How to write an R package and publish it on GitHub

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What is R package?

- Collection of code, data, documentation developed by R community
- Addresses particular problem with specialized statistical technique, graphical device, etc.
- Core set of packages come with base R
- > 15,000 Additional packages available from CRAN, Bioconductor, Omegahat, GitHub, etc.
- Popular R packages
 - dplyr
 - ggplot2



What is GitHub?

- Website that hosts software development and version control using Git
- Free basic services
- Truly open source
- "Facebook for programmers"



Why write R package?

- For yourself
 - Save time
 - Keep track of your functions
 - Have all your functions in one place
 - Document your work
 - For publishing papers
 - Increasingly, (bio)statistical journals ask for R package development of novel method
- For others
 - If package is useful to you, it is also useful to someone else
 - Readers of your paper can use proposed method
 - Advance science!

Why publish package on GitHub?

- Reproducibility
- Accessibility
- Collaboration
- Back-up method
- Version-control using git (more on this later)

What do you need to write R package?

- R Studio (https://rstudio.com/)
- devtools & roxygen2 packages
- Git (install from https://git-scm.com/)
- GitHub account (https://github.com/)

Some other useful packages

- here
- available



Let's begin!

Major steps

- 1. Open R Studio
 - i) New Project \to New Directory \to R Package \to Enter info \to Create Project
- 2. Write functions
 - i) Each function should be saved in its own file
 - ii) Write package description and document functions
- 3. Create new repo in GitHub
 - i) Repo name = package name
- 4. Connect to GitHub
- 5. Pull + Commit + Push
- 6. Use/share package with install_github()!

There is more than one way!

Naming R package

Some tips:

- Make it simple & short
- Make it unique (use available package see next slide)
- Must start with a letter & cannot end with a period
- Do not use special characters
- Trend towards using all lower case
- Hadley's book has more information

Naming R package

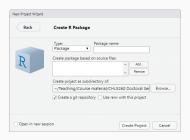
Check to see if name is unique, especially if you plan to submit to CRAN

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

```
#available("ayapack", browse = FALSE)
```

In RStudio

- New Project \rightarrow New Directory \rightarrow R Package
 - Enter package name
 - Optional: Select R scripts that include your functions (if you leave blank, a default function is included)
 - Select subdirectory where you want to save the package (location is not too important since the final product will be saved on Github)
 - Check 'Create a git repository'



In RStudio

This will create the following files and folders

- packagename.Rproj: This indicates that the directory is a project
- DESCRIPTION: This is where all the meta-data about your package goes – you can edit this file manually
- NAMESPACE: This file indicates what needs to be exposed to users for your R package – we will recreate this file using document() (see next slide)
- R: This is where all your R code goes for your package
- man: This is where the manuals for your functions will be saved
- Don't worry too much about the rest (.gitignore, .Rbuildingore)



In RStudio

Next, load the devtools package $\,$

library(devtools)

Then, delete the NAMESPACE file

Great resources

- Book by Hadley Wickham and Jenny Bryan
- devtools cheatsheet
- Blog post by MZES Social Science Data La

Turn this project into a package

```
devtools::create(here::here())
```

This will create 3 additional files

- DESCRIPTION: This is where all the meta-data about your package goes. You can edit this file manually.
- NAMESPACE: This file indicates what needs to be exposed to users for your R package. Do not edit this file.
- R: This is where all your R code goes for your package.

Open new R script and write your function

```
myfunc <- function(x){
  y <- x + x
  return(y)
}</pre>
```

Include @export tag above your function to indicate this function to be "exposed" to users.

```
#' @export
myfunc <- function(x){
  y <- x + x
  return(y)
}</pre>
```

Also, include documentation for your function when you go ?myfunc.

```
#' This is my function.
#'
#' This function returns a value from adding the parameters.
#' @param x
#' @return y
#' @export
myfunc <- function(x){</pre>
  y \leftarrow x + x
  return(y)
```

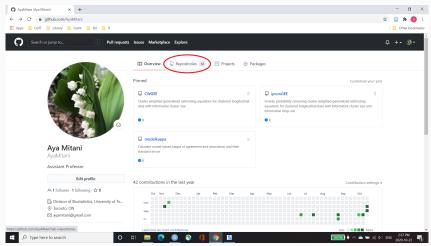
Now run

```
devtools::document()
```

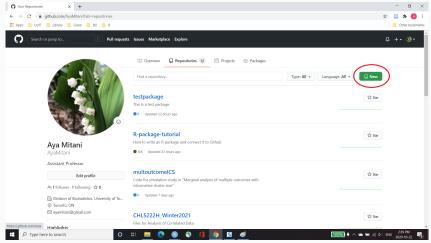
- This will create man directory that includes read-only file myfunc.Rd
- Note that NAMESPACE file has been updated

Add data set

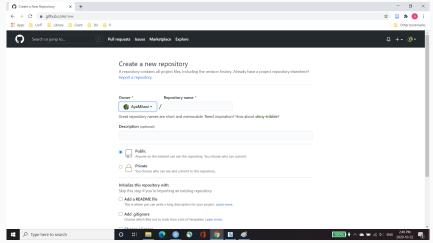
First, log in and go to Repositories



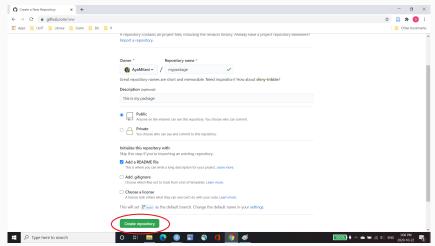
Then, create new repository



Repo name should be same as package name



Finish creating new repo



Finally, copy URL

