

## Lecture 14: Continue on R package intro

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## Before we start

- ▶ Open R project associated with empty R package skeleton we created last time
- ▶ If you don't have it, follow

```
library(usethis)  
  
# Specify path where you want to store the package  
path <- getwd() # current working directory  
# or explicit path  
# path <- "/Users/Irina/Documents/Teaching/TAMU/Computing"  
  
# Create full path  
fullpath <- file.path(path, "MyPackageTest1")  
  
# Create package skeleton  
create_package(fullpath)
```

## DESCRIPTION - Author

```
person("Irina", "Gaynanova",  
       email = "irinag@stat.tamu.edu",  
       role = c("aut", "cre"))
```

```
## [1] "Irina Gaynanova <irinag@stat.tamu.edu> [aut, cre]"
```

Tell that both author (*aut*) and maintainer (*cre*) is Irina Gaynanova.

From H. Wickham

*The full list of roles is extremely comprehensive. Should your package have a woodcutter ("wdc"), lyricist ("lyr") or costume designer ("cst"), rest comfortably that you can correctly describe their role in creating your package*

## DESCRIPTION - Multiple authors

```
c(person("Yunfeng", "Zhang",  
  email = "some email",  
  role = c("aut", "cre")),  
  person("Irina", "Gaynanova",  
    email = "irinag@stat.tamu.edu",  
    role = "aut"))
```

```
## [1] "Yunfeng Zhang <some email> [aut, cre]"
```

```
## [2] "Irina Gaynanova <irinag@stat.tamu.edu> [aut]"
```

# LICENSE

The most confusing part.

- ▶ **GPL 2** or **GPL-3** - open source, most commonly used to my knowledge, anyone who does derivatives of your work has to make the code available under the same license.
- ▶ **MIT** - also open source, the license should always be distributed with the code.

See [R packages](#) for more.

# Version

From Hadley again

*A released version number consists of three numbers,  $\langle \text{major} \rangle . \langle \text{minor} \rangle . \langle \text{patch} \rangle$ . For version number 1.9.2, 1 is the major number, 9 is the minor number, and 2 is the patch number. Never use versions like 1.0, instead always spell out the three components, 1.0.0.*

The fourth number always indicated the package in development, i.e. 9000 (cause many development may happen before the release)

## R package building and checks

Run checks. What does it complain about?

Can Install and Restart.

Can build binary and source files (the latter are needed for CRAN submission)

## Example function/documentation

Suppose you want to have a soft-thresholding function available within your package.

```
soft <- function(a, lambda){  
  sign(a) * max(abs(a) - lambda, 0)  
}
```

Save it as soft.R file within **R/** package folder.

Try to build a package using Build -> More -> Build Source Package.

Install and restart (from Build menu). Try using function soft. What happens?



## Where is function soft?

The package by default **masks** function soft because it has **not been exported**

You can still see that it's there by typing

```
getAnywhere(soft)
```

**getAnywhere()** is a very useful function to check other packages code for functions that are **masked** from the user, that is have not been exported/have no documentation

Go to **NAMESPACE** file. What do you see?

# Basics of documentation

!!! You may need to first install roxygen2 using

```
install.packages("roxygen2")
```

To make functions usable, they have to be documented (at least barely).

While one can write documentation completely manually, it is **strongly recommended** and **required for your project** to use Roxygen.

Go to your Build Toold (Build – > More – > Configure Build Tools). Make sure you have check mark next to Roxygen documentation. Also click on Configure, and add check mark to everything.

## Documentation with roxygen

Go to soft function you created, place a cursor anywhere, and then go to Code— >Insert Roxygen Skeleton

- ▶ Roxygen comments star with `#'`, and come before the function
- ▶ 1st sentence is a user-readable function title
- ▶ **@param** automaticall catch the input of the function. You will need to eventually provide description for each input
- ▶ **@return** here you should explain the function output
- ▶ **@examples** here you should provide example of the function use (**required** by CRAN for all functions that are made available to the user)

## Documentation with roxygen

Try to reinstall you new package (without messing with roxygen).  
Do you see function soft now?

## Documentation with roxygen

Try to reinstall you new package (without messing with roxygen).  
Do you see function `soft` now?

Go to **man** folder. Now you can see **soft.Rd** file. Open that file in Rstudio and hit Preview.

Go to **NAMESPACE** file. What do you see?

# Why roxygen?

Karl Broman:

*With Roxygen2, you write specially-structured comments preceding each function definition. These are processed to produce the .Rd files that R wants, and it also creates that painful NAMESPACE file for you.*

# Documentation with roxygen

Let's give some simple description to soft function within roxygen documentation.

# Documentation with roxygen

You can (and often should) add more to the documentation than required by the skeleton.

See [R packages](#) for some examples and also [R package primer](#)



# Basics of testing

The **testthat** package can be used to create an automatic testing directory for any of your packages.

**usethis** makes it easy to add testing to your package

```
library(usethis)  
use_testthat()  
use_test()
```

What happened?

## Basics of testing - creating a test

```
library(usethis)  
use_testthat()  
use_test()
```

- ▶ The first command created a testing infrastructure.
- ▶ The second command created a particular test skeleton for our soft function.

Modify the tests to create 2 test for soft-thresholding function.

Run the tests using Build— > More— > Test package

## Package dependencies

If your package needs other packages to work properly, i.e. **glmnet**, **ggplot2**, you should use DESCRIPTION file

**Imports:** glmnet, ggplot2

Some older packages have **Depends** but according to Hadley one should not need this unless one really knows why.

Specifying dependencies in DESCRIPTION via Imports will automatically result in installation of those packages with yours.

## Package dependencies

The packages listed in Imports will be **installed**, but not necessarily **attached**.

**What this means?** Use `glmnet::glmnet()`,  
i.e. `pck_name::function_name()` when using outside functions

## Getting it on GITHUB - reminder

You may want to make your package available to the public via GitHub.

**devtools** R package will allow others to seamlessly install any R package that is on GitHub.

**Reminder:** From local to GitHub

- ▶ Commit the work we have done, if not already.
- ▶ Setup a new repository on GitHub **without** **readme**
- ▶ Use

```
git remote add origin [URL path to created repo on GitHub]
git push --set-upstream origin master
```