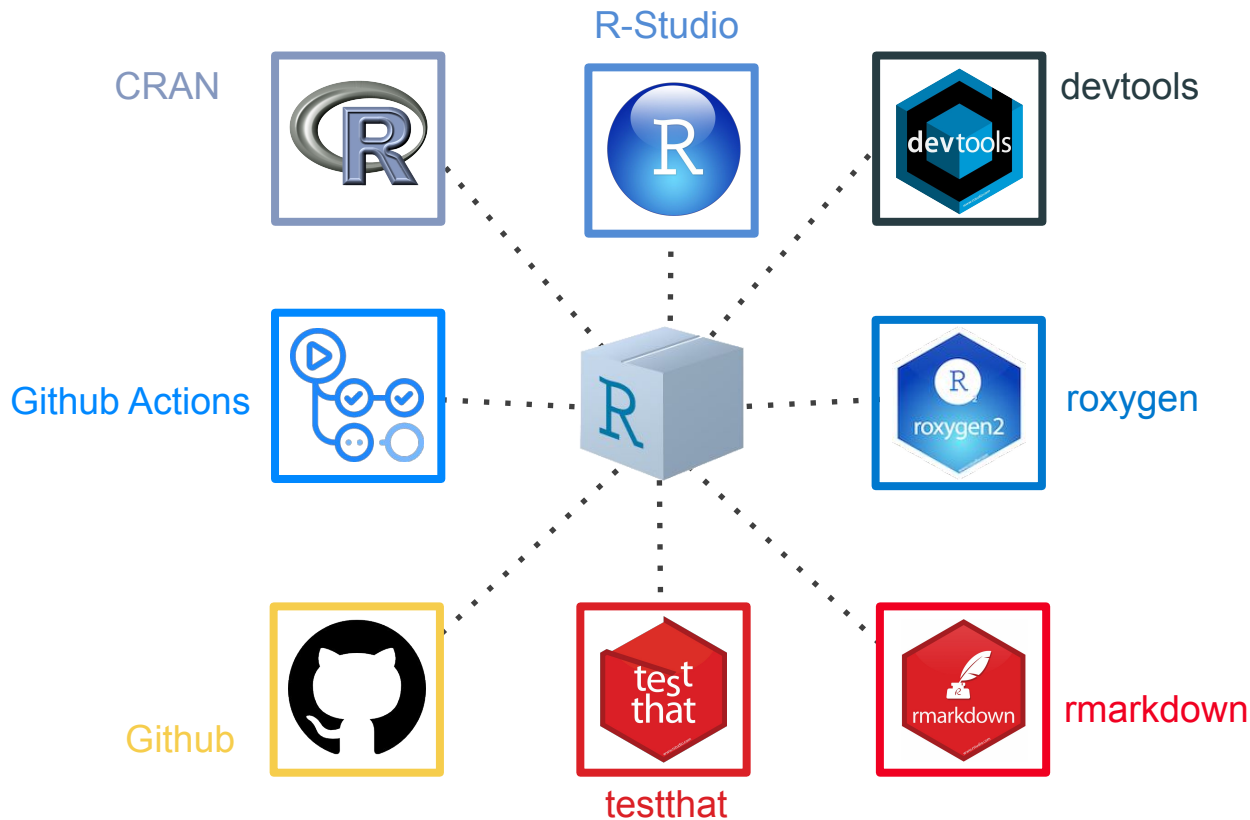
Russo F, Righelli D, Angelini C

Advantages and Limits in the Adoption of Reproducible Research and R-Tools for the Analysis of Omic Data

Lecture Notes in Computer Science 2016

- The scientific community needs to reproduce newly discovered insights
- Together with the article share
 - a. code and data
 - b. possibly an analysis report
- Main idea is
 - a. from raw data + code
 - b. produce tables/plots/processed data
 - c. write the article
 - d. share a supplementary report with the analysis
 - e. the “article reader” is able to entirely reproduce the analysis and so the described facts in the article
- As statistical model developers we want people to use our models
 - a. producing R packages

Some helpful components for reproducibility



Don't judge yourself and your code

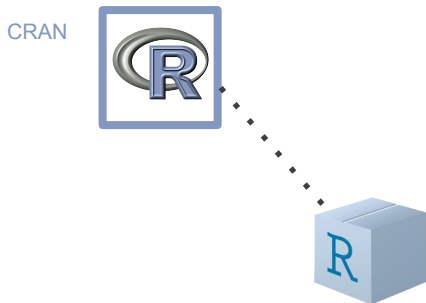


- Installing guidelines for R-devel:

- Linux:
 - <https://tinyurl.com/mspa6y56>
 - <https://tinyurl.com/4evytvny>
- OSX:
 - <https://tinyurl.com/3cncctct6>
 - use rswitch app (<https://rud.is/rswitch/>)
- Windows:
 - <https://tinyurl.com/bduh7k62>

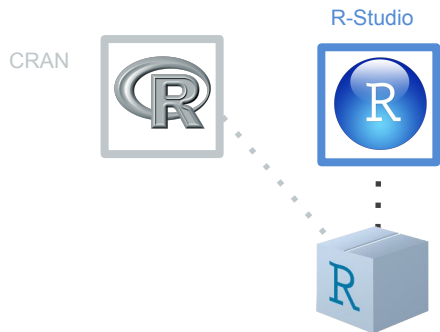
- Having some code is better than no code
- You can always improve/optimize it in a second moment
- Ask questions to your community, don't be shy!
 - colleagues, supervisors, friends, ecc
- Use the R development version
 - your R version dependency is not already old
 - dependencies bugs and problems
 - easier to solve them during your coding
 - you are ready to go on CRAN
- Your work is never “done”
 - continuous bug solving
 - broken dependencies along the package life
 - tip: keep your dependencies set small

A repository to collect [almost] them all



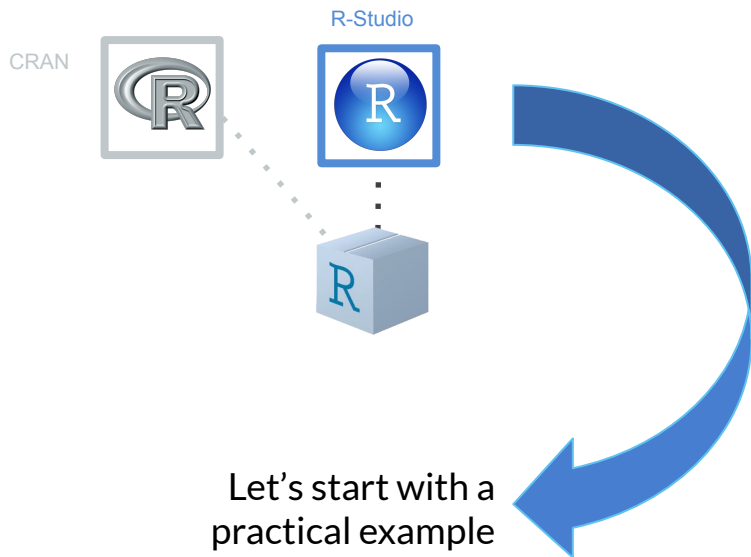
- The Comprehensive R Archive Network (CRAN)
- Official R packages repository
 - requires multiple check passing to enable a package to be submitted/accepted.
- It hosts one of the most relevant set of R packages:
 - tidyverse: borns from a group of people who contributed to let CRAN be what it is nowadays
 - organized in macro thematics
 - systematic check of packages
 - optimization of the checking packages with more informative outputs
 - constant improvement
- Writing R package guidelines:
 - <https://tinyurl.com/2p9eef9h>
 - good to know how to structure your code before starting to do it
- Submitting process guidelines:
 - <http://r-pkgs.had.co.nz/release.html>

A development environment dedicated to R



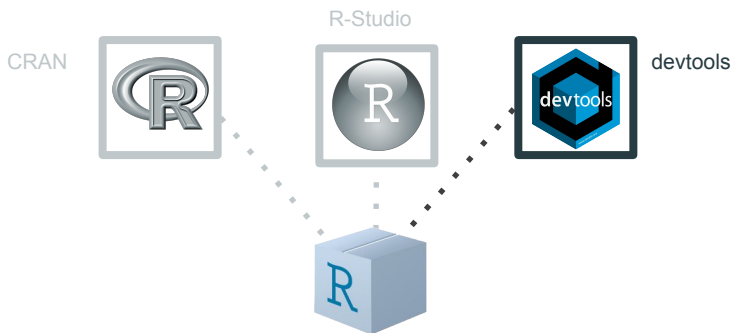
- Integrated Development Environment (IDE) for R
 - It also recognizes other programming languages
 - Python, SQL, Bash, Cpp, JSON, YAML, JavaScript, CSS
- Not the best IDE, but:
 - Easy creation of R package skeleton
 - Easy package documentation support (roxygen2)
 - rmarkdown support
 - code versioning support (git/svn)
 - integrated plot visualization
- Is organized in four different areas
 - code editing
 - visualization panes
 - terminal (shell, R)
 - environment (loaded data variables)
- Daily developed, upgraded and released
- More @ <https://www.rstudio.com>

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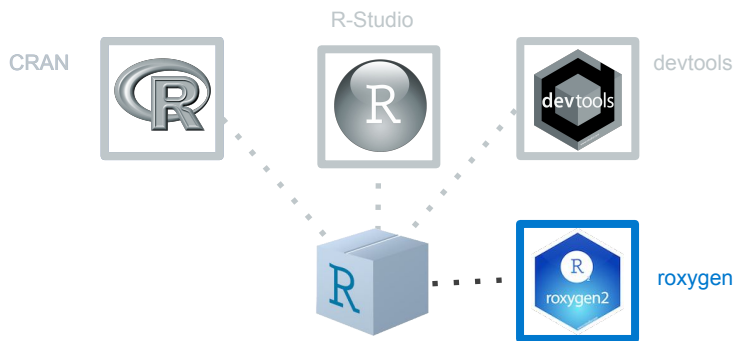
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A swiss-knife for package development



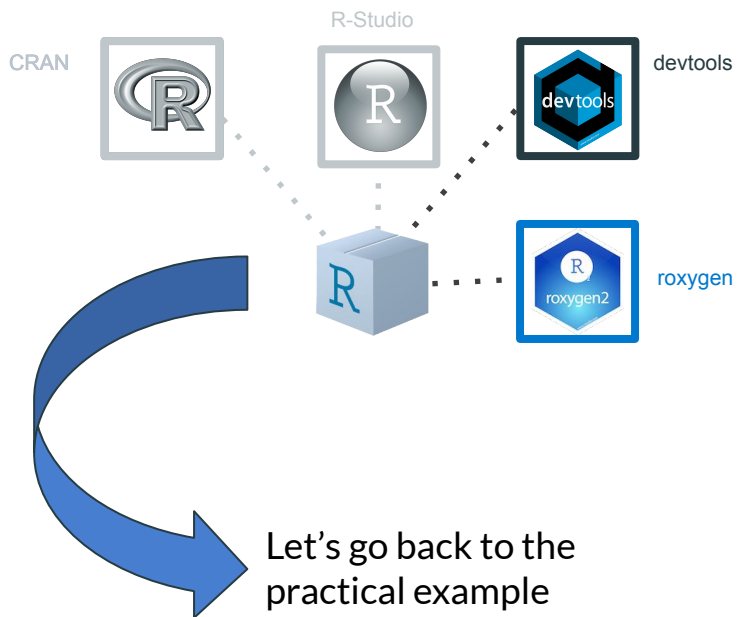
- Install and use the *devtools* package
- It gives some functionalities to speed up your development
- process:
 - `load_all()`
 - `install()`
 - `build()`
 - `check()`
- Once written the documentation of your function(s):
 - `document()`
- Once written the unit tests of your function(s):
 - `test()`
- Useful also to install packages from github (i.e.):
 - `install_github("hadley/dplyr")`
- <https://devtools.r-lib.org/>

Document your work



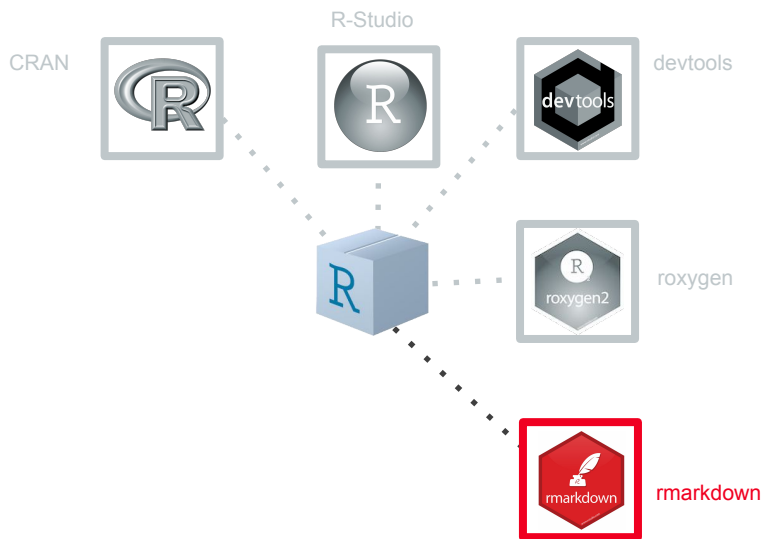
- Helps users to understand what your functions do
- Helps yourself to remember what your functions do
- *roxygen2* interprets special characters in the source code to write the function documentation
 - Use special characters `#'` to write your documentation in the code.
 - Use special keywords using `@` to tell roxygen how to handle the information
- R-studio can write a basic skeleton to document a function.
- roxygen+devtools:
 - Auto generate your documentation files!
- <https://tinyurl.com/yjj82xc>

Document your work



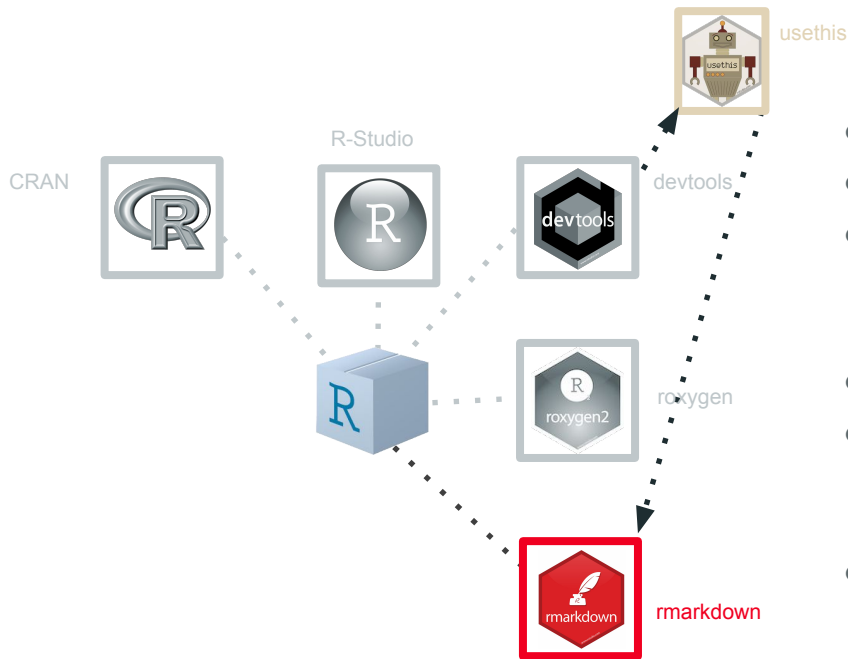
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Let people know how to use your package



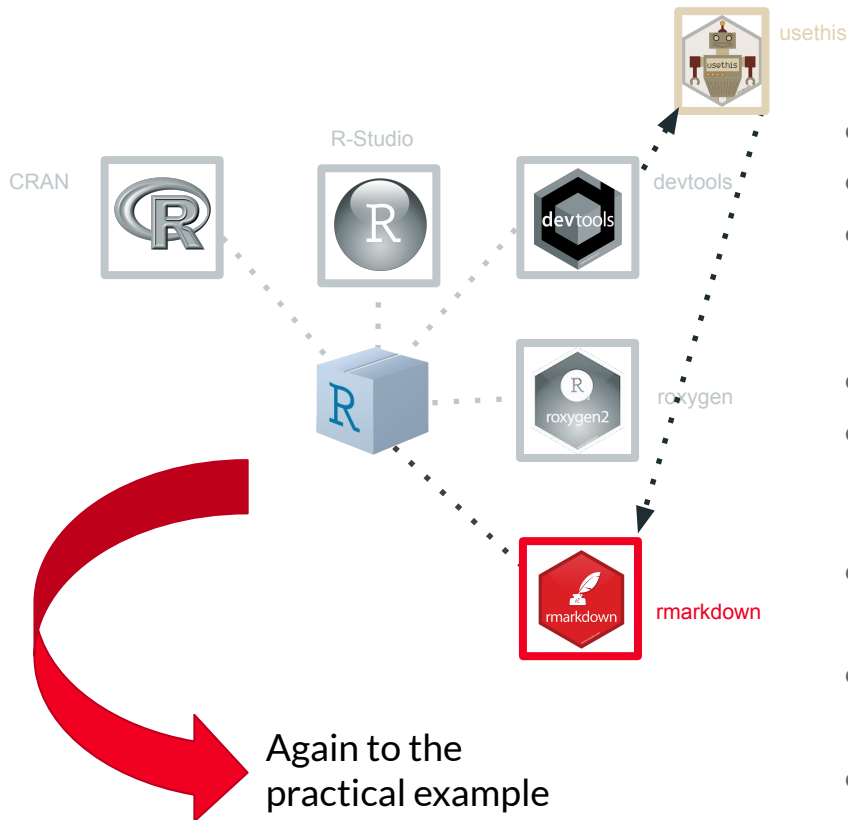
- *rmarkdown* is the markup language for R
- you can mix natural language with code chunks
- use it to write an usage example of the functions in you package
 - save the file in the vignettes folder
- helps other people to better understand your work
 - reproducibility and reusability
 - shareability
- otherwise you'll never remember how to use your own package in a couple of years from now
- tip: add a `sessionInfo()` code chunk at the end of your vignette
- <https://bookdown.org/yihui/rmarkdown/>

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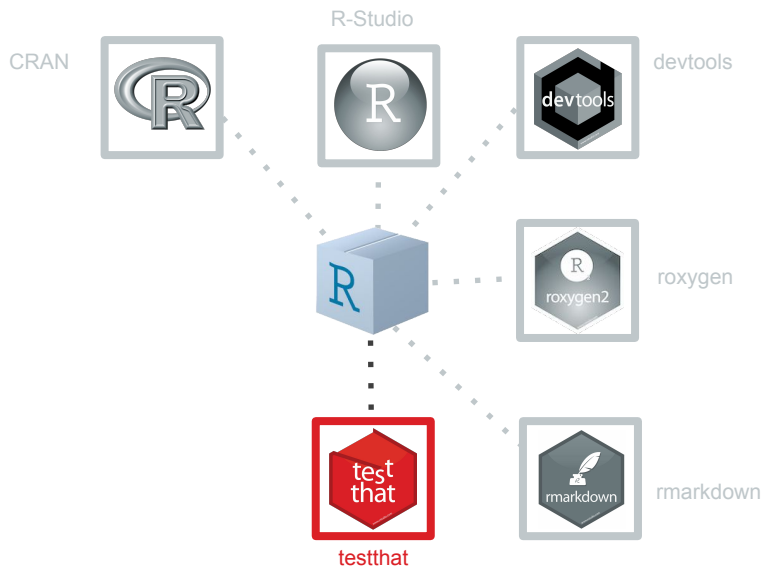
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Test the code for yourself

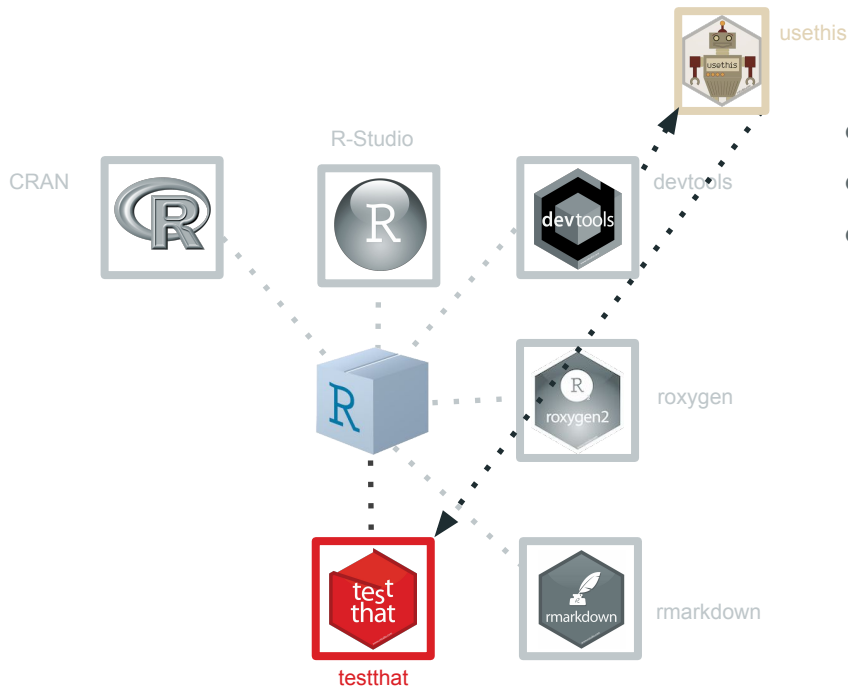


- *testthat* helps to create unit tests for your functions
- gives a series of functions to verify the input/output
- Ref: <https://testthat.r-lib.org/>

Full	Short cut
<code>expect_that(x, is_true())</code>	<code>expect_true(x)</code>
<code>expect_that(x, is_false())</code>	<code>expect_false(x)</code>
<code>expect_that(x, is_a(y))</code>	<code>expect_is(x, y)</code>
<code>expect_that(x, equals(y))</code>	<code>expect_equal(x, y)</code>
<code>expect_that(x, is_equivalent_to(y))</code>	<code>expect_equivalent(x, y)</code>
<code>expect_that(x, is_identical_to(y))</code>	<code>expect_identical(x, y)</code>
<code>expect_that(x, matches(y))</code>	<code>expect_matches(x, y)</code>
<code>expect_that(x, prints_text(y))</code>	<code>expect_output(x, y)</code>
<code>expect_that(x, shows_message(y))</code>	<code>expect_message(x, y)</code>
<code>expect_that(x, gives_warning(y))</code>	<code>expect_warning(x, y)</code>
<code>expect_that(x, throws_error(y))</code>	<code>expect_error(x, y)</code>

Table 1: Expectation shortcuts

Test the code for yourself

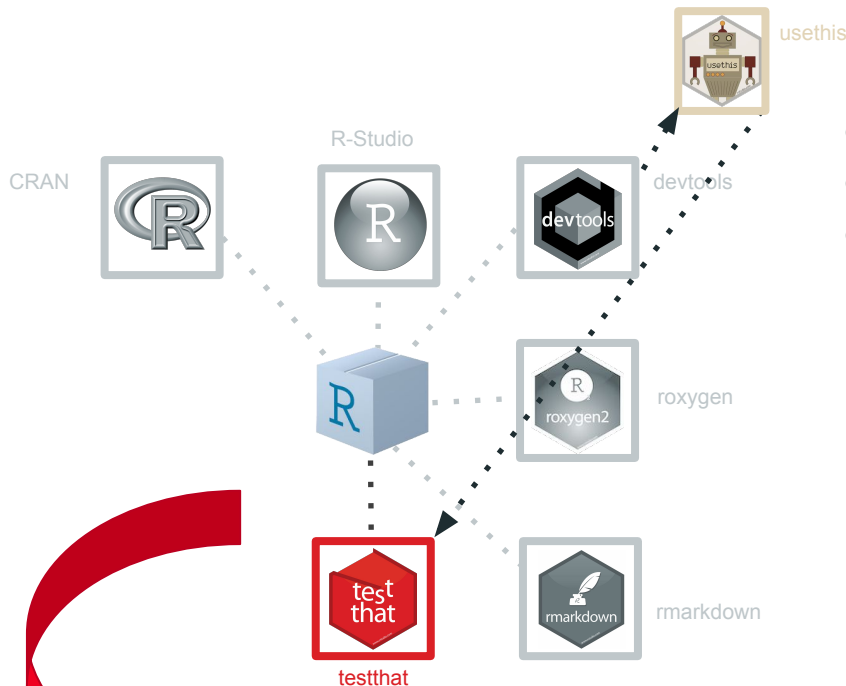


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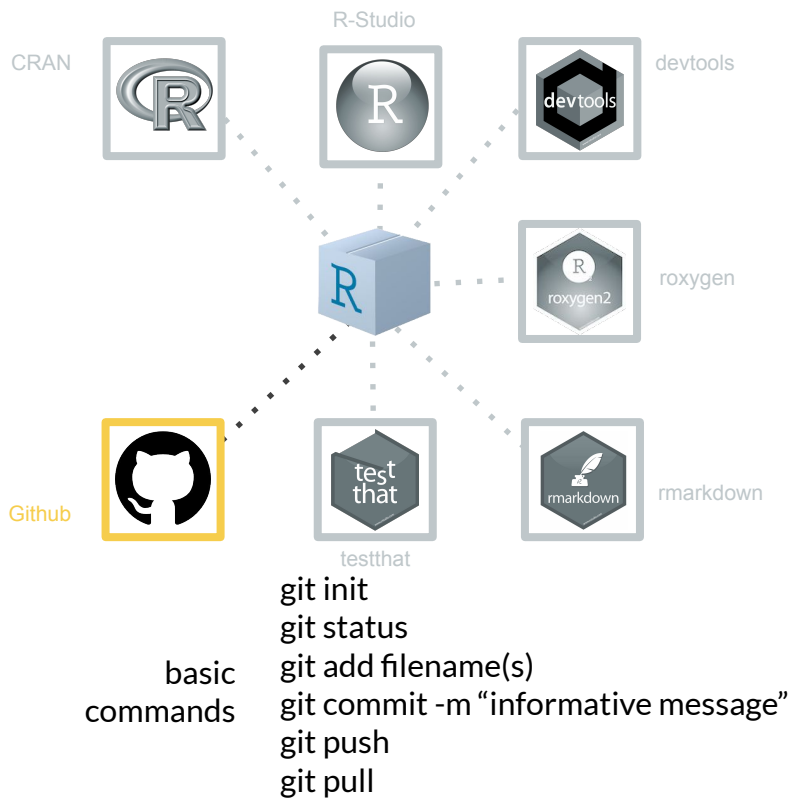
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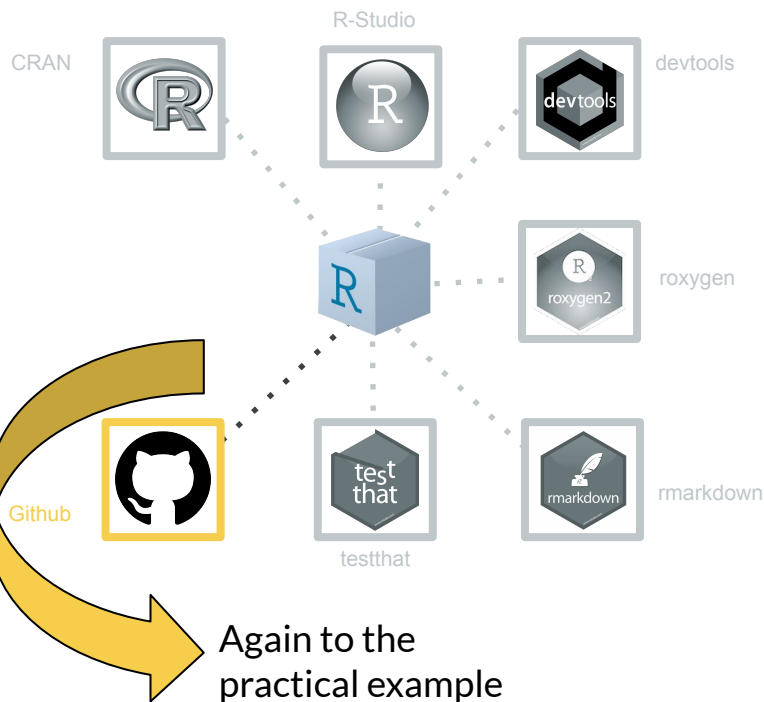
Again to the
practical example

Keep version of your changes



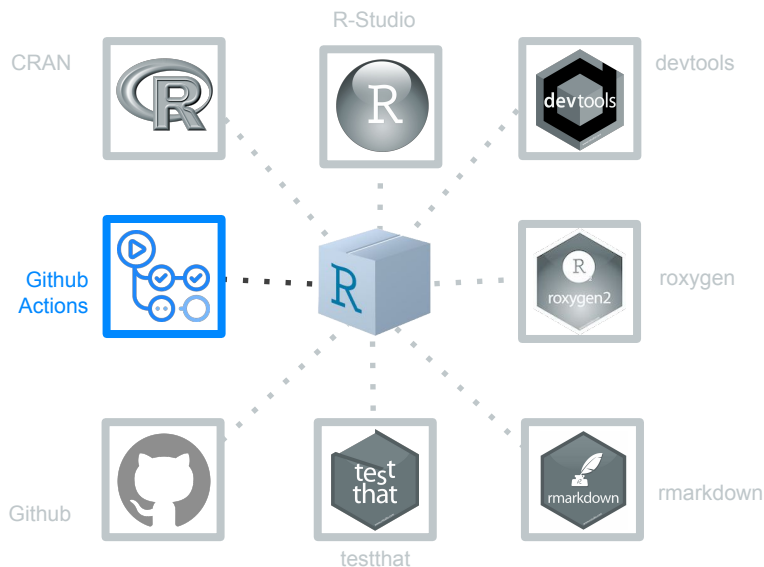
- **Github is not git!**
 - git is the code versioning software installed locally on your machine
 - github is an online (cloud) service for keeping track of your local git repositories
- Create a local git repository for each project you have
 - R package
 - your thesis
 - your CV
 - keep version control of each project individually
- Create a remote github repository for each local git repository you want to keep track on the cloud
 - then you can link local and remote git repositories
- github helps team-working
 - multiple users have access to the same remote repository
 - each of them can push/pull code on the same repository
 - git/github helps to keep track of changes/conflict in the code
- R-studio helps in creating and handling git repositories
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- Beginners tutorial: <https://tinyurl.com/4w563274>

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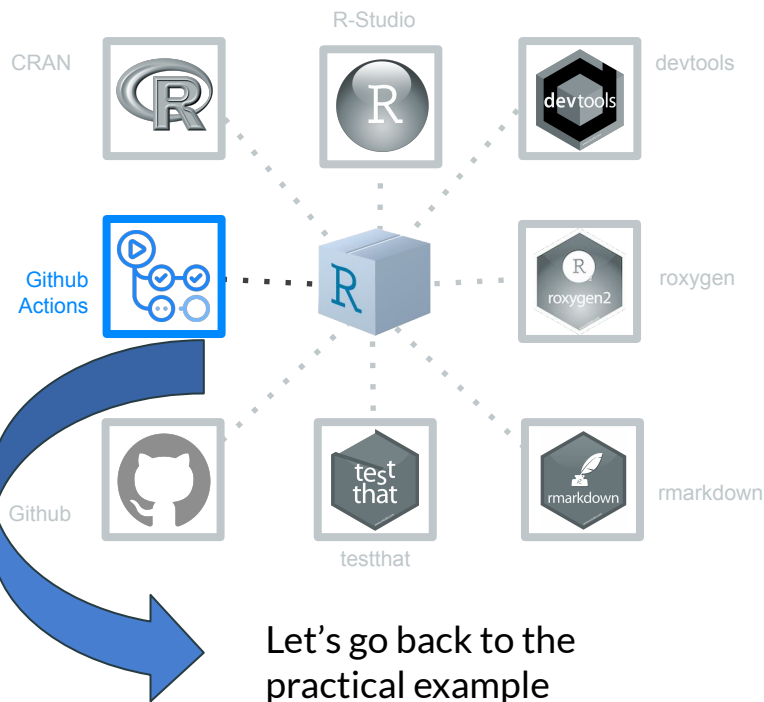
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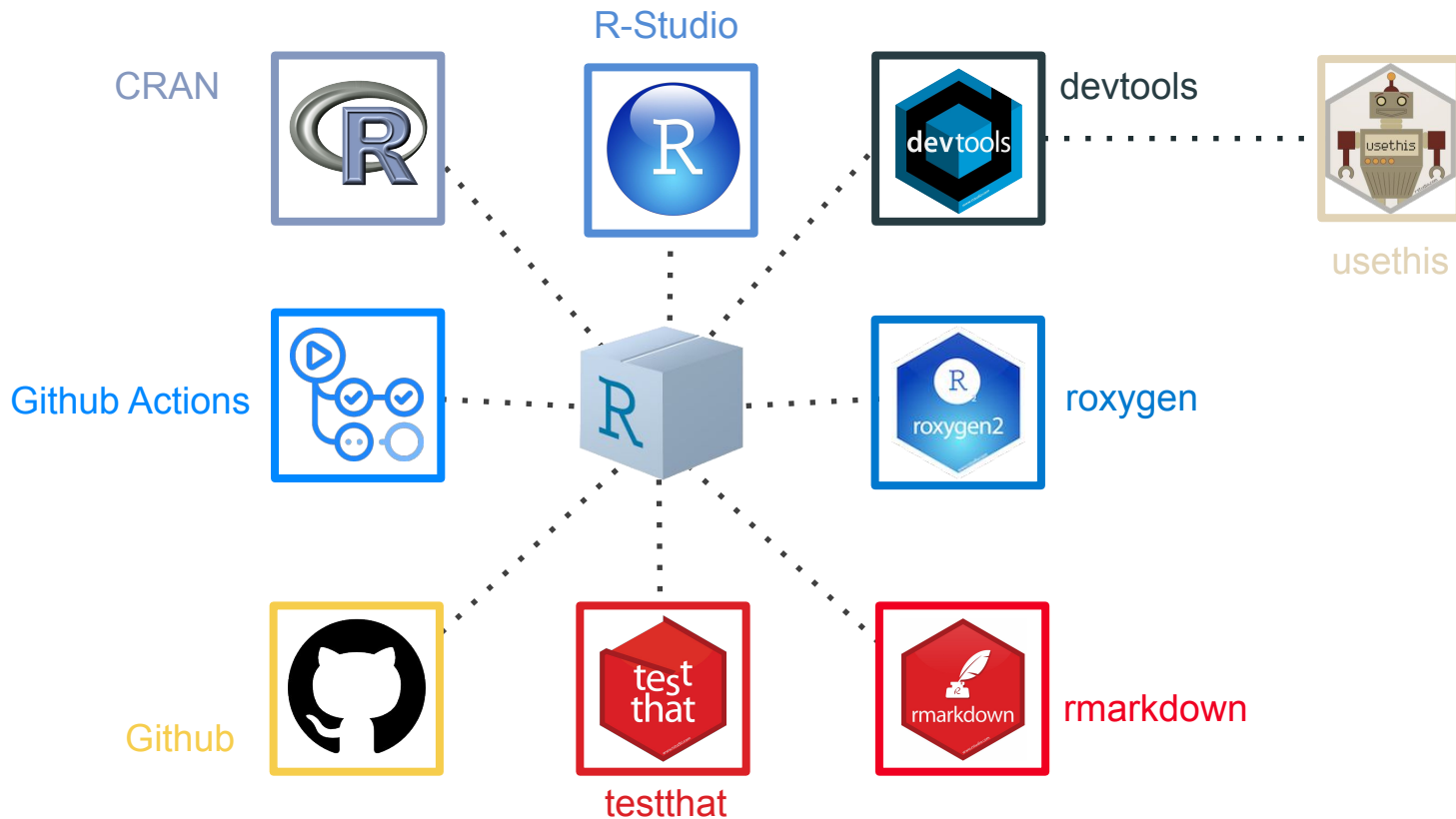
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 - optionally deploy you package to a specified platform
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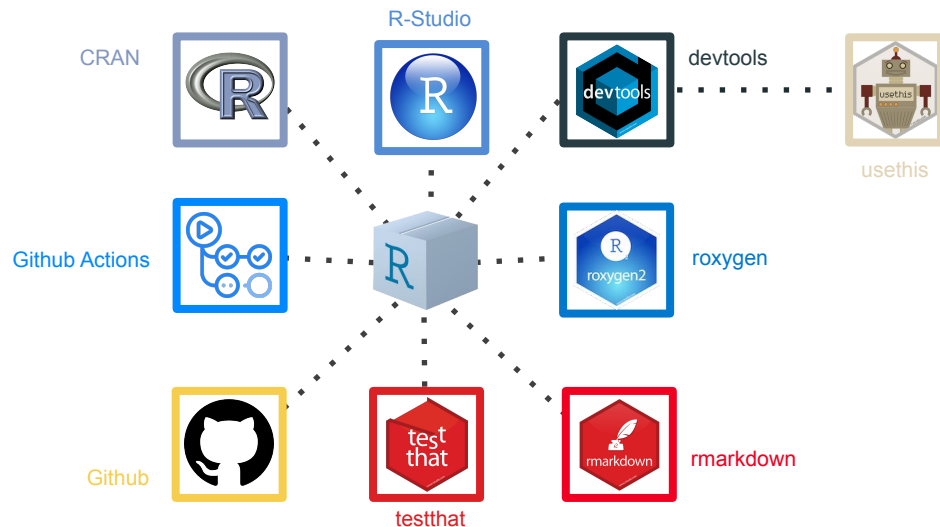


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Some helpful components for reproducibility bonus package



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