


# Introduction to Building packages in R



## Who am I?



- ▶ **Name:** Nicolas Attalides
- ▶ **Coding in  since:** 2005 (yes that's before RStudio!)
- ▶ **Profession:** Senior Data Scientist and trainer (6+ yrs.)
- ▶ **Education:** PhD in Statistical Science from UCL (2015)
- ▶ **R Status:** A never-ending evolving R dinosaur
- ▶ **Hobbies:** Tennis and coding (not at the same time)

# Workshop Setup:



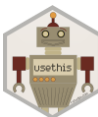
## Wi-Fi

- ▶ Network Name: N/A
- ▶ Password: N/A


## Resources

- ▶ R (version 3.6.3) 
- ▶ RStudio (version 1.4.1106)  Studio

## Packages

- ▶ **{tidyverse}** (version 1.3.0) 
- ▶ **{devtools}** (version 2.3.2) 
- ▶ **{usethis}** (version 2.0.1) 

# What is an R package?

An  package is like a **collection** of **code**, **data** and **documentation** that follow some standard rules and formats.

This is the best way for an R user to **share** their **work** and enable others to use the functionality that is developed.



# Comprehensive R Archive Network (CRAN)

The central repository of R packages is called the Comprehensive R Archive Network (**CRAN**). This contains an archive of R distributions and has more than 17,000 packages ready to be installed and used.

Available CRAN Packages By Name	
<a href="#">A</a> <a href="#">B</a> <a href="#">C</a> <a href="#">D</a> <a href="#">E</a> <a href="#">F</a> <a href="#">G</a> <a href="#">H</a> <a href="#">I</a> <a href="#">J</a> <a href="#">K</a> <a href="#">L</a> <a href="#">M</a> <a href="#">N</a> <a href="#">O</a> <a href="#">P</a> <a href="#">Q</a> <a href="#">R</a> <a href="#">S</a> <a href="#">T</a> <a href="#">U</a> <a href="#">V</a> <a href="#">W</a> <a href="#">X</a> <a href="#">Y</a> <a href="#">Z</a>	
<a href="#">A3</a>	Accurate, Adaptable, and Accessible Error Metrics for Predictive Models
<a href="#">aaSEA</a>	Amino Acid Substitution Effect Analyser
<a href="#">AATtools</a>	Reliability and Scoring Routines for the Approach-Avoidance Task
<a href="#">ABACUS</a>	Apps Based Activities for Communicating and Understanding Statistics
<a href="#">abbyyR</a>	Access to Abbyy Optical Character Recognition (OCR) API
<a href="#">abc</a>	Tools for Approximate Bayesian Computation (ABC)
<a href="#">abc.data</a>	Data Only: Tools for Approximate Bayesian Computation (ABC)
<a href="#">ABC-PAD</a>	Array-Based CpG Region Analysis Pipeline





Find out more about CRAN here: <https://cran.r-project.org/>

# Topics

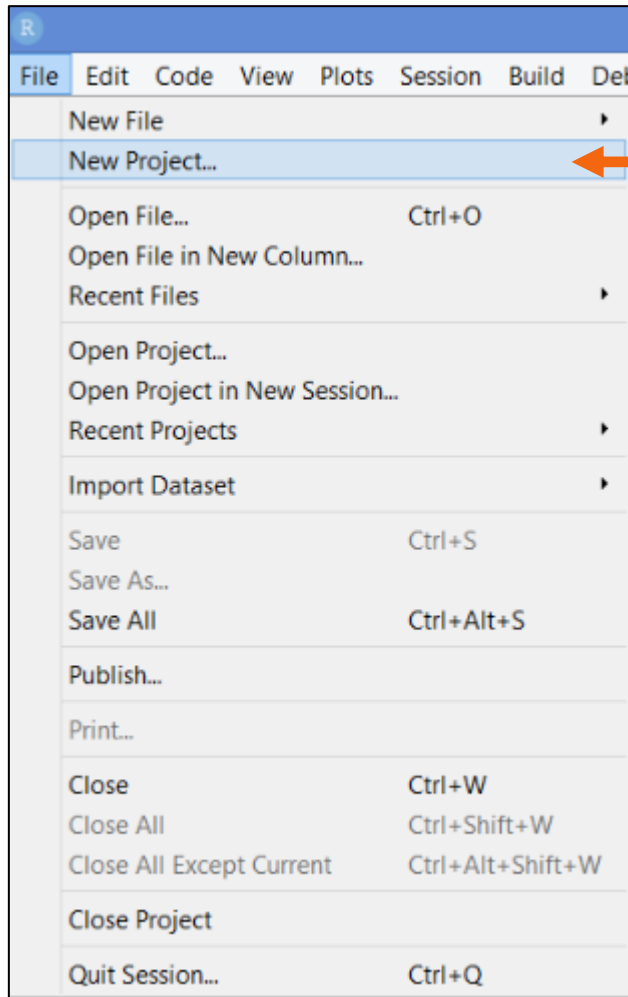
▶ Workshop aim:

Learn how to build an R package in an easy step-by-step approach.

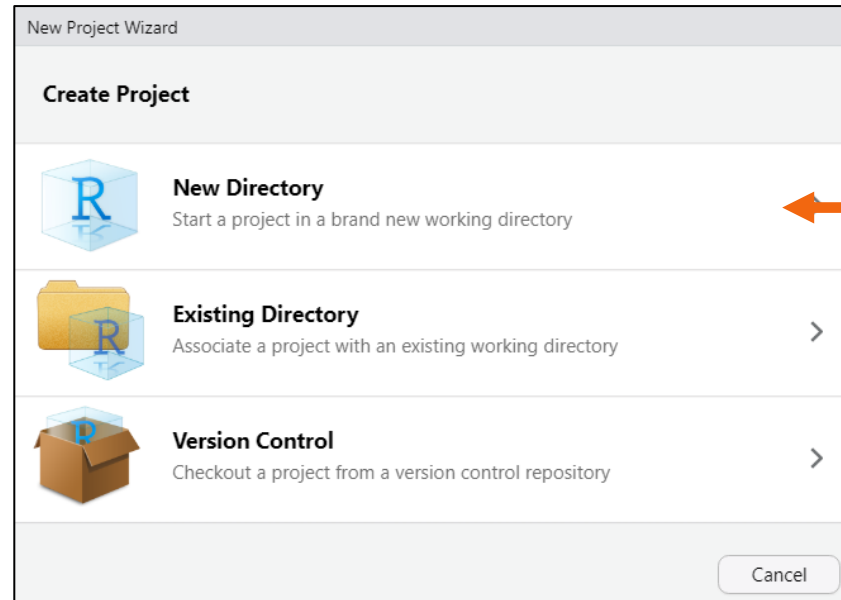
▶ Topics:

- Learn how to create an R package within  Studio®
- Understand the package structure and its various components
- Learn how to write, document and test functions in  for a package
- Learn how to check, build and install an R package

# Create an R package

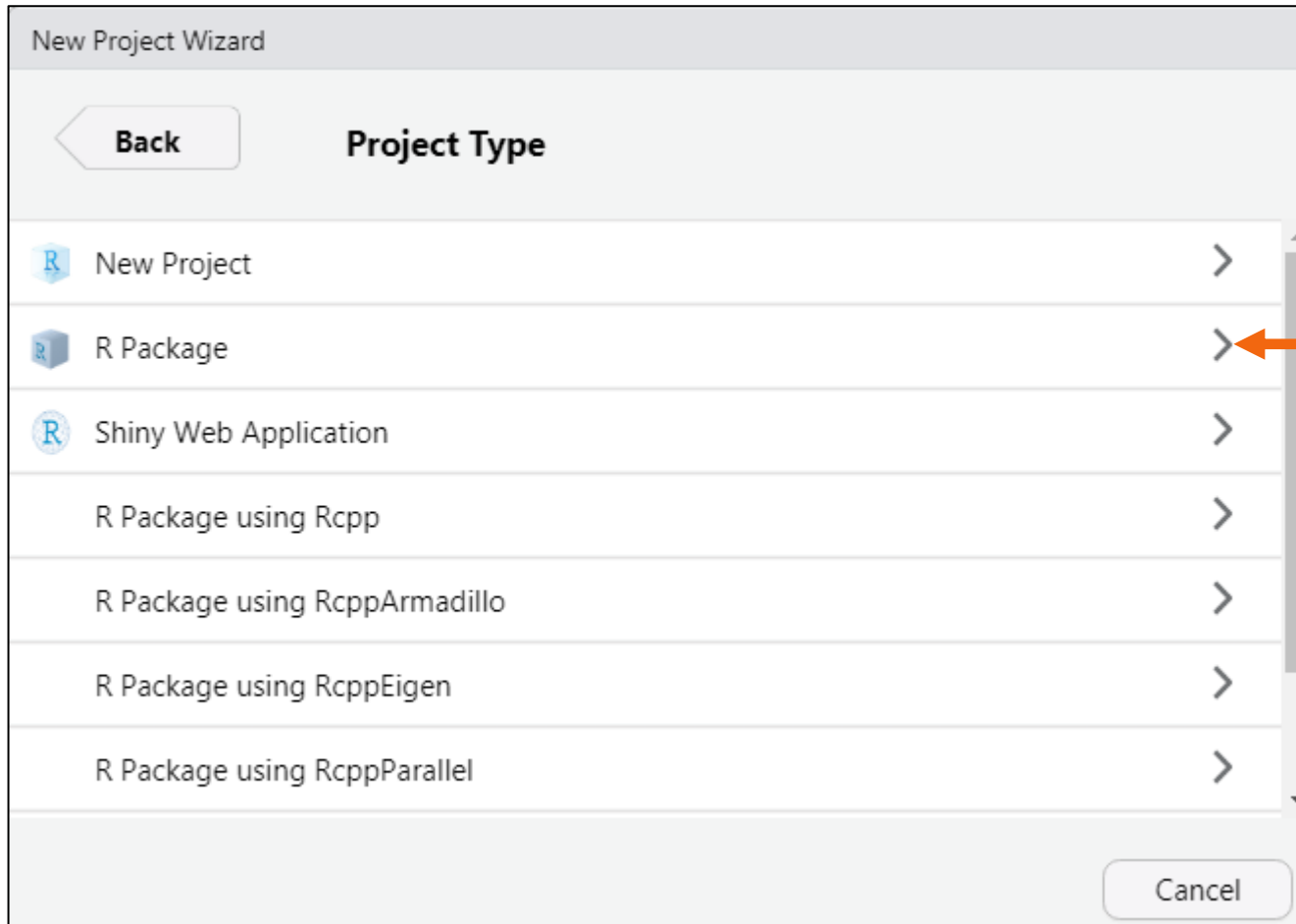


Open the File options and select  
"New Project..."



Then select  
"New Directory"

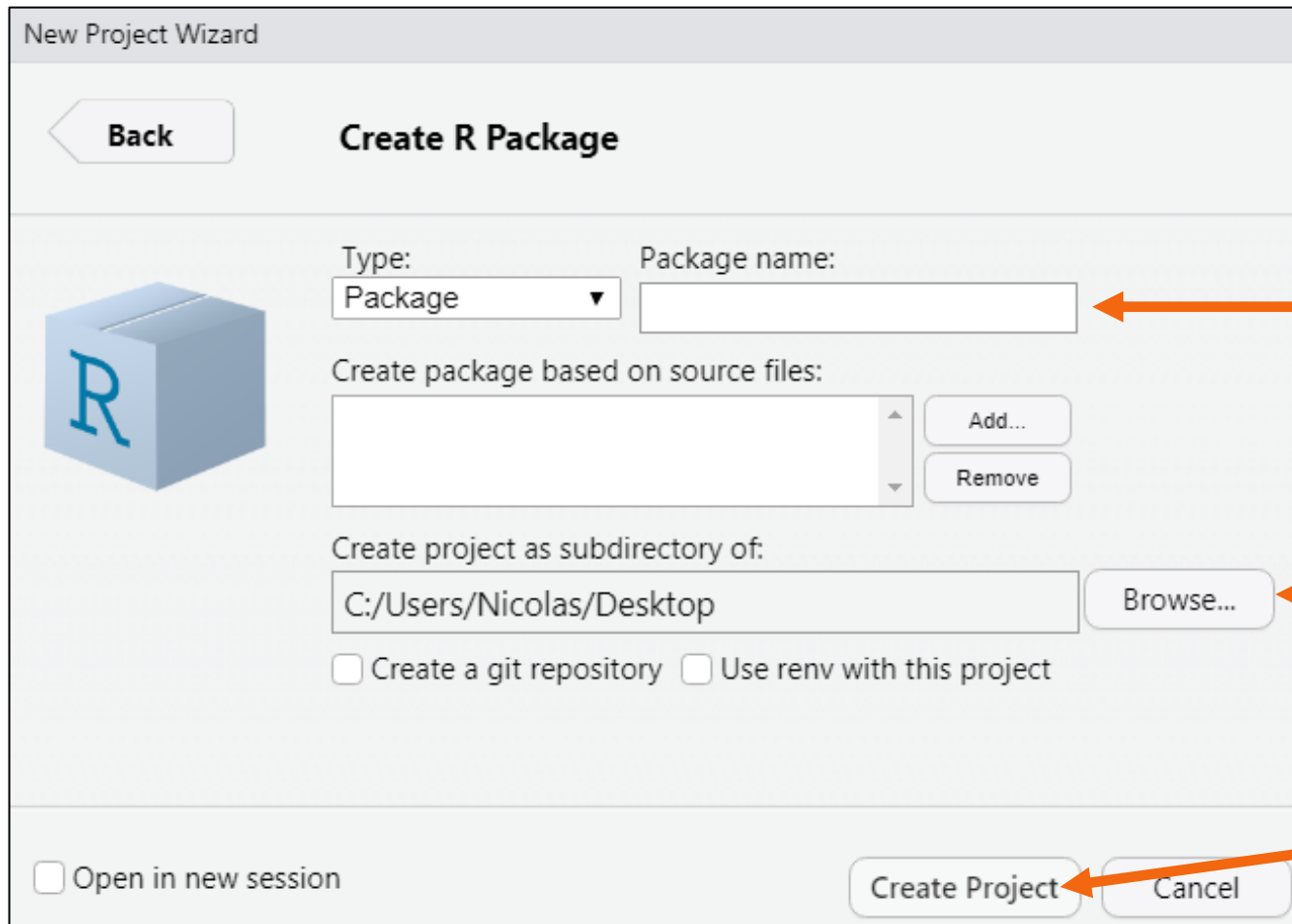
# Create an R package



Select "R Package"



# Create an R package










Type in package name

Select the location of the package

Select "Create Project"

# Congratulations! You created an R package\*!

	Name	Size	Modified
	 ..		
<input type="checkbox"/>	 man		
<input type="checkbox"/>	 R		
<input type="checkbox"/>	 test.Rproj	376 B	Mar 30, 2021, 10:28 AM
<input type="checkbox"/>	 .Rbuildignore	30 B	Mar 30, 2021, 10:28 AM
<input type="checkbox"/>	 DESCRIPTION	377 B	Mar 30, 2021, 10:28 AM
<input type="checkbox"/>	 NAMESPACE	32 B	Mar 30, 2021, 10:28 AM



\*An empty package with the basic structure...

# Create an R package

An alternative way to create an R package with the same result is to use the `create_package()` function from the `{usethis}` package.

```
library(devtools)
```

```
create_package("C:/Users/Nicolas/Desktop/myRpackage")
```



Package path

Package name



This opens a new RStudio session with the new project loaded

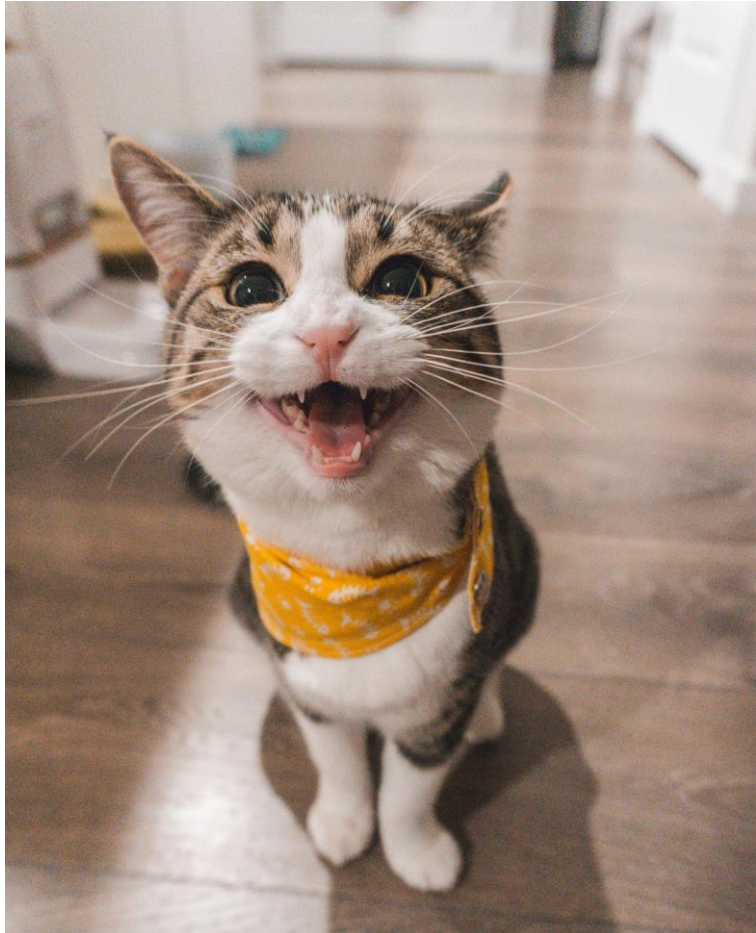
# Congratulations! You created an R package\*!

```
✓ Creating 'C:/Users/Nicolas/Desktop/test/'
✓ Setting active project to 'C:/Users/Nicolas/Desktop/test'
✓ Creating 'R/'
✓ Writing 'DESCRIPTION'
Package: test
Title: What the Package Does (One Line, Title Case)
Version: 0.0.0.9000
Authors@R (parsed):
  * First Last <first.last@example.com> [aut, cre] (<https://orcid.org/YOUR-ORCID-ID>)
Description: what the package does (one paragraph).
License: `use_mit_license()`, `use_gpl3_license()` or friends to
  pick a license
Encoding: UTF-8
LazyData: true
Roxygen: list(markdown = TRUE)
RoxygenNote: 7.1.1
✓ Writing 'NAMESPACE'
✓ Writing 'test.Rproj'
✓ Adding '^test\\.Rproj$' to '.Rbuildignore'
✓ Adding '.Rproj.user' to '.gitignore'
✓ Adding '^\\.Rproj\\.user$' to '.Rbuildignore'
✓ Opening 'C:/Users/Nicolas/Desktop/test/' in new RStudio session
✓ Setting active project to '<no active project>'
```



\*An empty package with the basic structure...

# Live Coding Example 1

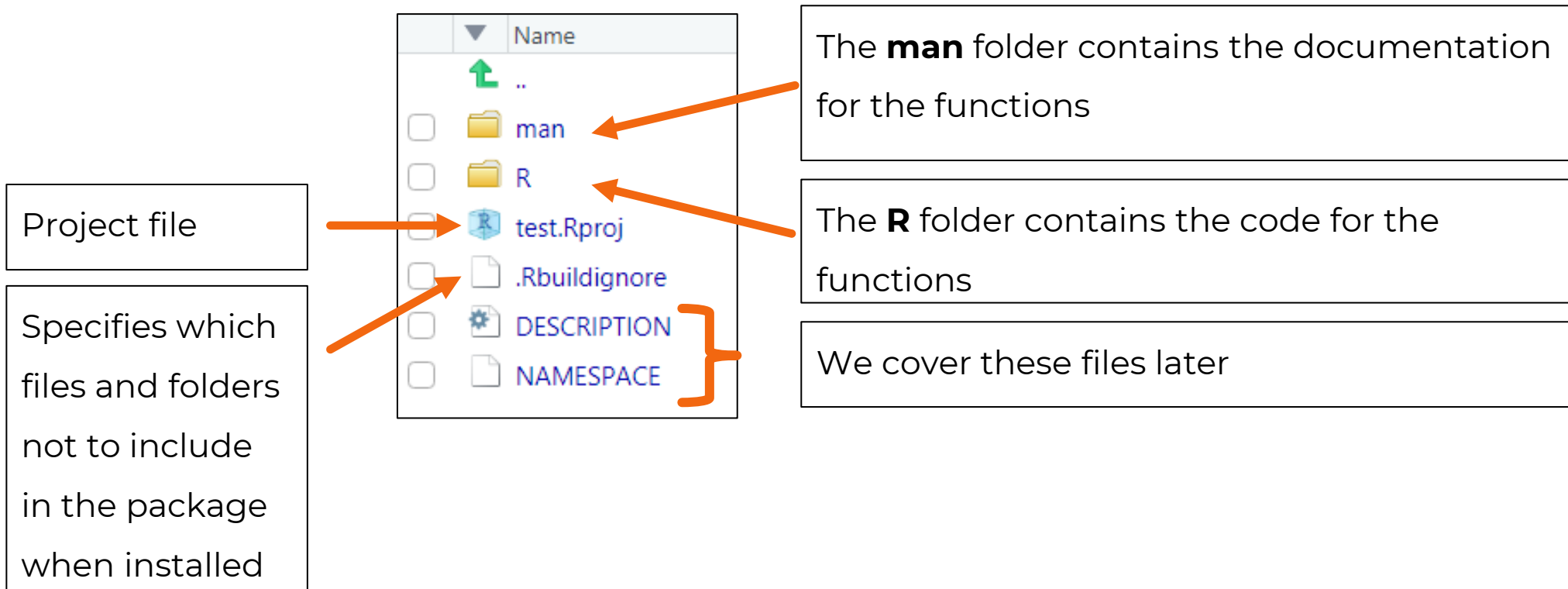



1. Load the `{devtools}` package
2. Create an empty R package called “myRpackage”

## Note:

- ❖ The functionality of this package will be kept simple
- ❖ The main goal is to help the learning process and practice the package development cycle!

# File and Folder structure of an R package



 The **man** folder will be missing when you use the `create_package()` function but it will be automatically created with the first documentation step

# Document an R package

The function **document()** from **{devtools}** is used to build all the documentation for a package.

```
document() # CTRL + SHIFT + D
```

For an empty package this function will simply create the “man” folder if it does not exist.

```
> document()  
Updating myRpackage documentation  
Loading myRpackage
```



# Check an R package

The function `check()` from `{devtools}` automatically builds and checks a package. It runs through a number of checks and will return a summary of the check results.

```
check() # CTRL + SHIFT + E
```

```
-- R CMD check results ----- test 0.0.0.9000 ----  
Duration: 8.9s  
  
> checking DESCRIPTION meta-information ... WARNING  
Non-standard license specification:  
  `use_mit_license()`, `use_gp13_license()` or friends to pick a  
  license  
Standardizable: FALSE  
  
0 errors ✓ | 1 warning x | 0 notes ✓
```



## Live Coding Example 2



1. Load the **{devtools}** package
2. Document the package  
**{myRpackage}**
3. Check the package  
**{myRpackage}**

# DESCRIPTION file

The **DESCRIPTION** file is used to store important **metadata** about the package. For example:

- ▶ What is the package title
- ▶ What is the package version
- ▶ Who to contact
- ▶ Who can use it (the license)
- ▶ What other packages are needed for it to work

```
Package: test
Title: What the Package Does (One Line, Title Case)
Version: 0.0.0.9000
Authors@R:
  person(given = "First",
         family = "Last",
         role = c("aut", "cre"),
         email = "first.last@example.com",
         comment = c(ORCID = "YOUR-ORCID-ID"))
Description: what the package does (one paragraph).
License: `use_mit_license()`, `use_gpl3_license()` or friends to
        pick a license
Encoding: UTF-8
LazyData: true
Roxygen: list(markdown = TRUE)
RoxygenNote: 7.1.1
```

# DESCRIPTION fields

Below is some important guidelines to follow for the DESCRIPTION file fields.

Field	Description
Title	This is typically a one line description of the package. It should be <b>plain text</b> , not more than <b>65 characters</b> long, <b>capitalised like a title</b> , and <b>NOT end in a period!</b>
Description	This is a more detailed text about your package. It can be multiple sentences but it is <b>limited to one paragraph</b> , each line must be no more than 80 characters wide and new lines must be indented with 4 spaces!
Imports	The packages that are listed in this field <b>must</b> be installed on your computer for your package to work because they are being used by your package. If any package is missing, it <b>will</b> be <b>automatically installed</b> when your package is installed.
Suggests	The packages that are listed in this field can be used by your package but they are <b>not required</b> . If any package is missing, it <b>will not</b> be <b>automatically installed</b> when your package is installed.

# Package version

The **Version** field of the DESCRIPTION file shows the package's version number.

- ▶ Typically a version number is made up of three numbers:

**<major>.<minor>.<patch>**

For example: 1.3.0

- ▶ A package that is **in-development** usually has ends with "9000"

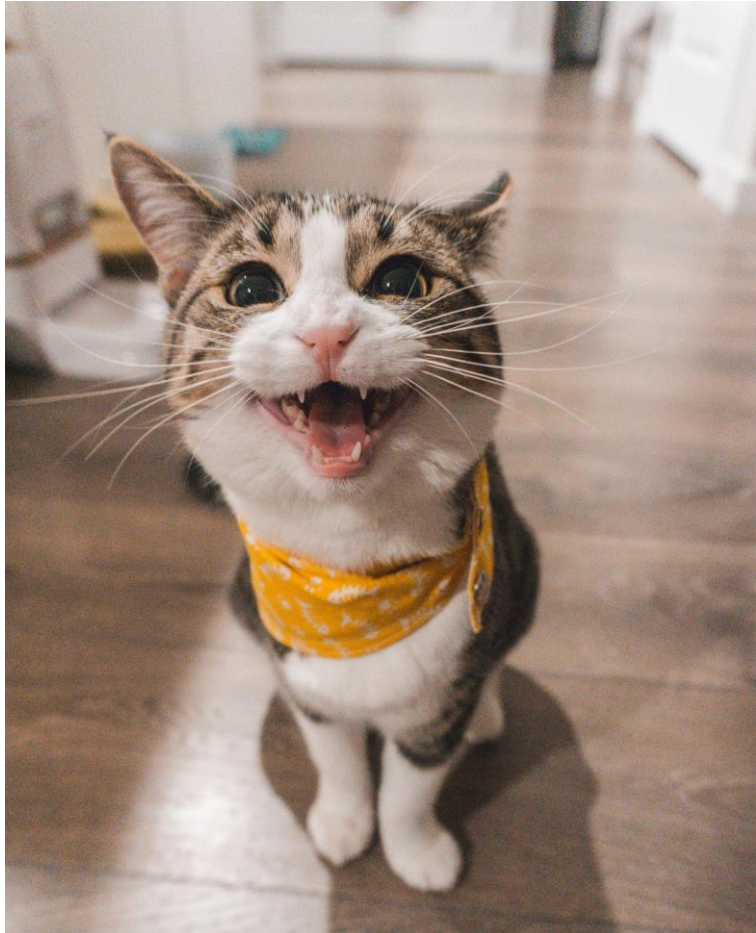
For example: 1.3.0.9000



Example of DESCRIPTION file of the **{devtools}** package

```
Package: devtools
Title: Tools to Make Developing R Packages Easier
Version: 2.3.2.9000
Authors@R:
  c(person(given = "Hadley",
           family = "Wickham",
           role = "aut"),
    person(given = "Jim",
           family = "Hester",
           role = c("aut", "cre"),
           email = "jim.hester@rstudio.com"),
    person(given = "Winston",
           family = "Chang",
           role = "aut"),
    person(given = "RStudio",
           role = "cph"),
    person(given = "R Core team",
           role = "ctb",
           comment = "Some namespace and vignette code extracted from base R"))
Description: Collection of package development tools.
License: GPL (>= 2)
URL: https://devtools.r-lib.org/, https://github.com/r-lib/devtools
BugReports: https://github.com/r-lib/devtools/issues
Depends:
  R (>= 3.0.2),
  usethis (>= 2.0.1)
```

# Live Coding Example 3



1. Edit the DESCRIPTION file with the metadata about the package (*Title* and *Description*) and your details (*Author*)
2. Save the changes
3. Document the package  
**{myRpackage}**
4. Check the package  
**{myRpackage}**



# NAMESPACE file

The **NAMESPACE** file can be quite confusing and is considered an advanced topic (hence the warning!).

In a simplified way, it is the file that controls the **communication between packages** and their functions. For example, it manages the **functions** to **export** (from your package) and functions to **import** (from other packages).



```
This document is read only.  
1 # Generated by roxygen2: do not edit by hand  
2  
3
```

# What is an R function

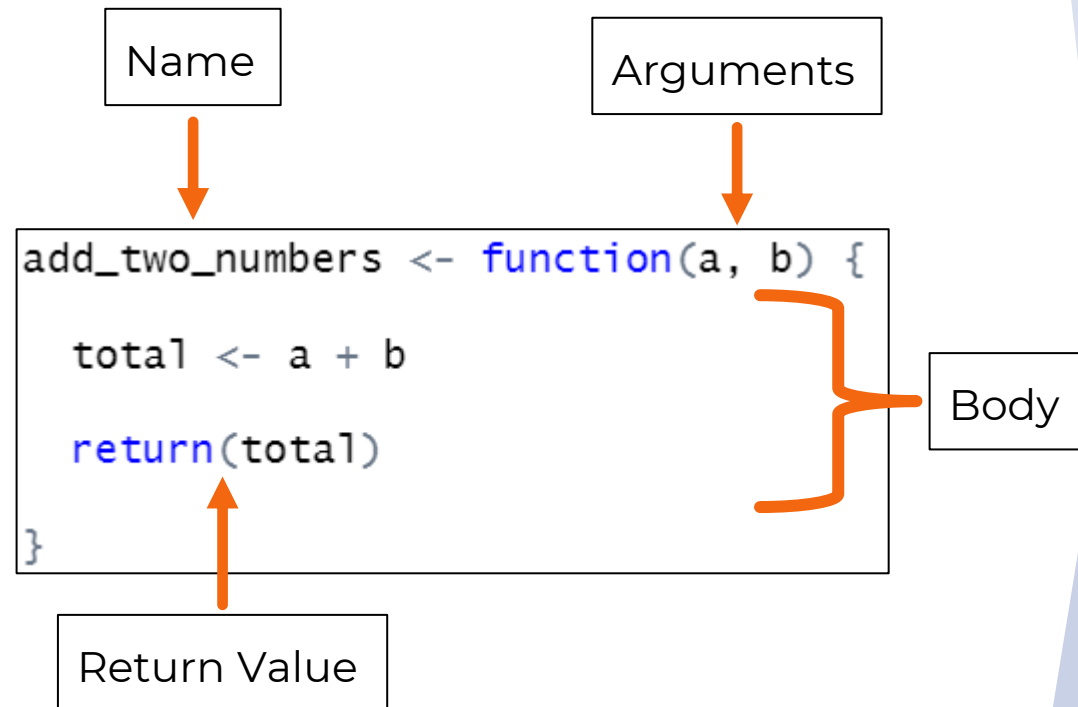
An R function is an **R object** that contains code to be executed.


- ▶ In a simplified way, an R function takes **inputs** and generates **outputs**
- ▶ R functions are useful when we have code that is **repeated** in a script and therefore help to avoid “copy-pasting” code
- ▶ An R function ideally **self-contains** a complex piece of code and is dedicated to solving one task

# Components of an R function

An R function is created just like any other object in R and follows a specific structure.

Component	Description
Name	The name of the function
Arguments	The values passed to the function (inputs)
Body	R code that the function executes
Return Value	The value/s the function is required to return (outputs)



 The return value is the last expression evaluated by the function. If the `return()` expression is used then the function will output the contents of `return()`



## Create a function

The package **{usethis}** makes it easy for us to **add** R functions to a package. The function **use\_r()** takes as input the **function name** and creates the “.R” file for that function inside the “R” folder.

```
# Create a function file in the R folder
use_r("add_two_numbers")

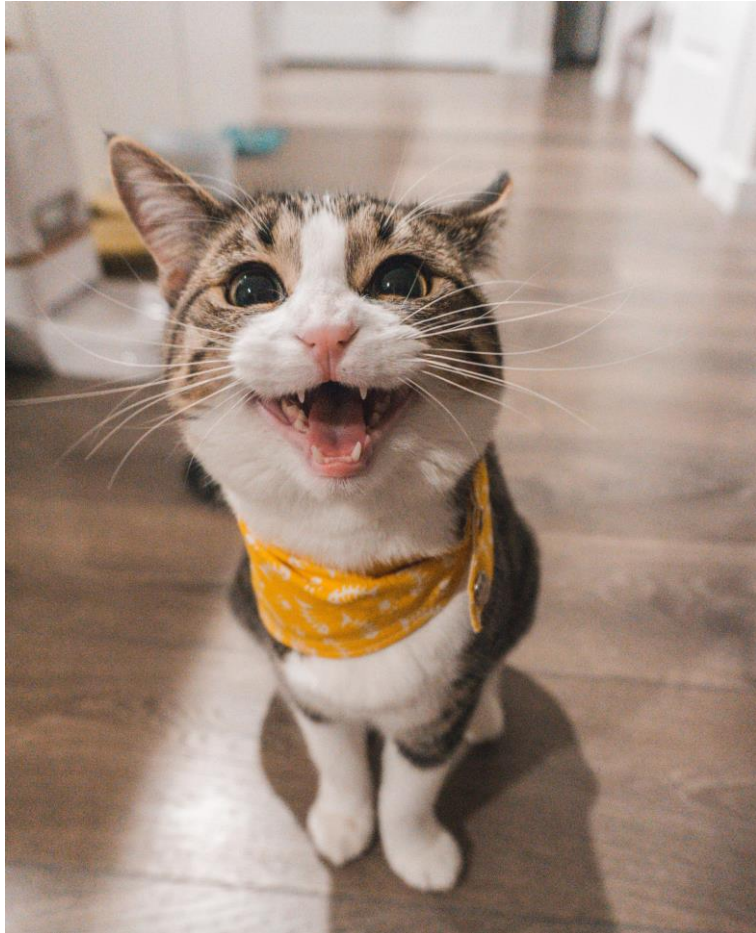
# Place the function code inside add_two_numbers.R
add_two_numbers <- function(a, b) {

  total <- a + b

  return(total)

}
```

# Live Coding Example 4



1. Create the following functions for the package **{myRpackage}**
  - `add_two_numbers()`
  - `subtract_two_numbers()`

# Live Coding Example 4

```
# Create a function file in the R folder
use_r("subtract_two_numbers")

# Place the function code inside add_two_numbers.R
subtract_two_numbers <- function(a, b) {

  total <- a - b


  return(total)

}
```

# Try out the function

Before we invest more time to properly document and test the function, it is a good idea to first **check if the function works!**

There are three ways to do this:

- ▶ (Messy way): Create the function arguments as objects and then run the code inside the function line by line
- ▶ (Script way): Source the “.R” file containing the function and call the function
- ▶  (Dev way): Use `load_all()` and call the function – more on this later

# Live Coding Example 5



Try out the function `add_two_numbers()` using the “Messy” and “Script” ways.

## Document a function

Function documentation can be somewhat fiddly to work with...

Typically each “.R” file in the “R” folder (containing a function) has its corresponding “.Rd” file in the “man” folder with the same name (containing the documentation).

The “.Rd” file that contains the **documentation** in an R-specific markup language ... luckily we do not have to worry about that!

# Document a function

What we need to do in order to document a function is write specially formatted comments above the function. These are called **roxygen** comments and the package **{roxygen2}** creates and edits the “.Rd” files for us!





# Roxygen comments

A **roxygen block** of comments above a function will determine the help provided to the user about the function.

```
#' Title  
#'  
#' Description  
#'  
#' @param  
#'  
#' @return  
#'  
#' @export
```

Comment	Description
Title	The first sentence and represents the title of the documentation
Description	The second paragraph and describes what the package does
@param	Argument name followed by a description and what it does
@return	Describes the output of the function
@export	Specifies that the function is usable outside of package

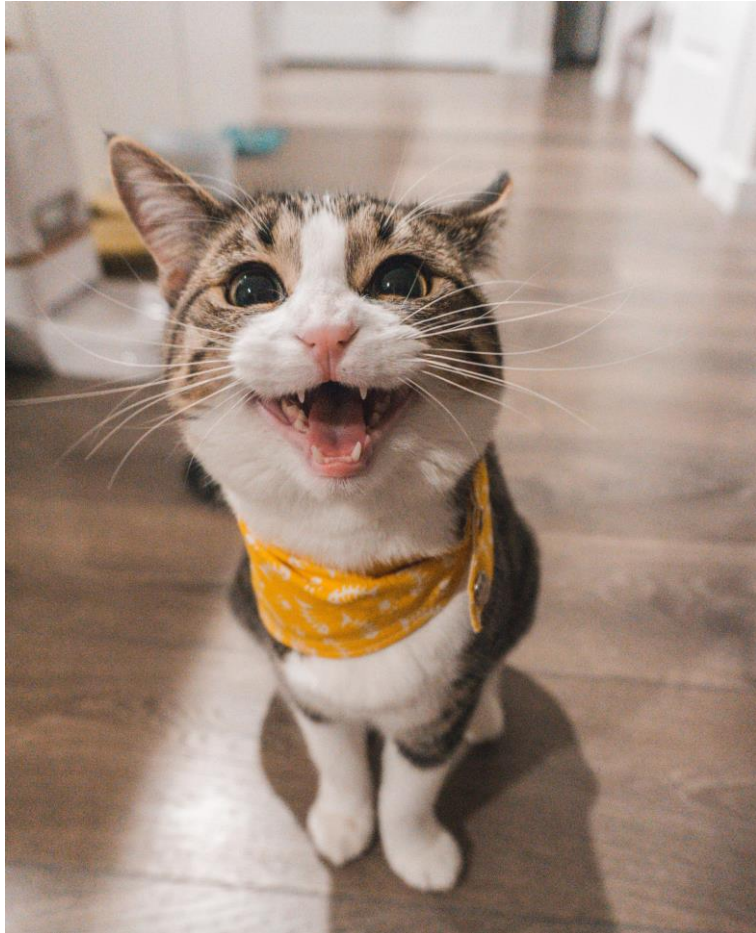
**Step 1:** Open the “.R” file of the function

**Step 2:** Place cursor somewhere in the function code

**Step 3:** Go to → Code → Insert Roxygen Skeleton or CTRL + ALT + SHIFT + R



# Live Coding Example 6



1. Documenting the function `add_two_numbers()`
2. Document the package `{myRpackage}`
3. Inspect help information of the function `add_two_numbers()`
4. Check the package `{myRpackage}`

# Live Coding Example 6

```
#' Add two numbers together
#'  
#' The purpose of this function is to take two numbers as inputs and add them  
#' together. The numbers can be positive or negative but not NA.  
#'  
#' @param a (numeric) A positive or negative number  
#' @param b (numeric) A positive or negative number  
#'  
#' @return The total sum of the two numbers  
#'  
#' @export  
#' @examples  
#' add_two_numbers(a = 1, b = 1)  
add_two_numbers <- function(a, b) {  
  
  total <- a + b  
  
  return(total)  
  
}
```

# Live Coding Example 6

```
add_two_numbers {demoConvertR}
```

R Documentation

## Add two numbers together

### Description

The purpose of this function is to take two numbers as inputs and add them together. The numbers can be positive or negative but not NA.

### Usage

```
add_two_numbers(a, b)
```

### Arguments

a (numeric) A positive or negative number

b (numeric) A positive or negative number

### Value

The total sum of the two numbers

### Examples

```
add_two_numbers(a = 1, b = 1)
```



## Add tests to a package

The next step is to add **tests**. This is a **formal** way to test the functionality of your package and that your functions work as you expect them to! First we need to initialise testing for the package.

```
use_testthat()
```

The function **use\_testthat()** from the **{usethis}** package adds “*Suggests: testthat*” to the DESCRIPTION file. It also creates the folders “tests/testthat/” and adds a generic script “testthat.R” in the “tests” folder.



Spend some time to write unit tests for your functions!

# Test a function

The `use_test()` function from `{usethis}` takes as input the function name and creates the “test-[name].R” file for that function inside the “testthat” folder with a generic test ready to edit.

**NOTE:** It is the developer’s responsibility to write the unit tests.

## Objects

```
expect_equal() expect_identical()  
expect_type() expect_s3_class()  
expect_s4_class()
```

## Vectors

```
expect_length()  
expect_lt() expect_lte() expect_gt()  
expect_gte()  
expect_named()  
expect_setequal() expect_mapequal()  
expect_true() expect_false()  
expect_vector()
```



Check out more information here: <https://testthat.r-lib.org/reference/index.html>

# Test an R package

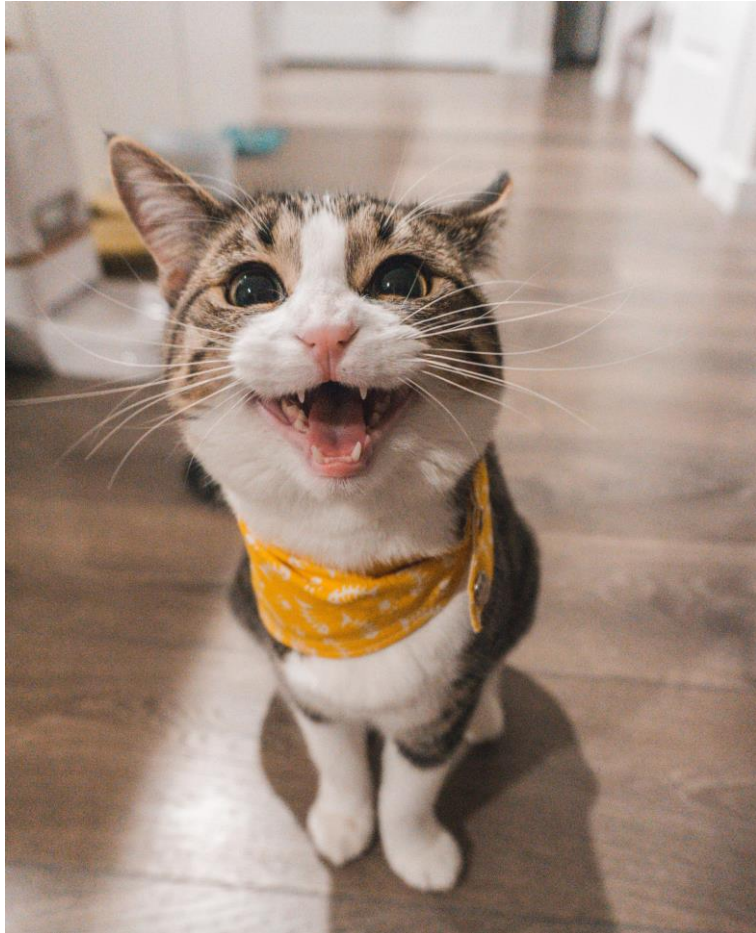
The function `test()` from the package `{devtools}` is used to run all of the tests of a package. It also prints out a test report about test **failures**, **warnings**, skipped tests and of course **passes**!

```
test() # CTRL + SHIFT + T
```





# Live Coding Example 7



1. Add testing to the package **{myRpackage}**
2. Write some unit tests for the function **add\_two\_numbers()**
3. Test the package **{myRpackage}**
4. Document the package **{myRpackage}**
5. Check the package **{myRpackage}**

# Live Coding Example 7

```
test_that("add_two_numbers returns the correct value and type", {
  expect_identical(object = add_two_numbers(a = 1, b = 2), expected = 3)

  expect_identical(object = add_two_numbers(a = 1, b = -1), expected = 0)

  expect_type(object = add_two_numbers(a = 1, b = 2), type = "double")
})

test_that("add_two_numbers returns NA if one of the arguments is NA", {
  expect_identical(object = add_two_numbers(a = NA, b = 2), expected = as.numeric(NA))
})

test_that("add_two_numbers returns Inf (+/-) if one of the arguments is infinite", {
  expect_identical(object = add_two_numbers(a = Inf, b = 2), expected = Inf)

  expect_identical(object = add_two_numbers(a = -Inf, b = 2), expected = -Inf)
})
```



# Live Coding Example 7

## ❖ Add testing components

```
✓ Adding 'testthat' to suggests field in DESCRIPTION
✓ Setting Config/testthat/edition field in DESCRIPTION to '3'
✓ Creating 'tests/testthat/'
✓ Writing 'tests/testthat.R'
```

## ❖ Add test for function

```
✓ Writing 'tests/testthat/test-add_two_numbers.R'
* Modify 'tests/testthat/test-add_two_numbers.R'
```

## ❖ Check DESCRIPTION file

```
Suggests:
  testthat (>= 3.0.0)
Config/testthat/edition: 3
```

## ❖ Run tests

```
Loading demoConvertR
Testing demoConvertR
✓ | OK F W S | Context
✓ | 6       | add_two_numbers [0.1 s]

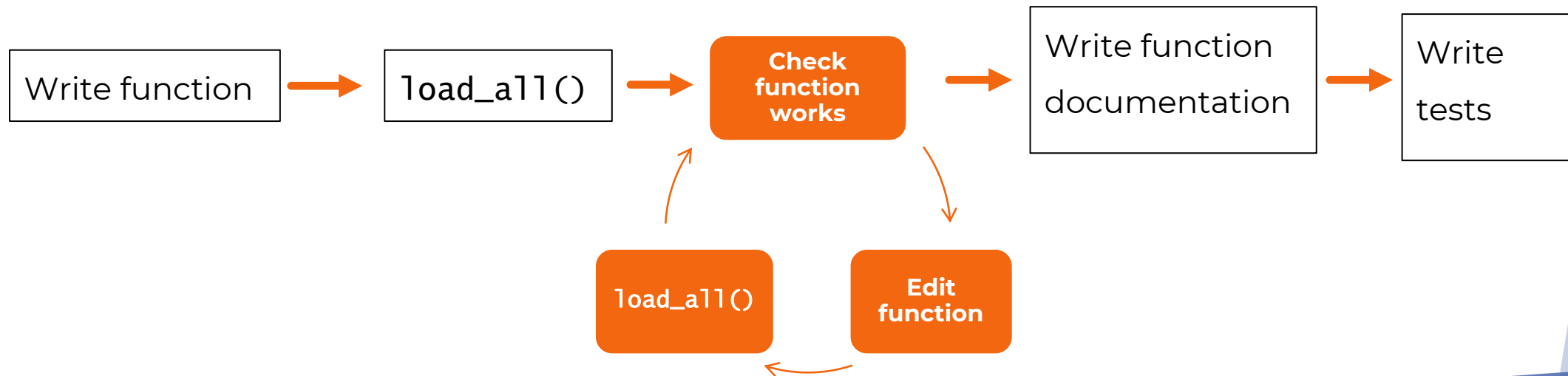
== Results ==
Duration: 0.2 s

[ FAIL 0 | WARN 0 | SKIP 0 | PASS 6 ]
```



# Test drive a package

It is a good idea to **regularly** test drive the functionality that we develop in our package. This means **fewer bugs** to worry about! An important function to use during the development cycle is the `load_all()` of the `{devtools}` package.



# Test drive a package

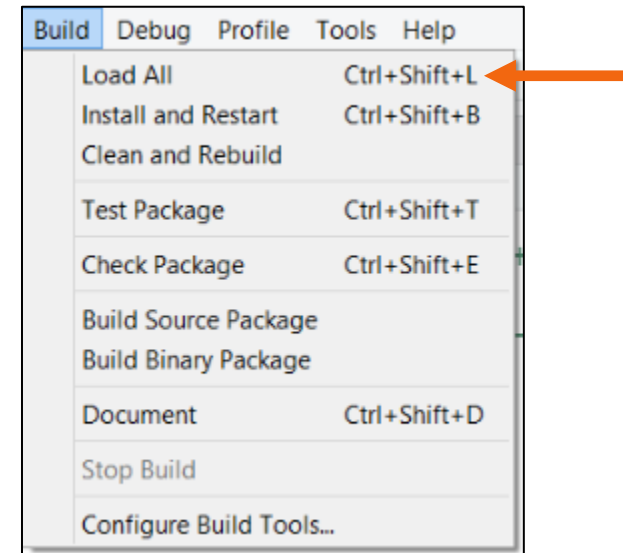
The function `load_all()` is very useful because it allows you to **interact** with your package and its functions.

You can think of it as a way to **simulate** what happens when a package is installed and loaded with `library()`.

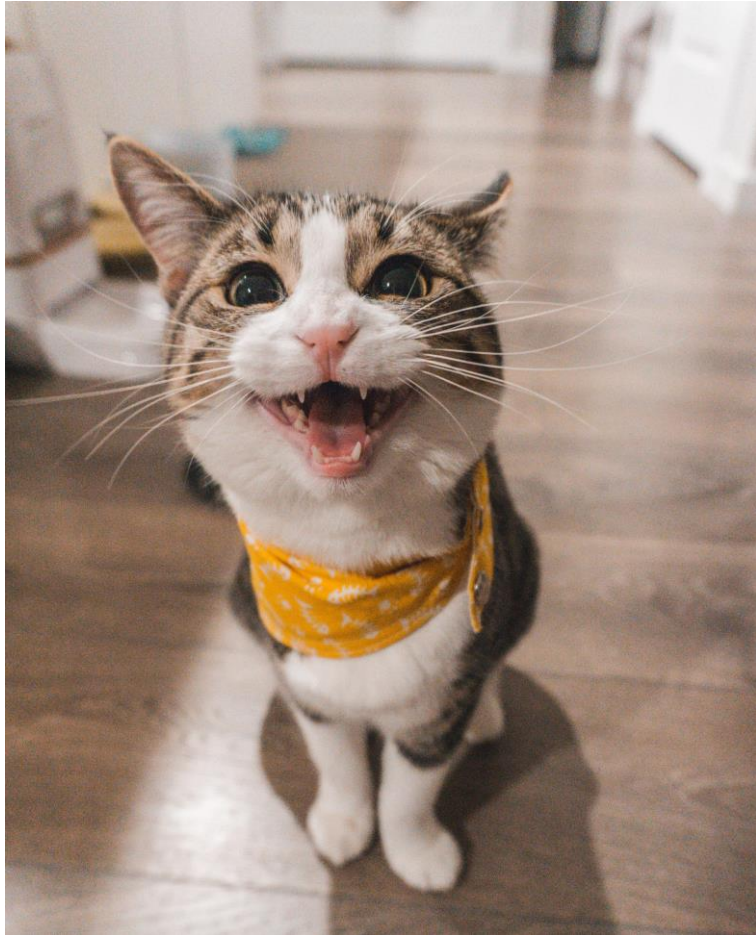
```
library(devtools)
```

```
# working directory is set at the top level of package
```

```
load_all() # CTRL + SHIFT + L
```



# Live Coding Example 8



1. Restart the R session

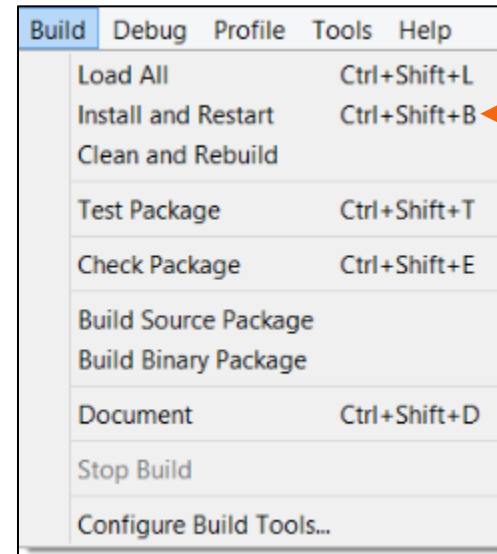
Go to → Session → Restart R

(or CTRL + SHIFT + F10)

2. Test drive the **{myRpackage}** package using the `load_all()` function

# Install a package

The function `install()` from the `{devtools}` package installs the package from the source state. More specifically the “R CMD INSTALL” command is executed behind the scenes.



```
install()
```

Alternatively the option “Install and Restart” (CTRL + SHIFT + B) installs the package, restarts R and loads it.

# Uninstall a package

An installed package is stored on your computer in a **library directory**.

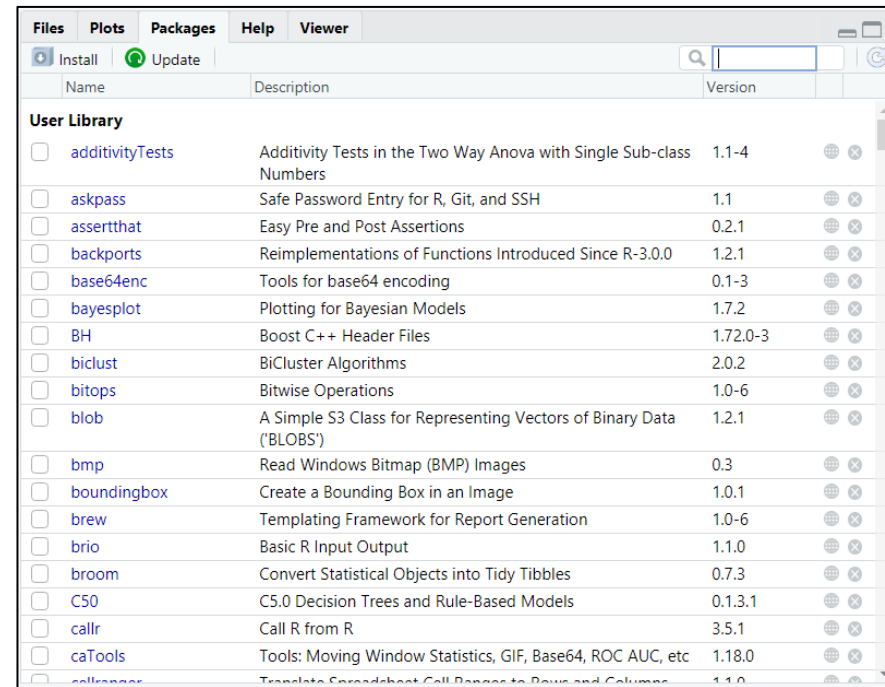
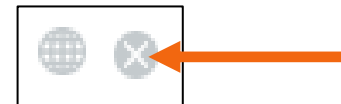
Typically we update, however it is a good idea to know how to uninstall them.







































The function `remove.packages()` from the `{utils}` package removes a package from the library directory.

```
remove.packages()
```



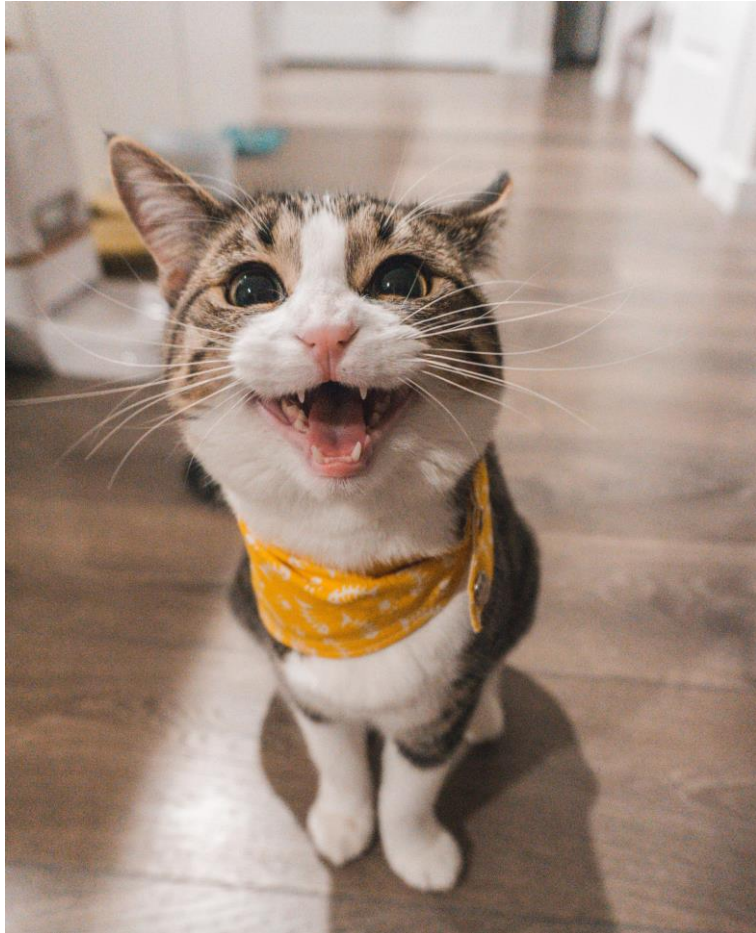
You can also navigate to the “Packages” tab, search for the package and click on the cross to uninstall the package



Name	Description	Version	
<b>User Library</b>			
<input type="checkbox"/> additivityTests	Additivity Tests in the Two Way Anova with Single Sub-class Numbers	1.1-4	 
<input type="checkbox"/> askpass	Safe Password Entry for R, Git, and SSH	1.1	 
<input type="checkbox"/> assertthat	Easy Pre and Post Assertions	0.2.1	 
<input type="checkbox"/> backports	Reimplementations of Functions Introduced Since R-3.0.0	1.2.1	 
<input type="checkbox"/> base64enc	Tools for base64 encoding	0.1-3	 
<input type="checkbox"/> bayesplot	Plotting for Bayesian Models	1.7.2	 
<input type="checkbox"/> BH	Boost C++ Header Files	1.72.0-3	 
<input type="checkbox"/> biclust	BiCluster Algorithms	2.0.2	 
<input type="checkbox"/> bitops	Bitwise Operations	1.0-6	 
<input type="checkbox"/> blob	A Simple S3 Class for Representing Vectors of Binary Data ('BLOBS')	1.2.1	 
<input type="checkbox"/> bmp	Read Windows Bitmap (BMP) Images	0.3	 
<input type="checkbox"/> boundingbox	Create a Bounding Box in an Image	1.0.1	 
<input type="checkbox"/> brew	Templating Framework for Report Generation	1.0-6	 
<input type="checkbox"/> brio	Basic R Input Output	1.1.0	 
<input type="checkbox"/> broom	Convert Statistical Objects into Tidy Tibbles	0.7.3	 
<input type="checkbox"/> C50	C5.0 Decision Trees and Rule-Based Models	0.1.3.1	 
<input type="checkbox"/> callr	Call R from R	3.5.1	 
<input type="checkbox"/> caTools	Tools: Moving Window Statistics, GIF, Base64, ROC AUC, etc	1.18.0	 
<input type="checkbox"/> callr	Translate Spreadsheet Cell Ranges to Rows and Columns	1.1.0	 



# Live Coding Example 9



1. Install the **{myRpackage}** package
2. Try out the package
3. Uninstall the **{myRpackage}** package



## Install a package from GitHub

We can also install an R package that is available on a **public GitHub repo**.

The **{devtools}** package offers the function `install_github()` which installs a package directly from GitHub (GitHub **username** and repository **name** are needed for this function).

For example:

```
install_github("tidyverse/dplyr")
```



This is useful when a package is not available to install from CRAN or you want to install the latest in-development version of a package.

# Add the pipe operator to your package

The pipe operator (`%>%`) from the **{magrittr}** package is extremely useful when writing code and especially for data transformations. To use the pipe operator within a package that you are developing call the function **`use_pipe()`** from **{usethis}** which carries out the necessary setup.

## Description

Does setup necessary to use magrittr's pipe operator, `%>%` in your package. This function requires the use roxygen.

- Adds magrittr to "Imports" in DESCRIPTION.
- Imports the pipe operator specifically, which is necessary for internal use.
- Exports the pipe operator, if `export = TRUE`, which is necessary to make `%>%` available to the users of your package.

## Use other packages in your package

It is very likely that you will want to use functionality from another package within your package. To do this you need to **add** the “external” package to the “**Imports**” field of the **DESCRIPTION** file.

An easy way to do this is by using the function `use_package()` from `{usethis}`. For example:

```
use_package("dplyr")
```

This adds the `{dplyr}` package to the “**Imports**” field of the **DESCRIPTION** file. It is recommended to use the `::` operator when using functions from other packages. For example: `dplyr::filter()`

# Keyboard Shortcuts

Below is the collection of keyboard shortcuts seen in this course that helps speed up the development cycle of an R package.

Shortcut	Description
CTRL + SHIFT + D	Runs <code>document()</code> – build all the documentation for a package
CTRL + SHIFT + E	Runs <code>check()</code> – builds and checks a package
CTRL + ALT + SHIFT + R	Go to “Code” then “Insert Roxygen Skeleton”
CTRL + SHIFT + T	Runs <code>test()</code> – run all the tests of a package
CTRL + SHIFT + L	Runs <code>load_all()</code> – enables test driving a package
CTRL + SHIFT + F10	Go to “Session” then “Restart R” to Restart R session
CTRL + SHIFT + B	Installs the package, restarts R and loads it

# Package Development cheat sheet

## Package Development: : CHEAT SHEET

### Package Structure

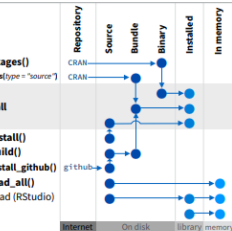
A package is a convention for organizing files into directories. This sheet shows how to work with the 7 most common parts of an R package:

- Package
- DESCRIPTION
- R/
- tests/
- man/
- vignettes/
- data/
- NAMESPACE

The contents of a package can be stored on disk as:

- source** - a directory with sub-directories (as above)
- bundle** - a single compressed file (.tar.gz)
- binary** - a single compressed file optimized for a specific OS

Or installed into an R library (loaded into memory during an R session) or archived online in a repository. Use the functions below to move between these states.



**install.packages()**  
install.packages(type = "source")

**R CMD install**

devtools:install()  
devtools:build()  
devtools:install\_github()  
devtools:load\_all()  
Build & Reload (RStudio)  
library()

devtools:use\_build\_ignore("file")  
Adds file to .Rbuildignore, a list of files that will not be included when package is built.

### Setup (DESCRIPTION)

The DESCRIPTION file describes your work, sets up how your package will work with other packages, and applies a copyright. You must have a DESCRIPTION file

- Add the packages that yours relies on with devtools:use\_package()
- Adds a package to the Imports or Suggests field

CCO	MIT	GPL-2
No strings attached.	MIT license applies to your code, and all code anyone bundles with it, if re-shared.	GPL-2 license applies to your code, and all code anyone bundles with it, if re-shared.

```

Package: mypackage
Title: Title of Package
Version: 0.1.0
Authors@R: person("Hadley", "Wickham", email = "hadley@rstudio.com", role = c("aut", "cre"))
Description: what the package does (one paragraph)
Depends: R (>= 3.1.0)
License: GPL-2
LazyData: true
Imports:
  dplyr (>= 0.4.0),
  ggvis (>= 0.2)
Suggests:
  knitr (>= 0.1.0)
    
```

Import packages that your package must have to work. R will install them when it installs your package.

Suggest packages that are not very essential to yours. Users can install them manually, or not, as they like.

### Write Code (R/)

All of the R code in your package goes in R/. A package with just an R/ directory is still a very useful package.

- Create a new package project with devtools:create("path/to/name")
- Create a template to develop into a package.
- Save your code in R/ as scripts (extension .R)

### Test (tests/)

Use tests/ to store tests that will alert you if your code breaks.

- Add a tests/ directory
- Import testthat with devtools:use\_testthat(), which sets up package to use automated tests with testthat
- Write tests with context(), test(), and expect statements
- Save your tests as .R files in tests/testthat/

**WORKFLOW**


1. Edit your code.
2. Load your code with one of devtools:load\_all()
  - Re-loads all saved files in R/ into memory.
  - Ctrl/Cmd + Shift + L (keyboard shortcut)
  - Saves all open files then calls load\_all().
3. Experiment in the console.
4. Repeat.

• Use consistent style with [r-pkgs.had.co.nz/r/htmlstyle](http://r-pkgs.had.co.nz/r/htmlstyle)

• Click on a function and press F2 to open its definition

• Search for a function with Ctrl + .

Visit [r-pkgs.had.co.nz](http://r-pkgs.had.co.nz) to learn much more about writing and publishing packages for R



## Document (man/)

man/ contains the documentation for your functions, the help pages in your package.

- Use roxygen comments to document each function beside its definition
- Document the name of each exported data set
- Include helpful examples for each function

### WORKFLOW

1. Add roxygen comments in your .R files
2. Convert roxygen comments into documentation with one of:
  - devtools:document()
  - Converts roxygen comments to .Rd files and places them in man/. Builds NAMESPACE.
3. Open help pages with ? to preview documentation
4. Repeat

**Ctrl/Cmd + Shift + D** (Keyboard Shortcut)

**.Rd FORMATTING TAGS**

<code>{emph}</code> (italic text)	<code>{email}</code> (name@foo.com)
<code>{strong}</code> (bold text)	<code>{href}</code> (url){display}
<code>{code}</code> (function(args))	<code>{url}</code> (url)
<code>{pkg}</code> (package)	<code>{link}</code> (=dest){display}
<code>{dntun}</code> (code)	<code>{linkS4}</code> (class){class}
<code>{dntshow}</code> (code)	<code>{code}</code> (link){function}
<code>{dnttest}</code> (code)	<code>{code}</code> (link){package}{function}

```

{deqn}a + b (block)
{eqn}a + b (inline)
{tabular}
  left {tab centered}
  right {tab}
  cr
  cell {tab cell}
  {tab cell} {cr}
    
```

### COMMON ROXYGEN TAGS

@aliases	@inheritParams	@seealso
@concepts	@keywords	@format
@describeIn	@param	@source
@examples	@rdname	@include
@export	@return	@slot
@family	@section	@field
		data
		S4
		RC

## Add Data (data/)

The data/ directory allows you to include data with your package.

- Save data as .Rdata files (suggested)
- Store data in one of data/, R/Sysdata.rda, inst/extdata
- Always use LazyData: true in your DESCRIPTION file.

devtools:use\_data()

Adds a data object to data/ (R/Sysdata.rda if internal = TRUE)

devtools:use\_data\_raw()

Adds an R Script used to clean a data set to data-raw/. Includes data-raw/ on .Rbuildignore.

Store data in

- data/ to make data available to package users
- R/Sysdata.rda to keep data internal for use by your functions.
- inst/extdata to make raw data available for loading and parsing examples. Access this data with system.file()

## Organize (NAMESPACE)

The NAMESPACE file helps you make your package self-contained; it won't interfere with other packages, and other packages won't interfere with it.

- Export functions for users by placing @export in their roxygen comments
- Import objects from other packages with package:object (recommended) or @import, @importFrom, @importClassesFrom, @importMethodsFrom (not always recommended)

### WORKFLOW

1. Modify your code or tests.
2. Document your package (devtools:document())
3. Check NAMESPACE
4. Repeat until NAMESPACE is correct

**SUBMIT YOUR PACKAGE**

[r-pkgs.had.co.nz/release.html](http://r-pkgs.had.co.nz/release.html)



## Next online R event!

Build Interactive {shiny} Apps to Share Your Work With Anyone!



Use  slo

**Build Interactive {shiny} Apps to  
Share Your Work With Anyone!**

**Speakers:** Andreas Botnen Smebye (NGI, Oslo)  
Christian Wilhelm Mohr (NIBIO, Ås)

**Time:** Thursday, 20 May 2021, 17:00 CET

**Place:** Zoom

<https://www.meetup.com/Oslo-useR-Group/events/277702734/>

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