

Introduction to Building R Packages

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Install R Packages

- ▶ Most common/popular repositories for R packages include:
 - ▶ CRAN: official repository maintained by many servers world-wide

```
install.packages("package")
```

- ▶ Bioconductor: specific packages for bioinformatics

```
source("https://bioconductor.org/biocLite.R")  
biocLite() # core packages  
biocLite("package")
```

- ▶ github: no review process

```
devtools::install_github()
```

- ▶ Some house-keeping tools

```
installed.packages() # check all installed packages  
update.packages() # update package  
remove.packages("package") # remove package
```

Load package

```
library(package)
require(package)           # no error if not installed

package::function()      # execute specific function only

detach("package", unload=TRUE) # unload package
```

Help files

```
?function
?package::function

# package overview
help(package = "packagename")

# vignette/tutorials
vignette(package = "packagename")
vignette("vignettename")
```

Why Building R Packages?

- ▶ platform-independent distribution of R code
 - ▶ alpha/beta versions on R-forge or github
 - ▶ finished projects on CRAN or Bioconductor
- ▶ archiving R code for a specific project and software documentation
- ▶ reproducible research: distribute data and software accompanying a publication
- ▶ maintenance of dependencies, and automated loading of required external code
- ▶ CRAN uses `R CMD check` to test package on various platforms; packages are tested daily

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A basic (but good) R package has the following structure:

DESCRIPTION what does the package? who can use it (license)?
who is responsible (maintainer)?

NAMESPACE which function should be seen by the user? which are
internal?

`R/` R functions

`man/` documentation, help files with syntax similar to \LaTeX

`data/` example data files

Additional (optional) files in R packages:

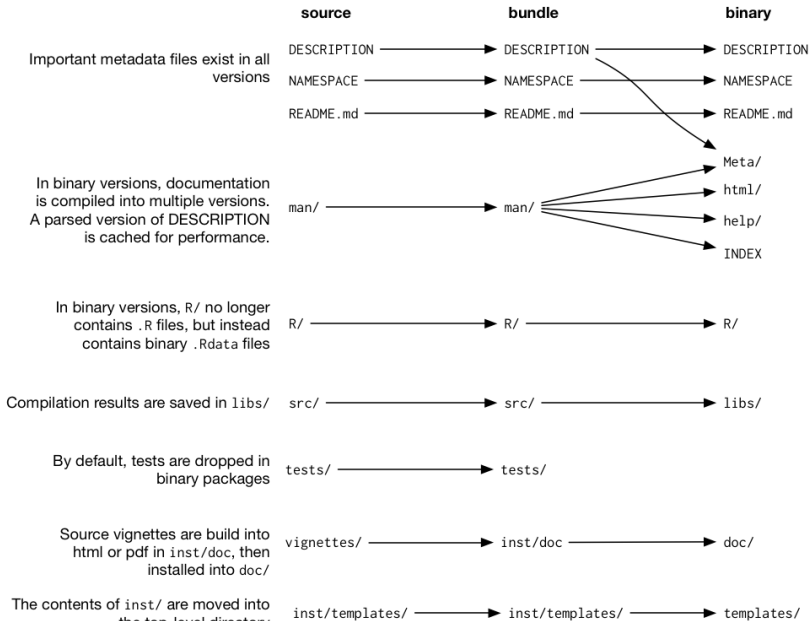
`src/` C, C++, FORTRAN source code

`tests/` tests

`vignettes/` vignette

`inst/CITATION` how should the user cite the package?

...



"There are only two hard things in Computer Science: cache invalidation and naming things." – Phil Karlton

Naming R Packages:

- ▶ can contain letters and numbers, but start with a letter
- ▶ avoid self-invented abbreviations, capital letters, ...
- ▶ should be identifiable in online search
- ▶ R package available

R/ **directory** contains all R code:

- ▶ each function in a separate file (good for small packages)
- ▶ everything in one file (ok for small packages)
- ▶ group related functions in a file with meaningful names (best solution for larger projects)

Example: Create R code and structure for myutils

The standard structure can be obtained automatically using `utils::package.skeleton()`:

```
# define some basic functions
add <- function(x, y){ x + y }
plusone <- function(x){ x + 1 }
# prepare some example data
dat <- data.frame(id=1:10, x=rpois(10, 5), y=rpois(10, 5))

# test your source code
add(10, 1)
plusone(4)
dat

# create standard structure
fdlist <- c("add","plusone","dat")
package.skeleton("myutils", fdlist)
```

```
Package: myutils
Type: Package
Title: What the package does (short line)
Version: 1.0
Date: 2019-01-24
Author: Who wrote it
Maintainer: Who to complain to <yourfault@somewhere.net>
Description: More about what it does (maybe more than one line)
License: What license is it under?
```

```
Package: ggplot2
Version: 3.1.0
Title: Create Elegant Data Visualisations Using the Grammar of
       Graphics
Description: A system for 'declaratively' creating graphics,
            based on "The Grammar of Graphics". [...]
Depends: R (>= 3.1)
Imports: digest, grid, gtable (>= 0.1.1), lazyeval, MASS, mgcv,
        plyr (>= 1.7.1), reshape2, rlang (>= 0.2.1), scales,[...]
Enhances: sp
License: GPL-2 | file LICENSE
URL: http://ggplot2.tidyverse.org
BugReports: https://github.com/tidyverse/ggplot2/issues
Collate: 'ggproto.r' 'ggplot-global.R' 'aaa-.r' .....
VignetteBuilder: knitr
RoxygenNote: 6.1.0
NeedsCompilation: no
Author: Hadley Wickham [aut, cre], Winston Chang [aut], [...]
Maintainer: Hadley Wickham <hadley@rstudio.com>
Date/Publication: 2018-10-25 04:30:25 UTC
Built: R 3.5.2: : 2019-01-07 06:31:07 UTC: unix
```

Package: name of the package

Title: description of the package (one line, < 65 characters)

Description: detailed description (one paragraph, multiple sentences)

Version: version number formatwise
major.minor-patchlevel or
major.minor.patchlevel.

Maintainer: name and e-mail of a person who wants to take over the responsibility

License: abbreviation of a software licence (GPL-2, BSD, MIT, ...)

Depends, Suggests, Imports, Enhances package dependencies

URL: for website of a package

Collate: order R files are loaded (default: alphabetically)

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- ▶ R documentation format is very \LaTeX -like output (\LaTeX installation required)

```
\name{add}
\alias{add}
\title{Add together two numbers}
\usage{ add(x, y) }
\arguments{
  \item{x}{A number}
  \item{y}{A number}
}
\value{
The sum of \code{x} and \code{y}
}
\description{ Add together two numbers }
\examples{
add(1, 1)
add(10, 1)
}
```


There are three steps in the transformation from roxygen comments in your source file to human readable documentation:

1. add roxygen comments to your source file
2. `roxygen2::roxygenise()` or `devtools::document()` converts roxygen comments to `.Rd` files
3. R CMD check converts `.Rd` files to human readable documentation

roxygen2: <http://cran.r-project.org/web/packages/roxygen2/vignettes/rd.html>

- ▶ roxygen comments start with #'
- ▶ tags like @param, @return, @author define parts in .Rd file
- ▶ tags like @includes, @export, @importFrom generate NAMESPACE und Collate
- ▶ tags like @method for OOP documentation

```
##' Add together two numbers
##'
##' @param x A number
##' @param y A number
##' @return The sum of x and y
##' @examples
##' add(1, 1)
##' add(10, 1)
add <- function(x, y) {
  x + y
}
```

`add {rvest}`

R Documentation

Add together two numbers

Description

Add together two numbers

Usage

```
add(x, y)
```

Arguments

x A number

y A number

Value

The sum of x and y

Examples

```
add(1, 1)  
add(10, 1)
```

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For building a R package `pkg` run the following commands in your console:

R CMD SHLIB `pkg` compiles C/C++/Fortran code in `pkg/src`

R CMD build `pkg` generates package bundle `pkg.tar.gz` or `pkg.zip`

R CMD INSTALL `pkg.tar.gz` installs package

R CMD check `pkg.tar.gz` runs CRAN validity checks (is `pkg` valid?)

In windows, installation of `Rtools` is required:

```
# On windows:  
R CMD INSTALL --build pkg
```

Example: Build and Check myutils

```
jmanitz@rladies$ R CMD build myutils_complete
* checking for file 'myutils_complete/DESCRIPTION' ... OK
* preparing 'myutils':
* checking DESCRIPTION meta-information ... OK
* installing the package to process help pages
* [...]
* building 'myutils_1.0.tar.gz'

jmanitz@rladies$ R CMD check myutils_1.0.tar.gz
* using log directory '/home/rladies/example/myutils.Rcheck'
* using R version 3.5.2 (2018-12-20)
* using platform: x86_64-pc-linux-gnu (64-bit)
* checking for file 'myutils/DESCRIPTION' ... OK
* [...]
* checking PDF version of manual ... OK
* DONE

Status: OK
```

Resources

- ▶ Hadley Wickham (2015). *R packages*. O'Reilly Media.
Available online: <http://r-pkgs.had.co.nz/>
- ▶ R-project manual: Writing R Extensions.
Available online: <https://cran.r-project.org/doc/manuals/r-release/R-exts.html>
- ▶ Friedrich Leisch. *Creating R Packages: A Tutorial*:
<http://cran.r-project.org/doc/contrib/Leisch-CreatingPackages.pdf>
- ▶ Internet search, R-help, other R packages, ...

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- ▶ although the NAMESPACE file looks like R code, it is not processed as R code
- ▶ specifies which variables in the package should be exported to make them available to package users, and which variables should be imported from other packages

```
import(foo, bar)      # all functions from foo and bar imported
importFrom(foo, f, g) # selected functions f and g from foo
export(f, g)          # export functions f and g
```

- ▶ for packages with many variables to export it may be more convenient to specify the names to export with a regular expression

```
exportPattern("\\^{ } [ \\^{ } \\textbackslash \\textbackslash . ] ")
```

Object-Oriented Programming (OOP)

- ▶ in OOP, computer programs are designed by making them out of objects that interact with one another
- ▶ a **class** defines the behaviour of **objects** by describing their attributes and their relationship to other classes.
- ▶ the class is also used when selecting **methods**, functions that behave differently depending on the class of their input.
- ▶ R has three OO systems: S3, S4, Reference classes (not covered by this lecture), and the system of base types

Picking a System

- ▶ majority of object-oriented code that I have written in R is S3
- ▶ S3 is sufficient for fairly simple objects and methods for pre-existing generic functions like `print()`, `summary()`, and `plot()`
- ▶ S4 may be more appropriate for more complicated systems of interrelated objects
- ▶ good example for S4 is the Matrix package by Douglas Bates and Martin Maechler

S3/S4 Object System Comparison

	S3	S4
defintion	not neccessary	<code>setClass('class_name', ...)</code>
generation of instances	<code>class(object) <- 'class_name'</code>	<code>new('class_name')</code>
inheritance	vector of class names (children before parents)	<code>contains='parental_cl'</code> in definition
test class	<code>inherits(object, 'class_name')</code>	<code>is(object, 'class_name')</code>
access slots	depends on base type: for lists <code>\$</code> or <code>[[]</code> .	new operator: <code>@</code>
list methods	<code>methods()</code>	<code>showMethods()</code>

Conventions: S3/S4 Classes For S3 and S4, there are the following conventions

- ▶ constructor functions should be named like the class itself, e.g. `lm()`, with exception if a class is the return value of a number of functions
- ▶ standard methods, which are available supplied for many classes:
 - `print` basic object information, also when using `<RET>` (S4 `show()`)
 - `summary` more detailed description of the objects instance
 - `plot` graphics
- ▶ every method should have the arguments of the correspondig generic (same order and defaults) and accept an arbitrary number of additional arguments (use `...`)

- ▶ ensure that the generics are imported and register the methods using `S3method` directives
- ▶ the function `print.foo` does not need to be exported

```
# example myutils
export(add)
S3method(print, add)
export(plusone)
S3method(print, plusone)
.
.

# example ggplot2
S3method(autoplot, default)
export(autoplot)
import(plyr)
importFrom(MASS, cov.trob)
.
.
```

- ▶ some additional steps are needed for packages which make use of S4 classes and methods
- ▶ package should depend on package methods (also DESCRIPTION file)
- ▶ you may need to import `graphics::plot` to make visible a function that can be converted into a implicit generic

```
exportPattern("^[[[:alpha:]]+") # regular pattern

import("methods")           # S4
importFrom(graphics, "plot") # S4 plot

# namespaces from dependencies
importFrom("utils", str, head, tail, assignInNamespace, capture.output)

# export methods and classes
exportMethods("cbind2", "rbind2", "plot", "show", "summary" )
exportClasses("denseMatrix", "sparseMatrix")
```

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R functions from devtools that simplifies R packaging:

`load_all()` simulates installing and reloading your package

`document()` updates documentation, file collation and
NAMESPACE.

`test()` reloads your code, then runs all testthat tests.

`run_examples()` will run all examples to make sure they work.

`check_doc()` runs most of the documentation checking
components of R CMD check

`check()` updates the documentation, then builds and checks
the package

`build()`, `build_win()` builds a package file from package
sources (only one R version)

"A unit testing system designed to be fun, flexible and easy to set up." (Wickham)

- ▶ Provides functions that make it easy to describe what you expect a function to do, including catching errors, warnings and messages.
- ▶ Displays test progress visually, showing a pass, fail or error for every expectation. If you're using the terminal, it'll even colour the output.

```
library(testthat)
library(yourpackage)

test_check("yourpackage")
```

`expect_that` describes expected result of your code (value, class, correct error message, computation time, etc.)

`test_that` is grouping a number of expectation's for one functions or a feature

`context()` is grouping a number of content-related tests

```
require(testthat)
test_that("trigonometric functions match identities", {
  expect_that(sin(pi / 4), equals(1 / sqrt(2)))
  expect_that(cos(pi / 4), equals(1 / sqrt(2)))
  expect_that(tan(pi / 4), equals(1))
})
```

testthat: Hadley Wickham (2011). *testthat: Get Started with Testing*. The R Journal 3(1).