What are R Packages?

R Packages



Starting Up

devtools

install.packages("devtools")

•

•

•

•

Checking Package Names

available::available

available::available("ggplot")

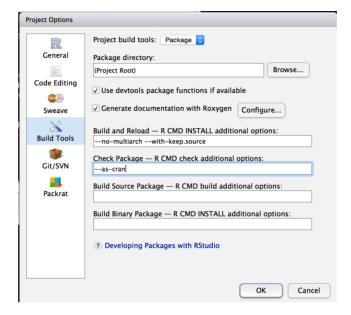
Urban Dictionary:

Not found.
Sentiment:???

```
## Warning: 'BiocInstaller' and 'biocLite()' are deprecated, use the 'BiocManager'
## — ggplot
## Name valid: 
## Available on CRAN: 
## Available on Bioconductor: 
## Available on GitHub: 
## Bad Words: 
## Abbreviations: http://www.abbreviations.com/ggplot
## Wikipedia: https://en.wikipedia.org/wiki/ggplot
## Wiktionary: https://en.wiktionary.org/wiki/ggplot
```

Setting Up

--as-cran



Setting Up

roxygen2

install.packages("roxygen2")

Roxygen Options			
Use roxygen to generate: ✓ Rd files ✓ Collate field			
✓ NAMESPACE file			
√ Vignettes			
Automatically roxygenize when running: R CMD check Source and binary package builds Build & Reload			
OK Cancel			

Modifying the Skeleton

R/hello.R

man/hello.Rd

Build \rightarrow Configure Build Tools \rightarrow Generate Documentation

with Roxygen Vignettes Build and

Reload

--as-cran Build → Configure

Build Tools

NAMESPACE # Generated

by roxygen2: do not edit by hand

DESCRIPTION file

•

function names

DESCRIPTION file

•

.

.

Authors

DESCRIPTION

Authors@R desc

DESCRIPTION file: additional fields

```
library require

base stats methods

ALL

examples vignettes
```

```
usethis::use_package("tidyr", type = "Imports")
usethis::use_package("dplyr", type = "Suggests")
```

Description

Description

Roxygen2

```
top = function(x, n) {
    xx = x[1:n, 1:n]
    hist(xx)
    print(xx)
}
```

top.R R/

Roxygen2

```
top = function(x, n) {
```

@title @description

```
#' @title
#' @description
#'

#' @param x
#' @param n
#'

#' @return
#' @export
#'
#' @examples
```

- · @param
- @return
- · @export

_

• @examples
 \dontrun{}

```
#' @title Print the top of a matrix
#' @description \code{top} is a small function to not just present the first rows
#' of a matrix, but also the first number of columns
# "
#' @param x a \code{matrix}
#' @param n Number of rows and columns to display of the matrix
# "
#' @return A \code{NULL}
#' @export
# "
#' @examples
#' mat = matrix(rnorm(100), nrow = 10)
\#' top(mat, n = 4)
#' \dontrun{
\#' top (mat, n = 10)
# 1 }
```

Functions: a little style

```
NULL is.null()
    #' @rdname

#' @inheritParams
verbose
message cat suppressMessages
...
do.call(FUNCTION, args = list_of_arguments)
```

NAMESPACE

NAMESPACE

· @export

-

• @import package @import PACKAGENAME

- ALL

- whole

package

library(PACKAGENAME)

NAMESPACE

· @importFrom function @import

PACKAGENAME func1 func2

-

- pkgA A pkgB A B @import pkgA A @import pkgB B A() pkgA

- base

stats quantile graphics hist

@importFrom graphics hist

Build and Reload

.

-

_

Using Data

data/ use_data

must

.RData rda

usethis::use_data(DATAOBJECT, compress = "xz")

DATAOBJECT.rda data

Making Data

data-raw

use_data

usethis::use_data_raw()

Documenting Data

DATAOBJECT

```
#' @title Some object to document
#'
#' @description A list containing things
#'
#' @format A list with 7 elements, which are:
#' \describe{
#' \item{x}{first thing}
#' \item{y}{second thing}
#' }
"DATAOBJECT"
```

Different kinds of data

```
inst/
blah.csv inst/
```

inst/extdata

find.package

find.package("readr")

[1] "/Library/Frameworks/R.framework/Versions/3.5/Resources/library/readr"

system.file

system.file("extdata", package = "readr")

[1] "/Library/Frameworks/R.framework/Versions/3.5/Resources/library/readr/extdata"

Different kinds of data

file.path

```
system.file("extdata", "challenge.csv", package = "readr")

## [1] "/Library/Frameworks/R.framework/Versions/3.5/Resources/library/readr/extdata/challenge

mustWork

system.file("extdata", "asdfsdf.csv", package = "readr", mustWork = TRUE)

## Error in system.file("extdata", "asdfsdf.csv", package = "readr", mustWork = TRUE): no file
```

Using the file system

Vignettes

•

how

· THIS IS EXACTLY WHAT A REPRODUCIBLE PAPER IS!

usethis::use_vignette("my-vignette")

•

Unit tests

testthat
 test-DESCRIPTOR.R

testthat

tests/testthat

usethis::use_testthat()

usethis::use test("name of test")

General Rule: Any package issue turns into a test.

Unit tests

testthat tests/testthat

test-DESCRIPTOR.R testthat

```
testthat::context("OVERALL DESCRIPTION OF TESTS IN THIS FILE")
testthat::test_that("Description of this test", { MYCODE })
testthat::expect_equal(OUTPUT, 1234.34535)
testthat::expect_identcal(OUTPUT1, OUTPUT2)
testthat::expect_true(SOME_OUTPUT)
testthat::expect_silent({ no_warning_error_code })
testthat::expect_message({ some_warn }, "a[test]regexp")
```



Building CI and README

```
usethis::use_git()
usethis::use_github() # must have GITHUB_PAT set up
usethis::use_github(protocol = "https") # must have GITHUB_PAT set up
usethis::use_readme_rmd()
usethis::use_appveyor()
usethis::use_travis()
```

Configuring Travis

.travis.yml

```
os:
   - linux
   - osx

warnings_are_errors: true
after_success:
   - Rscript -e 'covr::codecov(type = "all")'
```

Configuring Appveyor

appveyor.yml

environment:
 global:
 WARNINGS ARE ERRORS: 1

Adding to the README.Rmd

GITHUB_USERNAME/REPO

[![Travis-CI Build Status](https://travis-ci.com/GITHUB_USERNAME/REPO.svg?branch=master)](https://ci.appveyor.com/api/projects/status/github/GITHUB_USERNAME/REPO.svg?branch=master)]

```
usethis::use_badge("Travis-CI Build Status",
src = "https://travis-ci.com/GITHUB_USERNAME/REPO.svg?branch=master",
href = "https://travis-ci.com/GITHUB_USERNAME/REPO")
```

README.Rmd

S3, S4, Reference Classes

•		class(x) =	= "myS3Class"
	-		
	-		
•		new	
	-		
	-		-
•			
	-	class\$method	d ()
	_		

S3, S4 Methods

Compiled Code: C and C++

· src/

• cleanup configure

• configure make

Makevars Makefile

configure.win Makevars.win

· .Call

Rcpp

Questions?