

# R Packages and Wrap-Up

Computational Statistics

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# **Today**

# Distributing and Organizing Code

Workshop in creating an R package  $\,$ 

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# **Oral Examination Prep (Afternoon Session)**

What to think of during examination

# Organizing Code as an R Package

# **Organizing Code**

# Components

- Code for experiments
- Source code for functions (which we should be able to reuse)
- Tests
- Rcpp code
- Data

There are many ways to organize this. Which one to choose?

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- Tests
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# R Package

One way is to make an R package, makes it easy to

- connect to C++ code through Rcpp,
- set up automatic testing,
- document your code, and
- declare dependencies (other packages, R version).

# R Packages

Different approaches, but we will follow **R Packages** (Wickham and Bryan 2023), which is based around the **devtools** package.

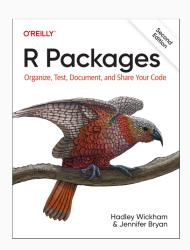


Figure 1: R Packages

### **Devtools**

Meta-package for various helpers that aid in developing R packages (and projects). First off, install and load **devtools**:

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

This loads other packages that will be useful for setting up your package, most importantly the **usethis** package.



## Rosenbrock Package

Let's build a simple package that solves the Rosenbrock optimization problem, i.e. find

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### What We Will Learn

Adding R functions to our package

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- Testing our code
- Interfacing with Rcpp
- Adding dependencies to other packages
- Licensing our package
- Documenting the code

### Create It

Call

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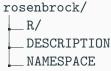
```
rosenbrock/
R/
DESCRIPTION
NAMESPACE
```

### Create It

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usethis::create_package("rosenbrock")
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Gives you a minimal package:



You may also have .Rbuildignore and .rosenbrock.Rproj depending on how you created the package.

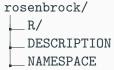
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You may also have .Rbuildignore and .rosenbrock.Rproj depending on how you created the package.

### Install It

Open up the package in your editor (R Studio $^a$ ).

devtools::install()

Voila, you have made an R package!

<sup>&</sup>lt;sup>a</sup>In which case it should alread be opened.

### R Code

### .R/

- All R code should live in .R-files in R/.
- These files should (almost) always contain only functions.
- Many ways to organize your files: one function per file, all functions of a certain S3 class in one file etc.

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- Many ways to organize your files: one function per file, all functions of a certain S3 class in one file etc.

Let's create a first file: R/objective.R. Use usethis::use\_r("objective") and insert this:

```
objective <- function(x, a = 1, b = 100) { (a - x[1])^2 + b * (x[2] - x[1]^2)^2}
```

We have created a first R file, but how do we use it? Two major options:

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Installs the package, like calling
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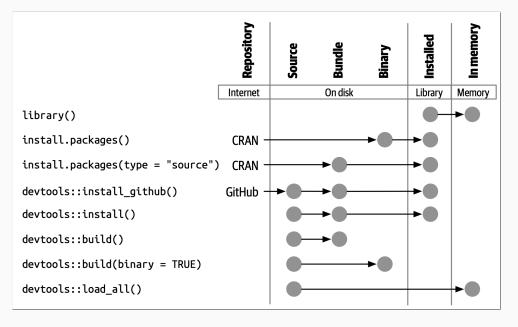
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Quick but not as robust.

# Try It

Try both options and see if you can call your newly defined function, objective().

 $<sup>{}^{</sup>a}\mathsf{Done}$  automatically in R Studio



**Figure 2:** The various states of a package and how to move between them.

# **Exporting Functions**

If you called devtools::load\_all() then everything is sourced and you can just call objective() directly.

But if you use devtools::install() and library(rosenbrock), the you would need to use rosenbrock:::objective(). The reason is that the function is not yet exported.

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Decides what functions you want exported. But right now it just contains a comment:

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### NAMESPACE

Decides what functions you want exported. But right now it just contains a comment:

```
# Generated by roxygen2: do not edit by hand
```

If you want to just export everything, you can remove this file and recreate it with this content:

```
exportPattern("^[[:alpha:]]+")
```

**roxygen2** is a package that helps with package documentation<sup>1</sup>, but it can also be used for handling the namespace.

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Go ahead and place this before your objective() definition. Then run devtools::document() to roxygenize your package.

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Now NAMESPACE will (should) contain this:

```
export(objective)
```

Reinstall the package and see if you can call objective() after loading it.

<sup>&</sup>lt;sup>1</sup>More on this later.

### **Tests**

### testthat

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This creates some new files and directories:

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Call use\_test("objective")<sup>2</sup> and insert this:

```
test_that("multiplication works", {
    # add a test using expect_equal()
})
```

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})
```

## **Check That Everything Works**

Run devtools::test(), and hopefully see:

```
[ FAIL O | WARN O | SKIP O | PASS 1 ]
```

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Now run devtools::check(). Is there a problem? Yes, let's fix it!

#### Metadata

The metadata for your package lives in DESCRIPTION. Right now it looks like this:

```
Package: rosenbrock
Title: What the Package Does (One Line, Title Case)
Version: 0.0.0.9000
Authors@R:
    person("First", "Last", , "first.last@example.com", role =
       c("aut", "cre"),
           comment = c(ORCID = "YOUR-ORCID-ID"))
Description: What the package does (one paragraph).
License: `use_mit_license()`, `use_gpl3_license()` or friends
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Encoding: UTF-8
Roxygen: list(markdown = TRUE)
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For now we'll leave most of these files alone, but let's fix one thing: the license

## Why Do You Need a License?

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So we need to pick a license: for now we'll pick the MIT license.<sup>3</sup>

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```
usethis::use_mit_license()
```

This will add new files to your package: LICENSE, LICENSE.md, and modify DESCRIPTION, in which you should see:

```
License: MIT + file LICENSE
```

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#### **Gradient**

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Let's say that we want to compute the gradient for the Rosenbrock function.

One way to do so is to use numerical differentiation through the **numDeriv** package:

```
gradient <- function(x, a = 1, b = 100) {
  numDeriv::grad(objective, x, a = a, b = b)
}</pre>
```

Now our package depends on numDeriv, so we need to add it to DESCRIPTION:

```
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In DESCRIPTION, you should now see this:

```
Imports:
   numDeriv
```

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We will rely on roxygen2. First, call usethis::use\_package\_doc() to set up a package doc file in R/rosenbrock-package.R.

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Then use usethis::use\_rcpp() to put the pieces in place:

```
rosenbrock/
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__slop-package.cpp/
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Then use usethis::use\_rcpp() to put the pieces in place:

```
rosenbrock/
L src/
L slop-package.cpp/
```

Now just need to run devtools::document() and devtools::load\_all() or devtools::install() and now your code is available (but not exported).

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Call your Rcpp function through an R wrapper:

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You can add roxygen2 comments in Rcpp code too:

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Saves you having to write and maintain an R function.

## **Documentation**

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# roxygen2

Primary purpose of the package. You write code is a special syntax and it converts it into manual files that R understands.

## **Types**

- Comments in code
- Manual (help files)
- Long-form articles (vignettes)

# roxygen2 Syntax

```
#' Function Title
# '
  Here you describe what the function does, possibly
#' using several lines.
# '
  Oparam x Explanation of argument x
# '
  Creturn Explanation of what the function returns
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```

#### Your Turn

Document objective() with roxygen2 syntax. No need for sensible documentation. Just make sure you have the bare minimum.

## **Documentation in This Course**

- Not making a package for CRAN, so lower standards.
- You don't need to document to benefit from building a package.
- But it's not a bad idea to do so anyway!

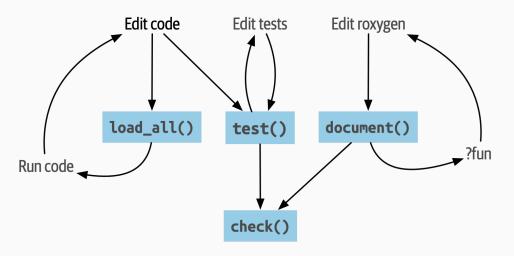


Figure 3: The whole game

# Projects

When you have a project, you typically need more things:

- scripts with simulations, etc, which produce output
- datasets stored in different formats
- notebooks (or latex sources)

These things do not naturally fit into a package framework.

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#### Two Choies of Structure

- 1. Just store these things directly into the package folder. Optionally, you can use .Rbuildignore to ignore these files when building the package.
- 2. Put your **package** into a **subdirectory** of your project. This cleanly separates the part of your project that contains reusable code (the package) and the part that is experiments and reports. But a little trickier to setup.

#### Rosenbrock

Continue building the **rosenbrock** package:

• Write a gradient descent (or stochastic gradient descent) implementation that minimizes the rosenbrock function.

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Continue building the rosenbrock package:

- Write a gradient descent (or stochastic gradient descent) implementation that minimizes the rosenbrock function.
- Write the code in Rcpp. If you want, you can first write it in R to see that everything is working, and then port it.
- Feel free to use generative AI to write the code.
- Export everything and document the package.

## An Assignment

Start trying to convert your work for one assignment into a package

 $\bullet$  Version control through git and github

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- Principled approaches to reproducibility (renv, containers)

# **Oral Examination Prep**

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#### **Examiners**

Me and Jonas Gyde Hermansen

It's possible that Niels will show up during one or two of the examinations.

## Remember the Five Points

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- Can you implement alternative solutions?
- Can the code be restructured e.g. by modularization, abstraction or object oriented programming to improve generality, extendability and readability?
- How does the implementation perform (benchmarking)?
- Where are the bottlenecks (profiling), and what can you do about them?

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- Use plots as much as possible
- Good to include math and code, but avoid overwhelming us.

### **Evaluation Criteria**

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Knowledge of fundamental algorithms for statistical computations and R packages that implement some of these algorithms or are useful for developing novel implementations.

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## Competence

Ability to select appropriate numerical algorithms for statistical computations and evaluate implementations in terms of correctness, robustness, accuracy and memory and speed efficiency.

# **Statistical Topics**

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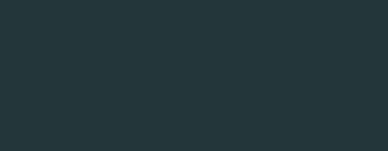
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## **Computational Topics**

- Debugging
- Profiling
- Benchmarking
- Debugging
- Writing performant code



Thank you (for real this time)!