

Reproducible projects in R: practical workshop

Marina Vabistsevits

What this workshop is about

1. **Project-oriented workflow** (organising your projects with .Rproj and renv)
 - 1.1. Rstudio efficiency tips
2. Using **Git** to track changes in your R projects
3. **R projects on UKB-RAP/All of Us** platforms with Git



Icon for R



Icon for RStudio



software
company who
develop
Rstudio

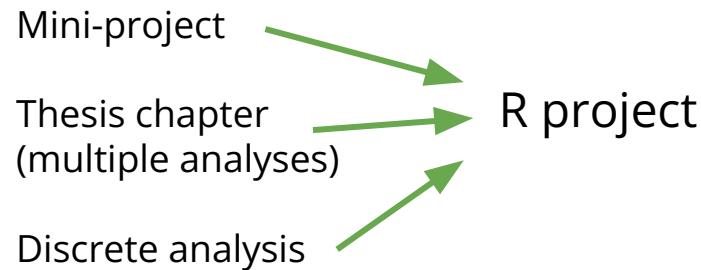


Positron -
new IDE from
Posit for
R/Python etc

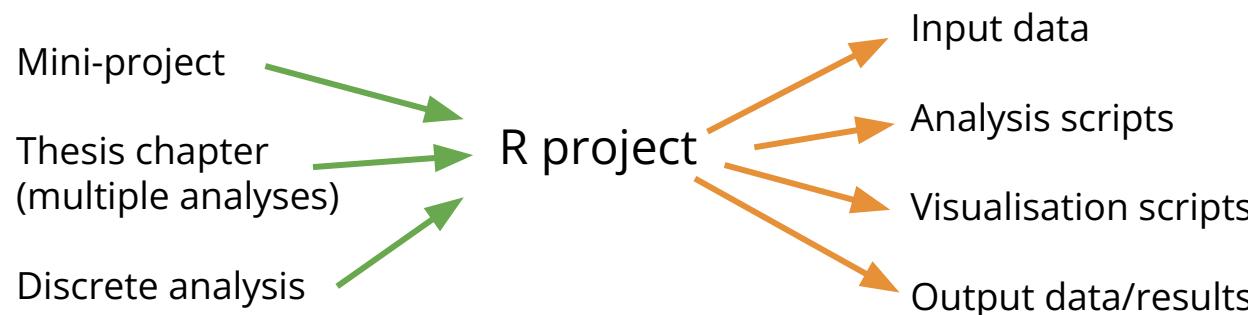
The slides will be shared after!

1. Project-oriented workflow

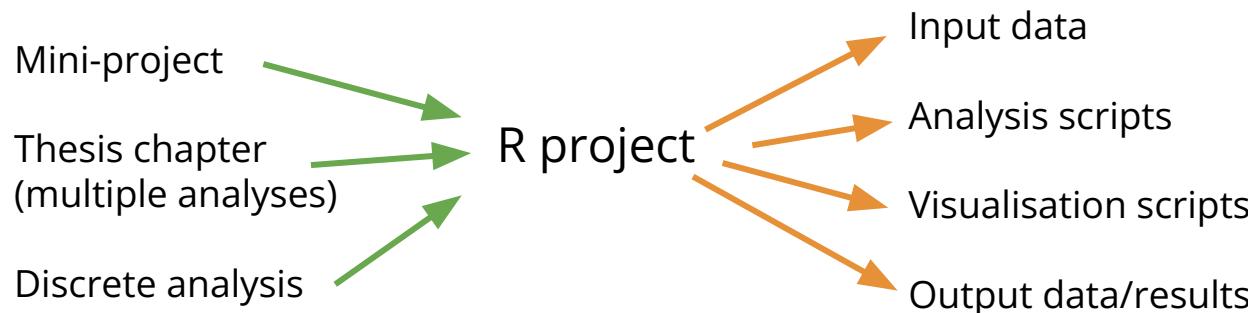
R projects



R projects



R projects



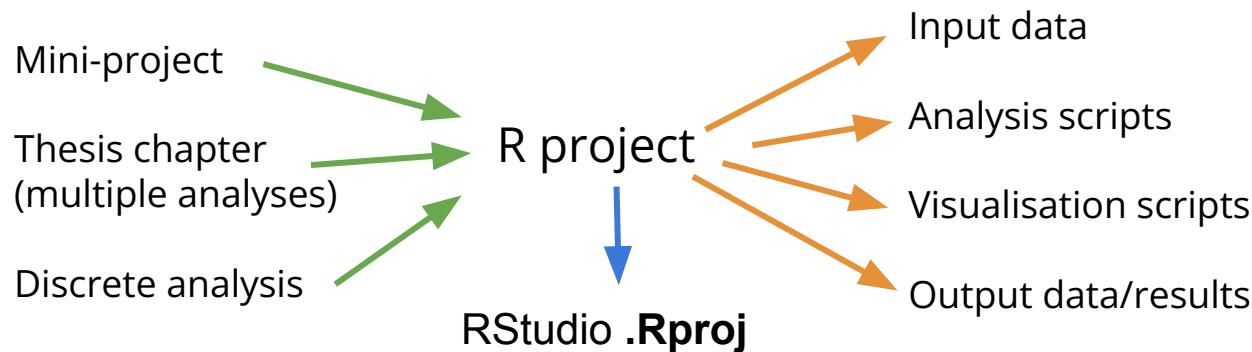
How you do organise your projects? scripts? folders?

Where do you store and access your data? results? figures?

Does it matter? Are your analysis/projects reproducible?

Can you organise your projects better and make life easier for future self/colleagues?

R projects



How you do organise your projects? scripts? folders?

Where do you store and access your data? results? figures?

Does it matter? Are your analysis/projects reproducible?

Can you organise your projects better and make life easier for future self/colleagues?

Reproducibility can be enhanced through intentionally organising projects with .Rproj, i.e. working in **project-oriented workflow**

Project-oriented workflow



2017/12/12

Jenny Bryan

I was honored to speak this week at the IASC-ARS/NZSA Conference, hosted by the Stats Department at The University of Auckland. One of the conference themes is to celebrate the accomplishments of Ross Ihaka, who got R started back in 1992, along with Robert Gentleman. My talk included advice on setting up your R life to maximize effectiveness and reduce frustration.

Two specific slides generated much discussion and consternation in #rstats Twitter:

If the first line of your R script is

```
setwd("C:\\Users\\jenny\\path\\that\\only\\I\\have")
```

I will come into your office and SET YOUR COMPUTER ON FIRE 🔥.

If the first line of your R script is

```
rm(list = ls())
```

I will come into your office and SET YOUR COMPUTER ON FIRE 🔥.

Jenny Bryan

@JennyBryan

Software engineer @rstudio, humane #rstats, adjunct prof @UBC where I created @STAT545, part of @ropensci, she/her

Vancouver, BC jennybryan.org

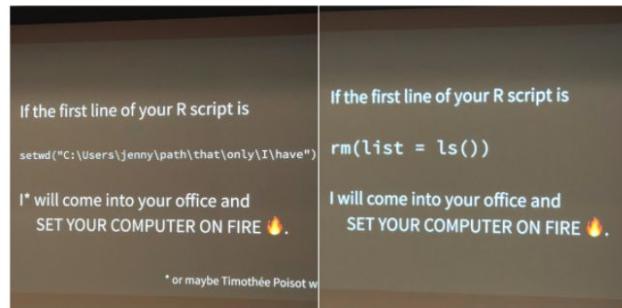


Hadley Wickham

@hadleywickham

Follow

The only two things that make @JennyBryan . Instead use projects + here::here() #rstats



4:50 PM - 10 Dec 2017

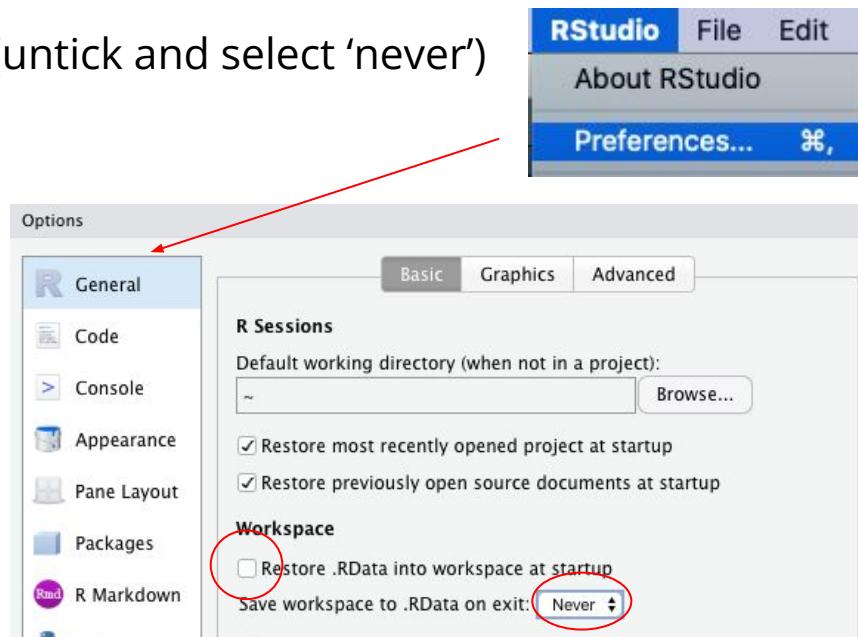
290 Retweets 950 Likes



Save your source code, not the workspace

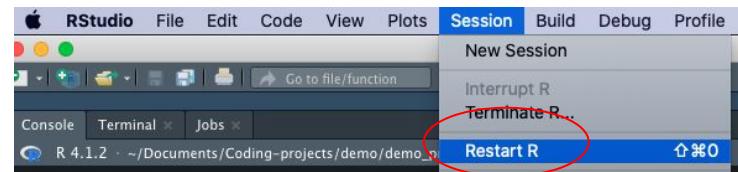
Do not save your .Rdata workspace:

(untick and select 'never')



Do not use `rm(list = ls())`

Restart R daily* to ensure a clean environment:



Save important objects and intermediates to files and scripts, in a modular way

If there is no source, it's not reproducible

Why setting directory does not work

Don't use `setwd()`

```
setwd("path/that/only/works/on/my/machine")
```

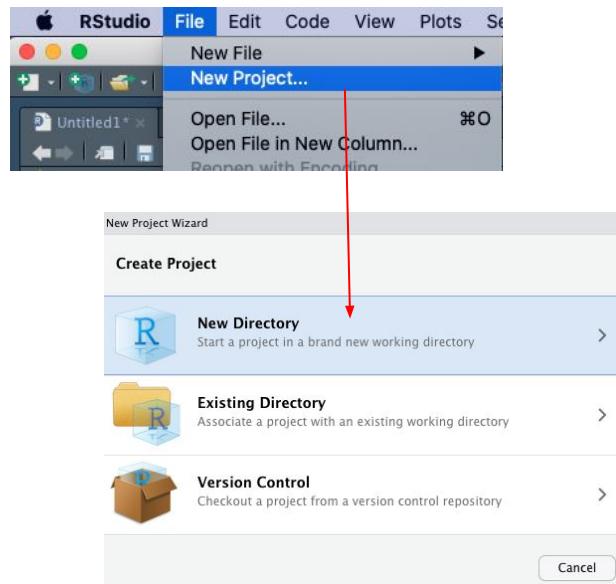
- Only works on your system
- Unlikely to work in a few years or on another computer
- Relying on hard-coded paths make a project easy to break and potentially not reproducible (shared data paths on the server is a different matter)

The solution to both:

- keep your work as an .Rproj (i.e. project-oriented workflow) to help you manage your workspace env and file paths

Project-oriented workflow

Use Rstudio / .Rproj for your data analysis projects



The screenshot shows the RStudio interface with a new project named 'demo_project1' created. The 'Create New Project' dialog is shown at the top right, with 'New Directory' selected. The main RStudio window shows the 'demo_project1 - RStudio' session. The 'Console' tab displays the R environment setup. The 'Global Environment' pane at the bottom right is empty, with the text '.Rproj file contains settings for the current project' overlaid. A red arrow points from the 'New Project Wizard' dialog to the 'demo_project1.Rproj' file in the 'Files' pane.

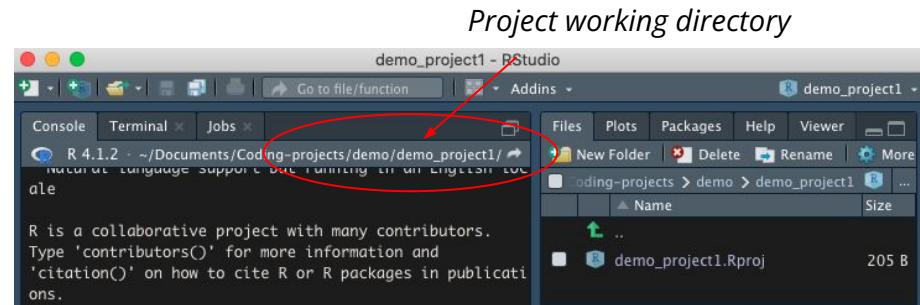
This means that you are essentially compartmentalizing your current project

Using .Rproj for your data analysis projects:

File system discipline: project directory stores all your data, scripts, figures

File path discipline: The working directory is set to the project directory (e.g. `demo_project1/`), so you don't need to `setwd()` or specify full paths to data (only internal subfolders that are relative to top directory)

Working directory intentionality: when working on project A, your working directory is set to project A's folder

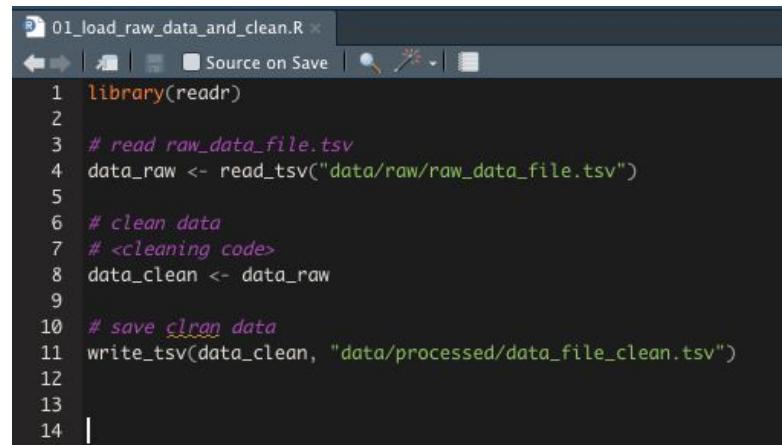


- The project creates everything it needs, within its workspace/folder, and touches nothing it did not create
- Any scripts are written assuming they will be run from a fresh R session within the project
- The project folder can be moved *anywhere*, and everything will still work (no paths will be broken)

You don't need to use setwd()

Keeping your work as an **.Rproj** will help you manage your file paths:

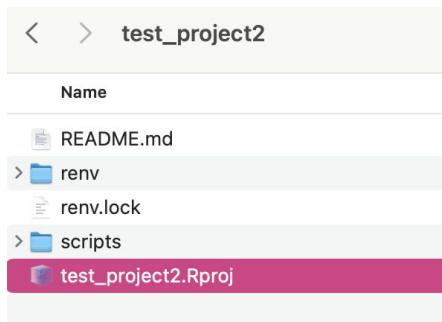
```
(base) [demo_project1] tree -L 3
.
├── README.md
└── data
    ├── processed
    │   └── data_file_clean.tsv
    └── raw
        └── raw_data_file.tsv
├── demo_project1.Rproj
└── scripts
    ├── 01_load_raw_data_and_clean.R
    ├── 02_explore_data.R
    ├── 03_run_analysis.R
    ├── 04_make_plots.R
    └── 05_project_summary.Rmd
```



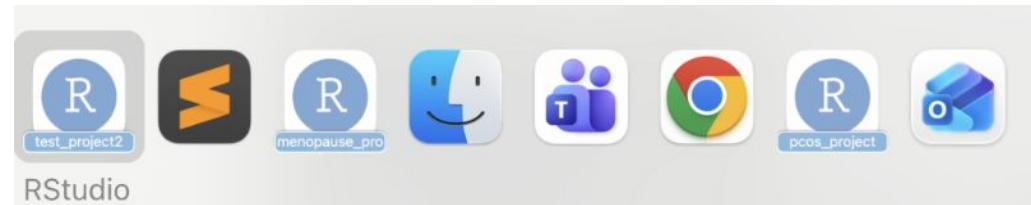
```
01_load_raw_data_and_clean.R
library(readr)
# read raw_data_file.tsv
data_raw <- read_tsv("data/raw/raw_data_file.tsv")
# clean data
# <cleaning code>
data_clean <- data_raw
# save clean data
write_tsv(data_clean, "data/processed/data_file_clean.tsv")
```

You don't need to use `rm(list=ls())`

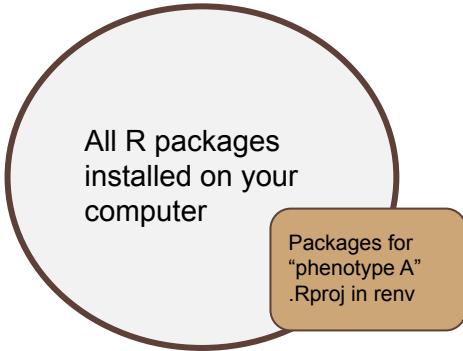
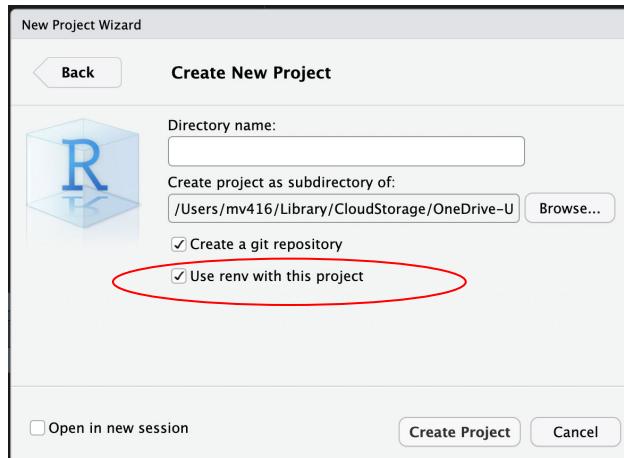
Click on .Rproj file in project directory to open a new clean session in your project:



You can also have multiple non-conflicting projects open at the same time:



renv for managing package versions within projects



renv package helps you create reproducible environments for your R projects



- Keeps track of all your installed packages
- Installs them in a new location (i.e. easy to share)
- Useful for working on UKB-RAP/AoU platforms (as a part of .Rproj)

Setting up your `renv`

```
install.packages("renv")
```

```
renv::init() - initialise your project env
```

NB this is the same as

Use renv with this project

This creates:

- **renv/**: stores packages for the project.
- **renv.lock**: records packages and the exact versions used
- **.Rprofile**: ensures renv activates when the project opens

		.Rprofile	26 B
		renv	
		renv.lock	402 B
		test_project1.Rproj	205 B

When `renv` is initialised, your project gets its own project library - a private folder (**renv/library/**) where packages are installed just for this project.

This means package versions are isolated and recorded for each project, so updating a package in one project won't break others.

This makes collaboration and reproducibility much easier.

```
renv::snapshot() - update your env with newly installed packages (saves to renv.lock)
```

```
renv::restore() - when project is re-opened, load all your required packages
```

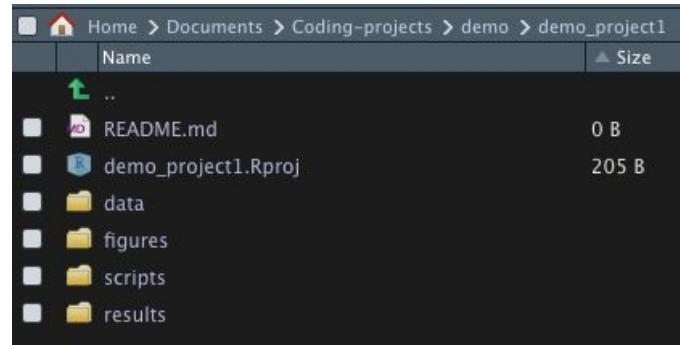
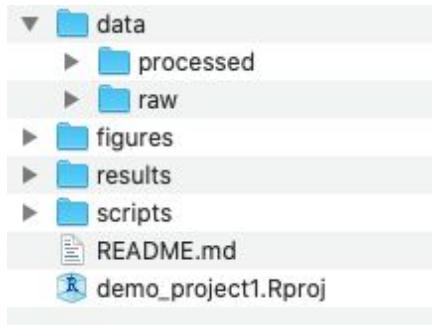
- should not be necessary to run in the local Rstudio, only when cloning a project from Git (eg on UKB-RAP)

Practical part (15 mins)

1. In your local Rstudio, create a new Project (e.g. *test_project1*) with **renv** and **git** enabled in a new session (window)
 - a. Create a new R script in the project folder; install package “sp” in it
 - b. Check the “sp” package loads in your project, then *renv::snapshot()* it
 - c. Check the “sp” package loads in your other ongoing open Rstudio session
 - d. Close your new project Rstudio window
 - e. Re-open by clicking on *test_project1.Rproj* file
 - f. Test the “sp” package still loads (do not install again!)
2. Download ‘*demo_project1.zip*’ from the email and uncompress it
 - a. Open the project and explore the folders/files
 - b. Has the project been created with *renv*?
 - c. Review *00_set_up_renv.R*; restore the env and install a package a new package (eg “sp”)

1.1. Hands-on tips for your R projects

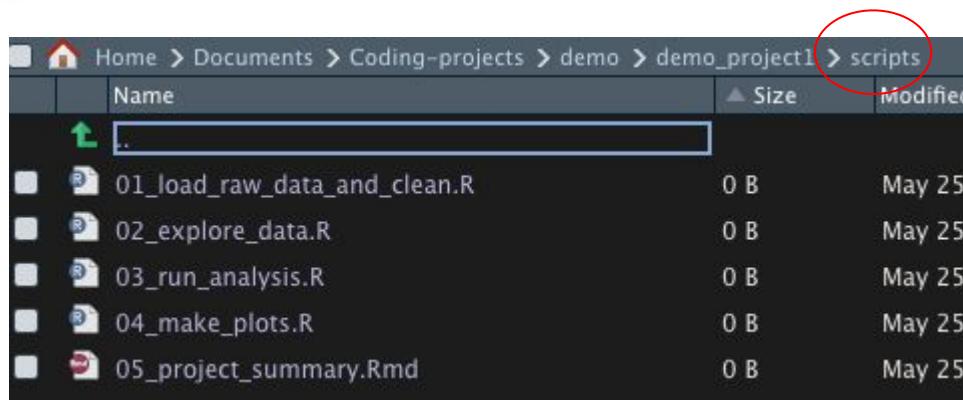
Organise your projects intentionally



Take advantage of default ordering

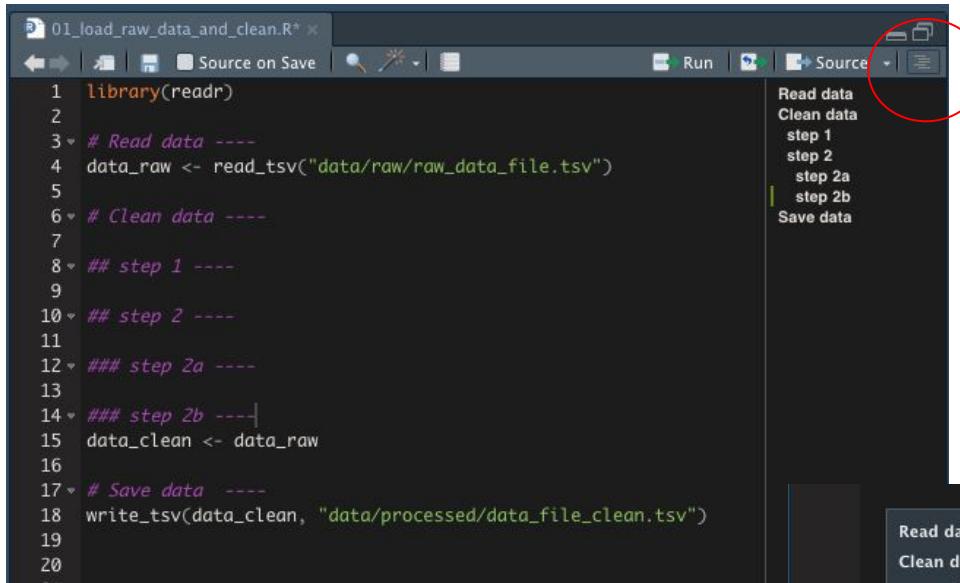
-  01_load_raw_data_and_clean.R
-  02_explore_data.R
-  03_run_analysis.R ←
-  04_make_plots.R
-  05_project_summary.Rmd

Can have many parts of the analysis separately - save interim results as files and re-read them in the next script



"How to name files" by
Jenny Bryan - [link](#) (5 mins)

Name your code sections and use them for quick navigation



A screenshot of the RStudio interface. On the left is a code editor window titled "01_load_raw_data_and_clean.R". The code is a script for data processing:library(readr)
Read data ----
data_raw <- read_tsv("data/raw/raw_data_file.tsv")

Clean data ----
step 1 ----
step 2 ----
step 2a ----
step 2b ----
data_clean <- data_raw

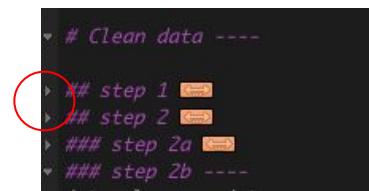
Save data ----
write_tsv(data_clean, "data/processed/data_file_clean.tsv")

```
On the right side of the interface, there is a vertical navigation bar. A red circle highlights the "Source" tab at the top of this bar. Below it is a dropdown menu with the following items: "Read data", "Clean data", "step 1", "step 2", "step 2a", "step 2b", and "Save data".
```

- Use section headings:

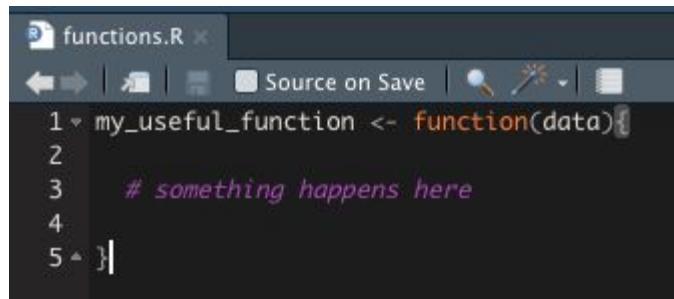
```
# section ----  
## subsection ----  
### subsubsection ----
```

- Great for navigating in long scripts
- Can fold sections



Jump to function definition or open data frame

```
• ## step 2 ----  
  
output <- my_useful_function(input)
```



To navigate to a function definition or review a function from a package:

- **Cmd + click** on the function name

> to see what it does internally / check arguments

Keyboard shortcuts

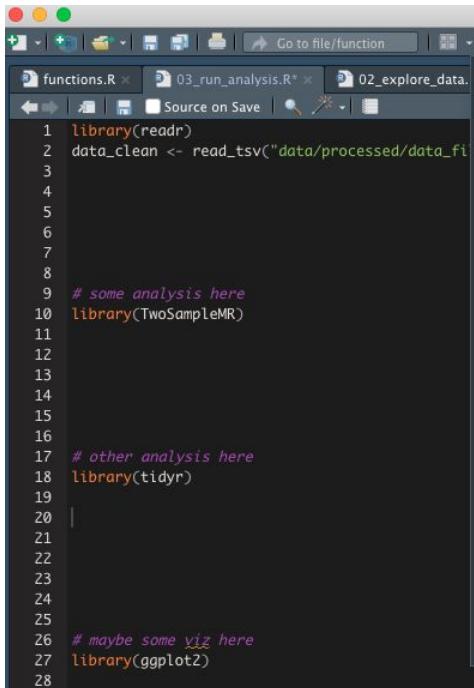


- (option + Enter)
- (option/alt + " - ")
- (%) (control + shift + M)
- (control + shift + I)

...

Keyboard Shortcuts		
Name	Shortcut	Scope
Insert Pipe Operator	Ctrl+Shift+M	Editor

Move all libraries to the top



```
library(readr)
data_clean <- read_tsv("data/processed/data_file_clean.tsv")
# some analysis here
library(TwoSampleMR)

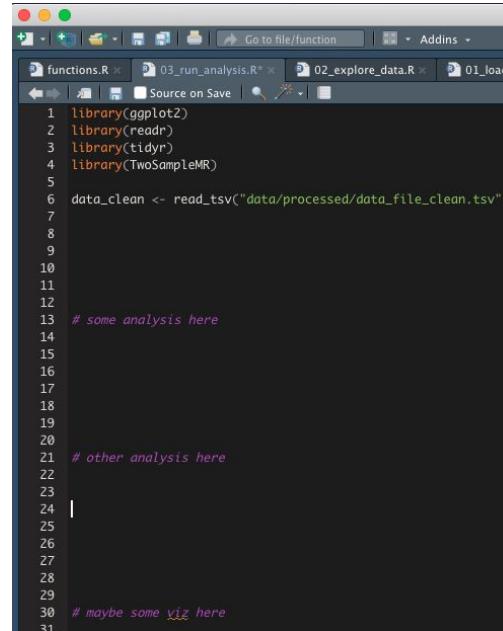
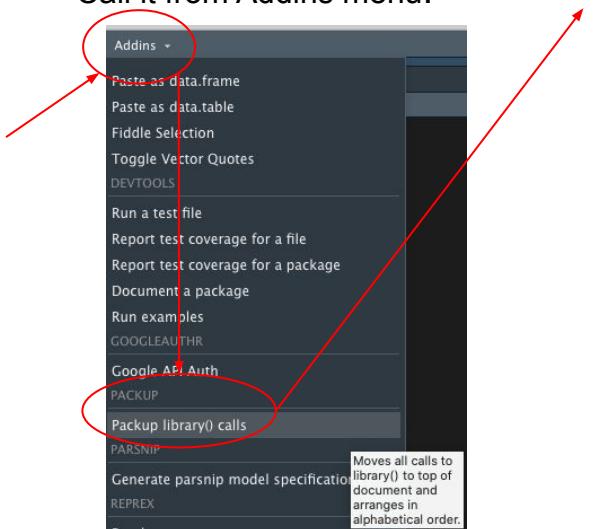
# other analysis here
library(tidyverse)

# maybe some viz here
library(ggplot2)
```

Install *packup* add-in:

```
devtools::install_github("milesmc Bain/packup")
```

Call it from Addins menu:

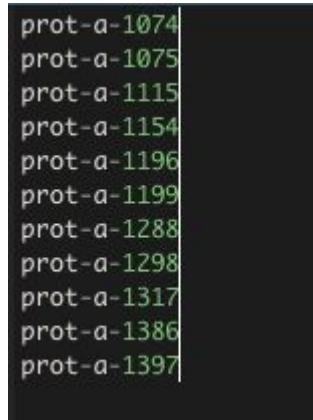


```
library(ggplot2)
library(readr)
library(tidyverse)
library(TwoSampleMR)

data_clean <- read_tsv("data/processed/data_file_clean.tsv")
# some analysis here
# other analysis here
# maybe some viz here
```

Vertical selection

(hold *option* or *alt* and drag cursor down to select vertically)



```
prot-a-1074
prot-a-1075
prot-a-1115
prot-a-1154
prot-a-1196
prot-a-1199
prot-a-1288
prot-a-1298
prot-a-1317
prot-a-1386
prot-a-1397
```

A screenshot of a terminal window with a vertical selection highlighted by a thick black border. The selected text consists of eleven lines of code, each starting with "prot-a-".

Great for e.g.

- commenting out a block of code with #
- adding " " around a column of ids

Any other R tips to share?

2. Using Git to track changes in your R projects

<https://happygitwithr.com/>

Checklist before we start

- GitHub account
- Personal access token
- Git is available on your laptop
- SSH-key (linking Git on your laptop with your GitHub account)

Git brief intro

Git - tool for code version control

- Tracking code changes
- Keeping older versions of the script
- Coding collaboration (with others or yourself in multiple locations)
- Tracking who and when made changes



If used consistently!

Why use it?

- Can facilitate data/code integrity
- Improves reproducibility (e.g. keeps record of changes)
- Enables code sharing with colleagues / as a part of publication
- Important skill for anyone working with data (e.g. can showcase your work during job applications)

Github / Gitlab etc

- Platforms where you can store your coding projects using Git

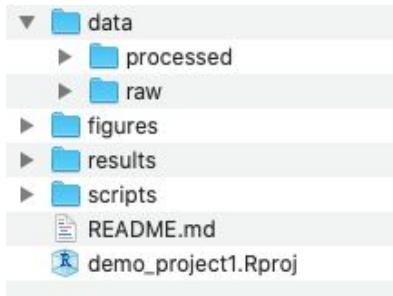
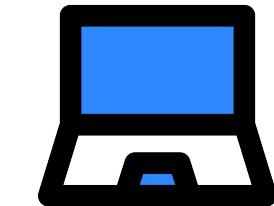


git

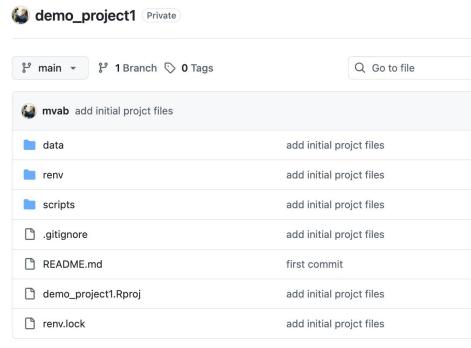


Why might you want to use git? (scenario 1)

Your R project
on local Mac



Your R project
as GitHub repo

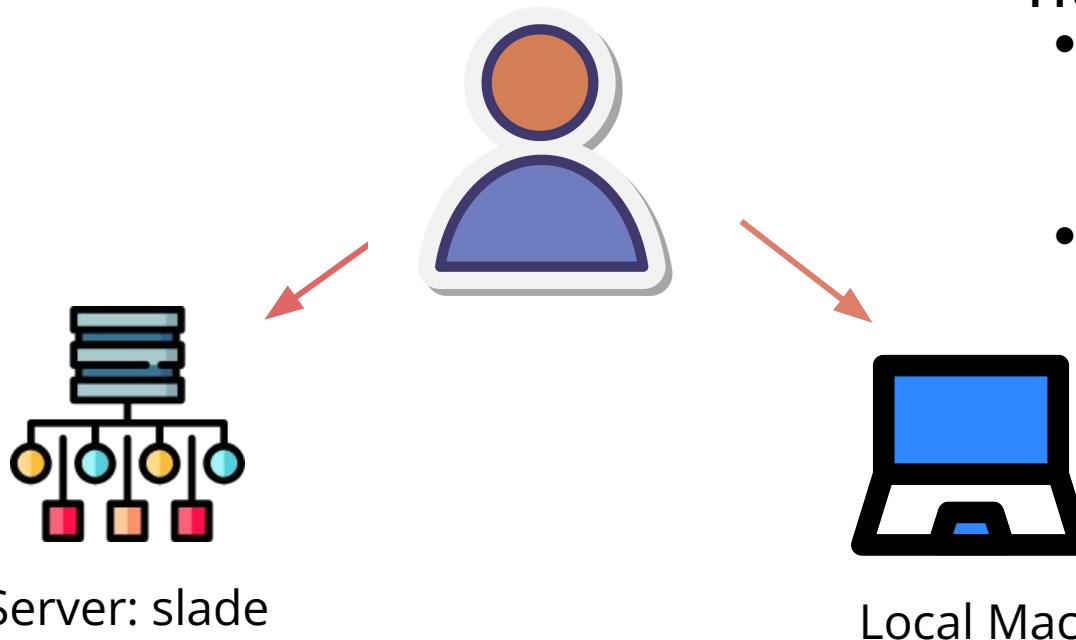


A screenshot of a GitHub repository page for "demo_project1". The repository is private. The commit history shows the following activity:

- mvab add initial project files (first commit)
- add initial project files

- Using git helps you keep track of your daily changes
- All your work is backed-up (as long as you do it regularly and intentionally)
- “Collaborating with yourself” in one place

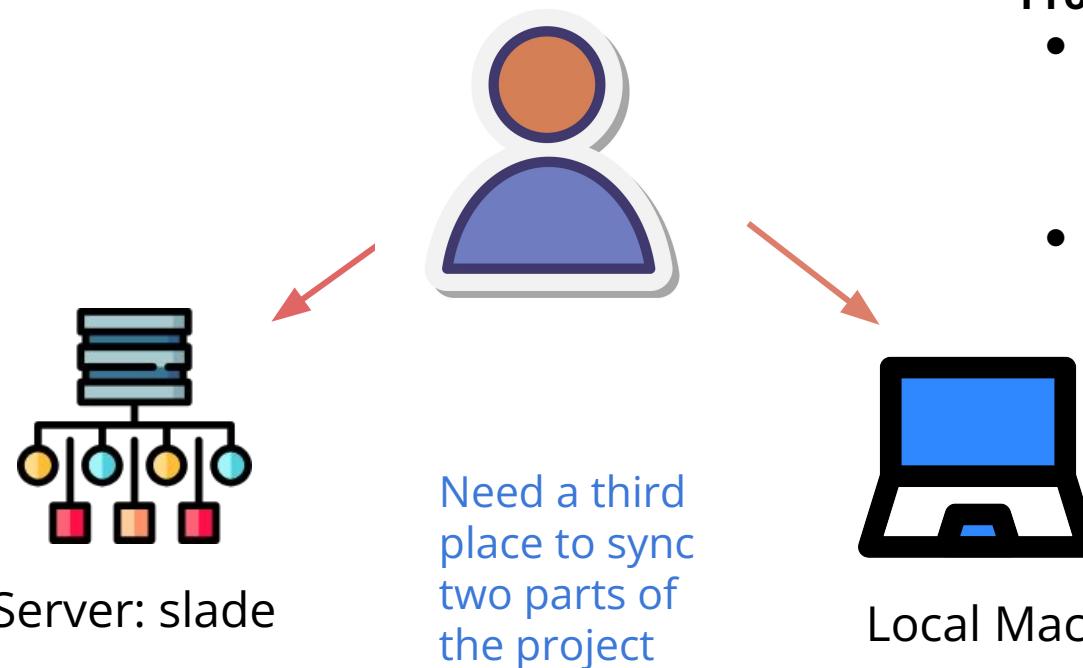
Why might you want to use git? (scenario 2)



Problem:

- The same project is both locations (e.g due to data restrictions / tool availability)
- Don't want to copy over scripts all the time, as it might get messy!

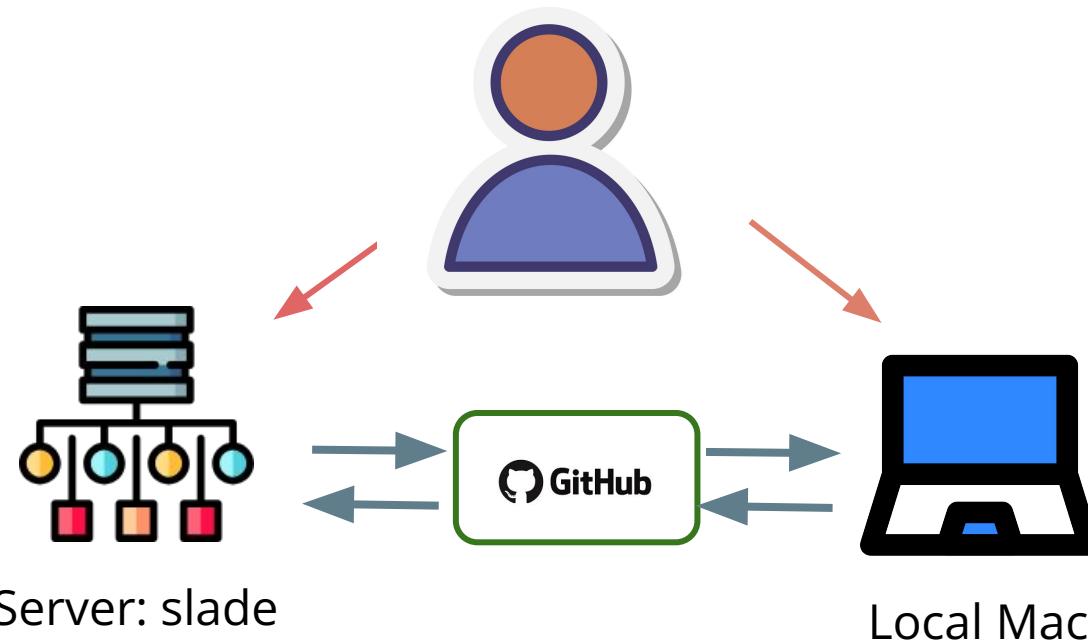
Why might you want to use git? (scenario 2)



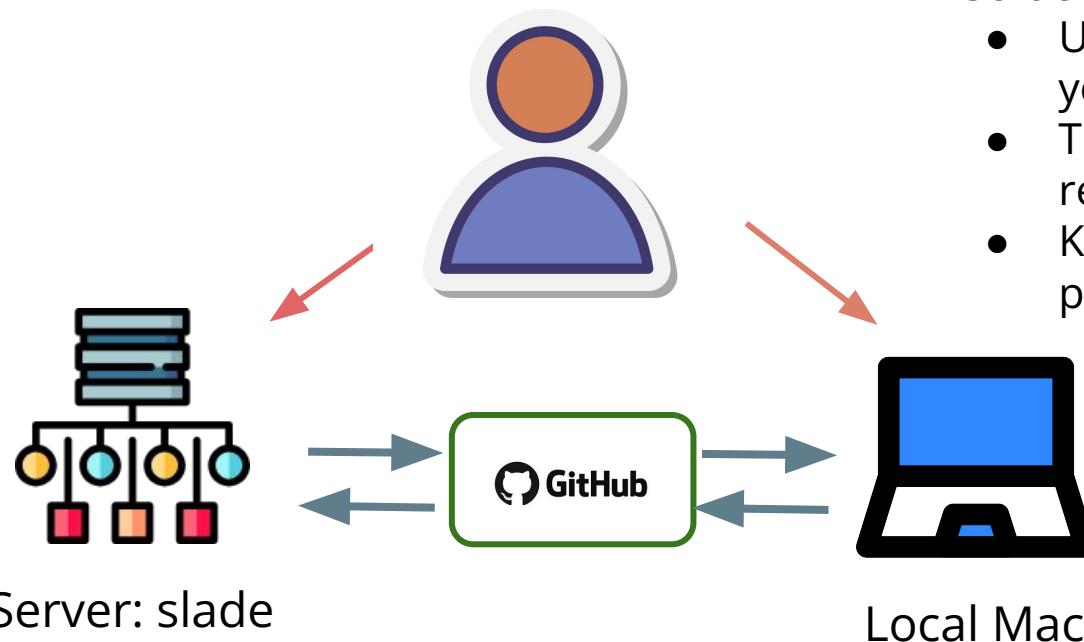
Problem:

- The same project is both locations (e.g due to data restrictions / tool availability)
- Don't want to copy over scripts all the time, as it might get messy!

Why might you want to use git? (scenario 2)



Why might you want to use git? (scenario 2)



Solution:

- Use git to “collaborate with yourself” in two places
- This enhances project reproducibility
- Keeps all analysis code in one project folder/git repository

Why might you want to use git? (scenario 3)

Your R project
on



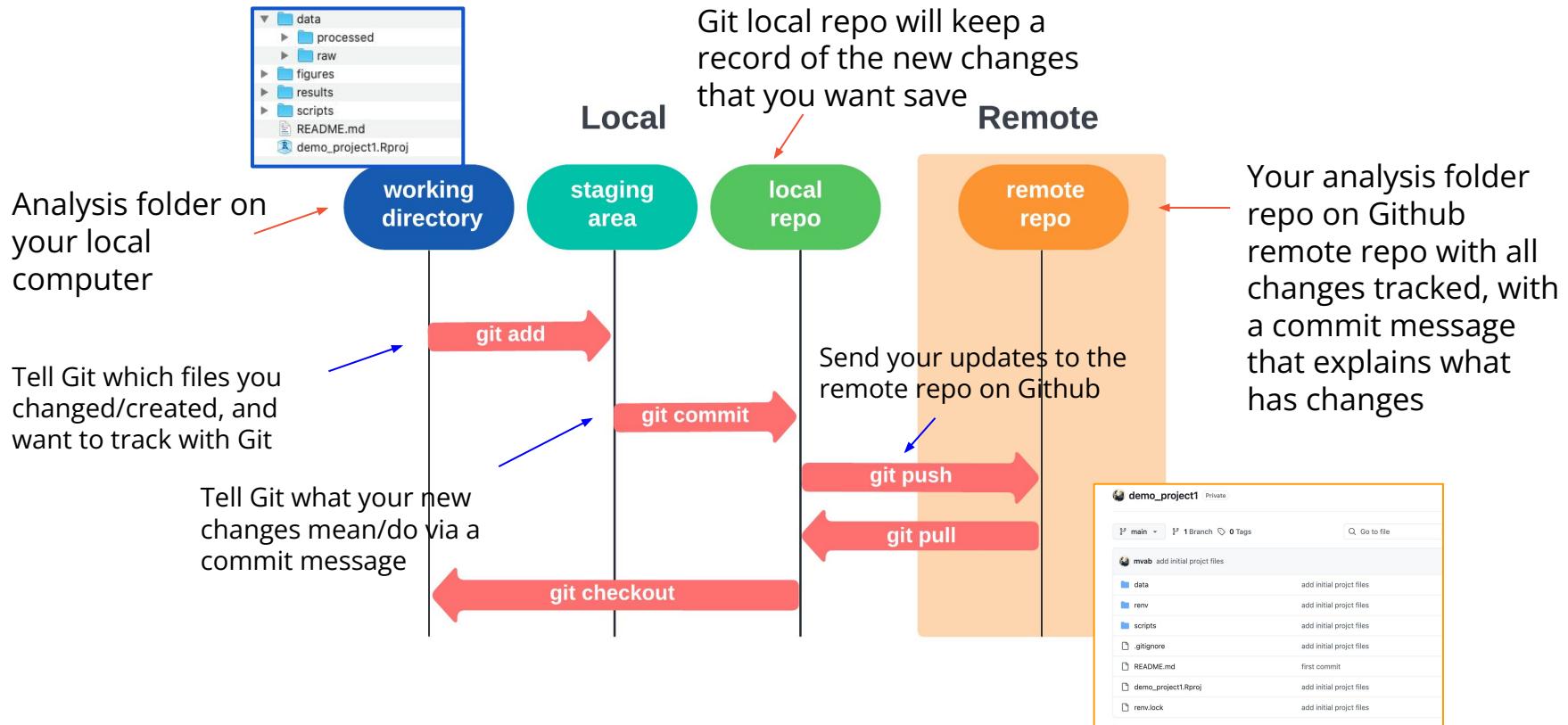
biobank



Your R project
as GitHub repo

- Using git helps you keep track of your daily changes
- All your work is backed-up (you are 'forced' to use Git daily to back up your work, because you start a new R instance every time you work on your project)

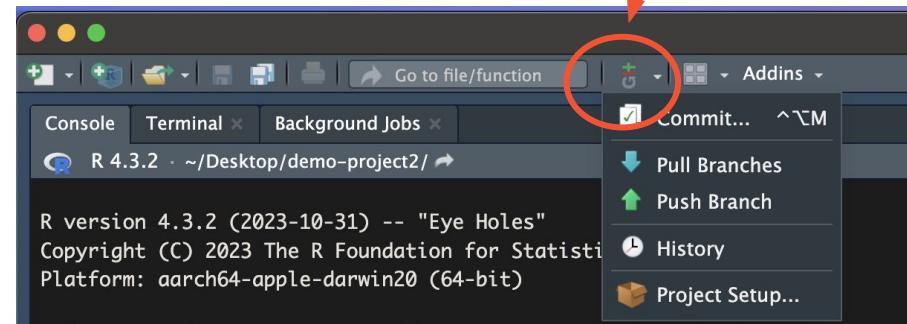
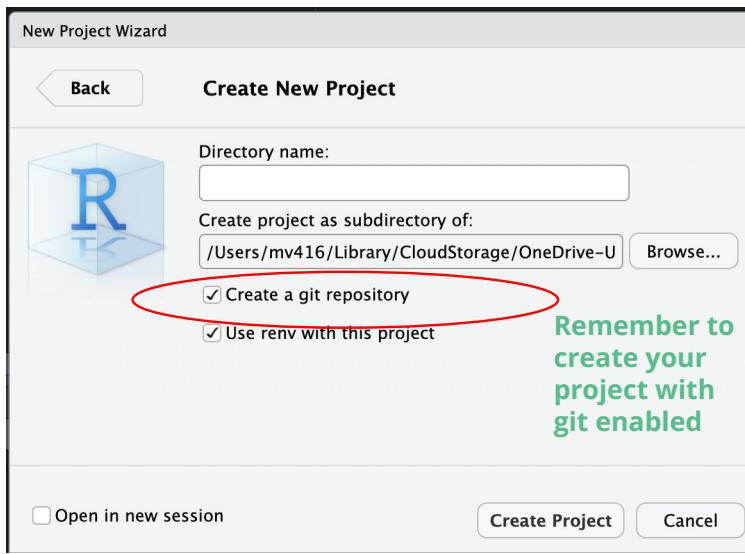
Basic git commands / actions



Follow along if you can

Using git with your .Rproj

- Rstudio user interface makes it easy to get started with git
- (but also can use the terminal to run git commands)



>> Next need to link your local project folder to a GitHub repo

Create remote repo

After logging in your Github account:

To create a new repository:

Repositories > New

Create a new repository

Repositories contain a project's files and version history. Have a project elsewhere? [Import a repository](#).

Required fields are marked with an asterisk (*).

1 General

Owner * mvab Repository name * test_project1

Give repo the same name as your .Rproj folder

test_project1 is available.

Great repository names are short and memorable. How about [legendary-memory](#)?

Description

0 / 350 characters

2 Configuration

Choose visibility * Public

Don't change any configurations to keep it simple (except visibility)

Start with a template No template

Add README Off

Add .gitignore No .gitignore

Add license No license

Create repository

test_project1 is available.

Great repository names are short and memorable. How about [legendary-memory](#)?

0 / 350 characters

Public

No template

Off

No .gitignore

No license

GitHub makes it easy to link your .Rproj to the new repo:

The image shows a GitHub repository creation interface and an RStudio terminal window.

GitHub Repository Creation Interface:

- Header:** test_project1 Public
- Set up GitHub Copilot:** Use GitHub's AI pair programmer to autocomplete suggestions as you code. [Get started with GitHub Copilot](#)
- Add collaborators to this repository:** Search for people using their GitHub username or email address. [Invite collaborators](#)
- Quick setup — if you've done this kind of thing before:**
 - Set up in Desktop or HTTPS SSH https://github.com/mvab/test_project1.git
 - Get started by [creating a new file](#) or [uploading an existing file](#). We recommend every repository include a [README](#), [LICENSE](#), and [.gitignore](#).
- ...or create a new repository on the command line:**

```
echo "# test_project1" >> README.md
git init
git add README.md
git commit -m "first commit"
git branch -M main
git remote add origin https://github.com/mvab/test_project1.git
git push -u origin main
```

A red circle highlights this section, and a red arrow points from it to the RStudio terminal.

Please note - the 'password' it will ask for is your 'personal access token'

- ...or push an existing repository from the command line:**

```
git remote add origin https://github.com/mvab/test_project1.git
git branch -M main
git push -u origin main
```

RStudio Terminal:

- Console Terminal Background Jobs
- Terminal 1 ~ /Downloads/test_project1
- (base) FY2JTCXL4H:test_project1 mv416\$

A red arrow points from the GitHub command line section to the RStudio terminal, indicating where the copied command should be pasted.

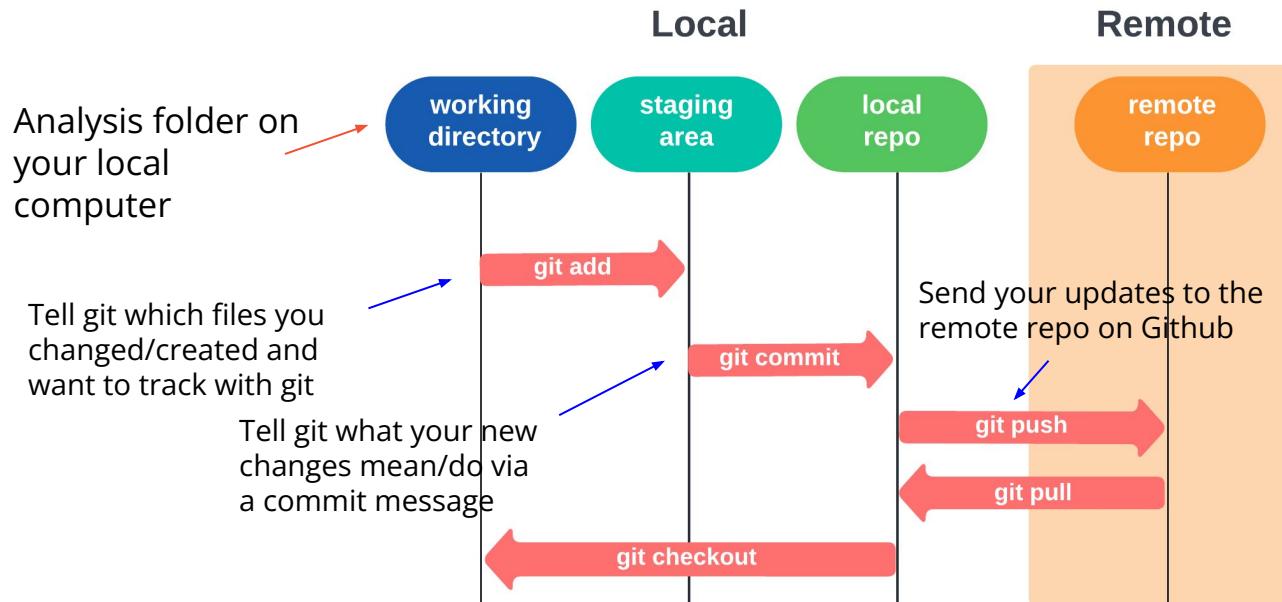
Remote repo after first commit

A screenshot of a GitHub repository page for a private project named "test_project2". The repository has one branch, "main", and one commit by user "mvab" titled "first commit". The commit hash is 40ea328, it was made at "now", and there is 1 Commit. The file "README.md" is listed under the commit, showing the commit message "first commit" and the date "now". The file "README" is also listed but is not yet committed. The repository name "test_project2" is displayed at the bottom.

Only
README file
has been
added

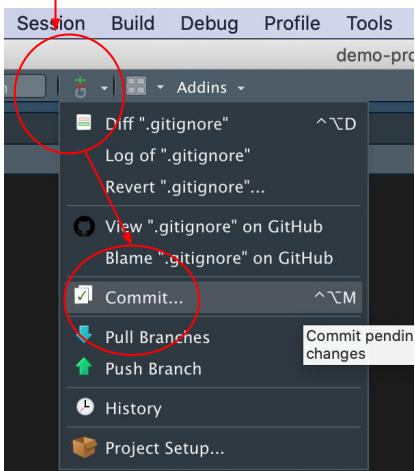
```
echo "# test_project1" >> README.md  
git init <- initialises your project repo  
git add README.md  
git commit -m "first commit"  
git branch -M main  
git remote add origin https://github.com/mvab/test_project1.git  
git push -u origin main
```

} initialises your project repo

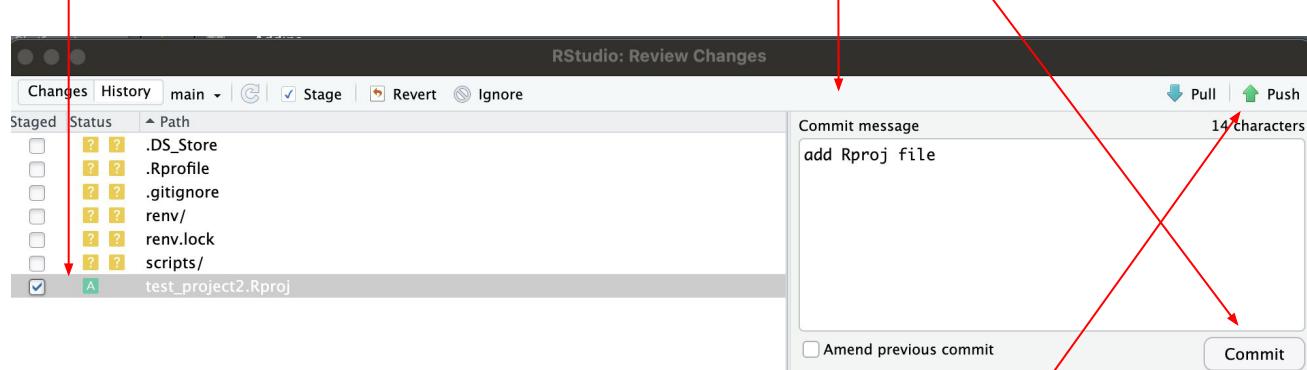


Time to add other files!

- 1) Git button -> Commit



- 2) Select the files you want Git to add (track):



- 3) write your 'commit message', ie description of your change
(tip: write it as an instruction), then press "**Commit**"

- 4) Press "**Push**" to send the change to remote repo

Your change has been added:

 **test_project2** Private Watch 0

 main  1 Branch  0 Tags Go to file Add file Code

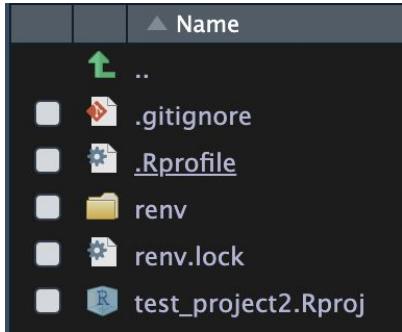
File	Commit Message	Time
 mvab add Rproj file	85cc036 · 18 minutes ago	2 Commits
 README.md	first commit	37 minutes ago
 test_project2.Rproj	add Rproj file	18 minutes ago

 **README** 

test_project2



.gitignore file - list files you don't want to store in remote repo

A screenshot of a code editor window titled ".gitignore*". The file contains the following content:

```
1 .Rproj.user
2 .Rhistory
3 .RData
4 .Ruserdata
5
6 data/
7 .DS_Store
```

A red arrow points from the "Name" header of the first screenshot towards the ".gitignore" file in the code editor.

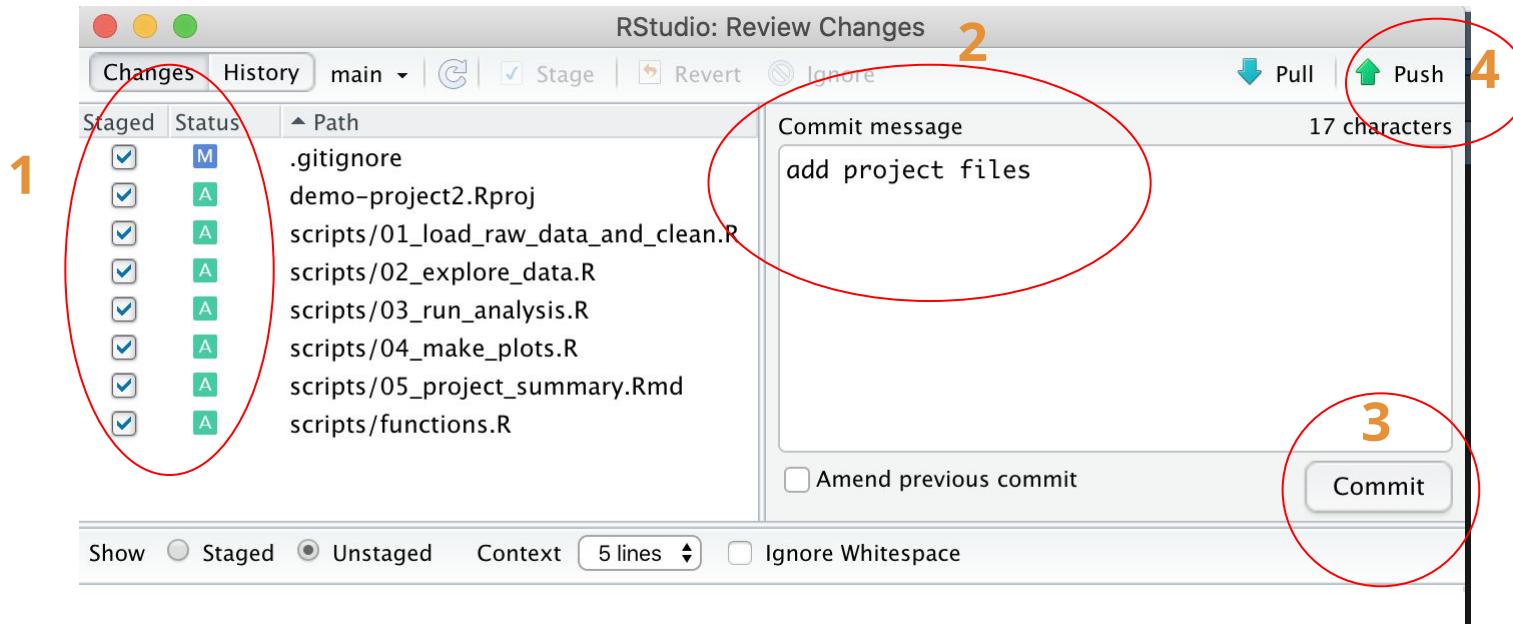
Add **data/** folder to *.gitignore* file so that your data files (if large or sensitive) are not committed to your project repo on Github

- list files and folders in your project folder in the *.gitignore* file - they will be ignore by Git (this prevents them from being accidentally added to GitHub repo)
- save and '**add**' file
- **commit** and **push** the updated *.gitignore* file to your repo

Now add more files:

Commit changes:

Add message:



**test_project2**

Private

Watch 0

main ▾

1 Branch

0 Tags

Go to file

t

Add file ▾

Code ▾

**mvab** add all scripts

6da936d · now

5 Commits



renv

add renv files

now



scripts

add all scripts

now



.Rprofile

add renv files

now



.gitignore

add gitignore

1 minute ago



README.md

first commit

3 days ago



renv.lock

add renv files

now



test_project2.Rproj

add Rproj file

3 days ago

README

test_project2

Your changes
on Github:

Practise adding a specific change to a script:

1

```
library(readr)
# Read data ----
data_raw <- read_tsv("data/raw/raw_data_file.tsv")
# Clean data ----
## step 1 ----
c = 1 + 2 # adding this change
## step 2 ----
output <- my_useful_function(input)
### step 2a ----
```

2

You will see what's changed here

3

4

Commit message
calculated c

Amend previous commit

Commit

Write a message that describes your change, then commit and push

```
@@ -5,10 +5,13 @@ data_raw <- read_tsv("data/raw/raw_data_file.tsv")
# Clean data ----
## step 1 ----
c = 1 + 2 # adding this change
## step 2 ----
output <- my_useful_function(input)
### step 2a ----
```

main ▾

demo-project2 / scripts /



mvab calculated c

..

01_load_raw_data_and_clean.R	calculated c
02_explore_data.R	add project files
03_run_analysis.R	add project files
04_make_plots.R	add project files
05_project_summary.Rmd	add project files
functions.R	add project files

Good practice principles

1. **Add** and **commit** your changes often and in small chunks
 - o E.g. separate commits for different files and sections in your analysis steps
2. Write descriptive commit messages of what you've changed
 - o "fix typos" is ok, but "update" is not
3. Use `.gitignore`
4. Keep a project README
 - o describes what your scripts do, and where external data is stored

3. R projects on UKB-RAP/All of Us platforms with Git

Start two new Rstudio instances now!

Working in Rstudio in on-demand instances

The problem:

- You are forced to scratch from a clean env every time you work a project:
 - All packages need to be installed every time
 - All intermediate data tables need to be regenerated
 - To save progress, you have to copy your R/Rmd scripts back and forth with your User project



The solution:

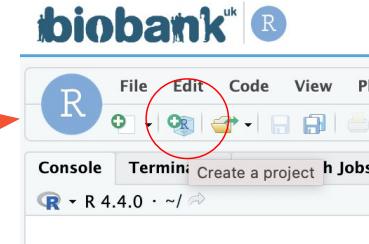
- Organise your R code as *Rproj* with *renv*, and use *Git* to back it up:
 - *renv::restore()* brings back all your packages
 - Intentional and modular file organisation helps you go back your intermediate files quicker
 - Committing your ongoing changes to your project repo on GitHub keeps your work backed up
 - *git clone* brings back your entire project to your clean Rstudio instance

Follow along if you can

You already know how to make it work

Start a new project in Rstudio instance
(instance #1 you started):

- 1) Create a new project (with renv + git)
- 2) On Github, create a new private repo
- 3) On Rstudio Terminal, authenticate:
 - o *git config --global user.email "you@example.com"*
 - o *git config --global user.name "user"*
- 4) Run the set up commands from GitHub on Rstudio Terminal
 - o use your token as password!
- 5) Push all other files to your remote repo
- 6) Upload or create *00_renv_setup.R* file project folder
(attached in the email)
- 7) Add any other dummy R files you want

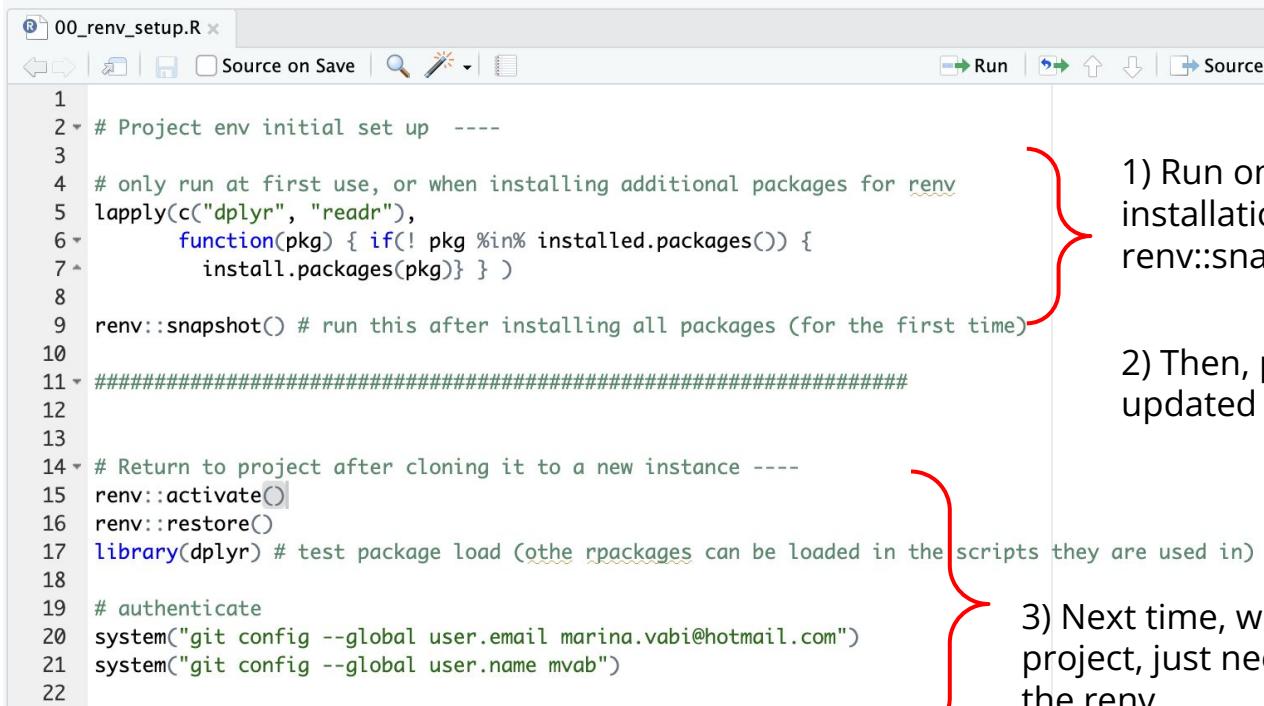


...or create a new repository on the command line

```
echo "# test_project_ukrap" >> README.md
git init
git add README.md
git commit -m "first commit"
git branch -M main
git remote add origin https://github.com/mvab/test_project_ukrap.git
git push -u origin main
```

A screenshot of a GitHub staging area. It's a table with columns for 'Staged', 'Status', and 'Path'. There are seven files listed, each with a green checkmark in the 'Staged' column and an 'A' in the 'Status' column. The paths are: '.Rprofile', '.gitignore', 'renv.lock', 'test_project_ukrap2.Rproj', 'renv/.gitignore', 'renv/activate.R', and 'renv/settings.json'. The 'renv.lock' file is currently selected, as indicated by a grey background.

renv setup



The screenshot shows an RStudio interface with a code editor containing the file `00_renv_setup.R`. The code is as follows:

```
1 # Project env initial set up ----
2
3 # only run at first use, or when installing additional packages for renv
4 lapply(c("dplyr", "readr"),
5       function(pkg) { if(! pkg %in% installed.packages()) {
6         install.packages(pkg)} } )
7
8
9 renv::snapshot() # run this after installing all packages (for the first time)
10 #####
11 #####
12 #####
13 #####
14 # Return to project after cloning it to a new instance ----
15 renv::activate()
16 renv::restore()
17 library(dplyr) # test package load (the rpackages can be loaded in the scripts they are used in)
18
19 # authenticate
20 system("git config --global user.email marina.vabi@hotmail.com")
21 system("git config --global user.name mvab")
```

A red curly brace on the right side of the code highlights the first two sections (lines 1-8). Another red curly brace on the right side of the code highlights the last section (lines 14-21).

1) Run once, only on the first package installation only + `renv::snapshot()` to save it to `renv.lock`

2) Then, push the new file and the updated `renv.lock` to GitHub:

Staged	Status	Path
<input checked="" type="checkbox"/>	A	00_renv_setup.R
<input checked="" type="checkbox"/>	M	renv.lock

3) Next time, when going back to this project, just need to activate and restore the renv

Tip: keep a copy of `00_renv_setup.R` file in all projects

Ending your R instance working on Rproj

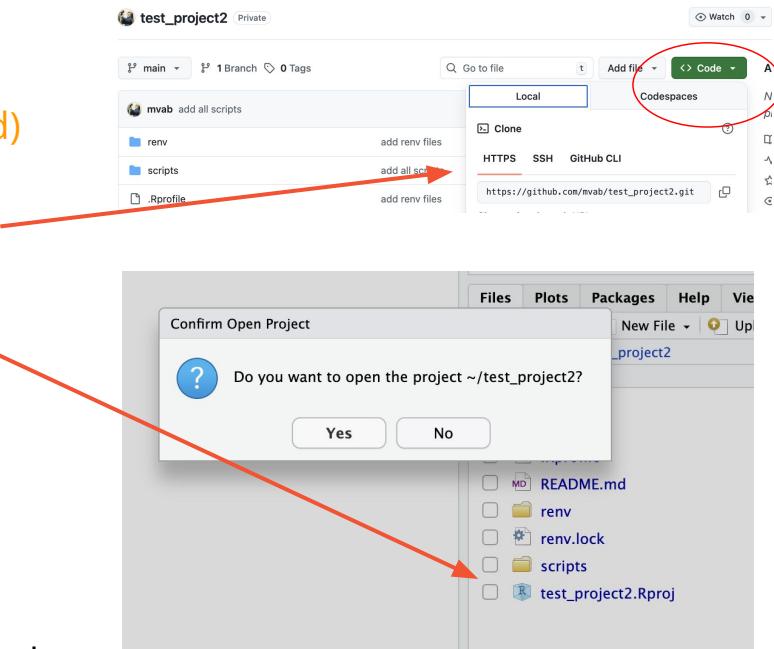
At the end of your session, save and push all your changes to GitHub (check it's all there)

- Remember to run `renv::snapshot()` - so that the `renv.lock` file is updated (push it to GitHub too!)
- Be especially careful if working with individual level data (tip: only save your data files in to folder `data/`, and list that folder in your `.gitignore` - so it does not get added to GitHub by accident)
- Once all code you wrote in the current session is on GitHub, it's safe to terminate **instance #1**

Returning to your R project

- 1) Starting from a clean instance (**instance #2 you started**)
- 2) On Rstudio terminal, *git clone* your project
 - o *git clone https://github.com/mvab/test_project2.git*
 - o Authenticate with your token
- 3) **Important:** click on Rproj file in the project folder to activate the project
- 4) Open *00_renv_setup.R* file and test that *library(dplyr)* currently does not work!
- 5) In *00_renv_setup.Rrun*:
 - o *renv::activate()*
 - o *renv::restore()*
 - o Test dplyr again!

Any other packages you install - remember to run *renv::snapshot()* - so that the *renv.lock* file is updated (and push to your GitHub repo)





Summary: using .Rproj for organising work

- “Reproducible project in R” means:
 - **File system discipline:** all files related to a single project are stored in a designated folder;
 - **Working directory discipline:** intentionally work in project directory when opening Rproj
 - **File path discipline:** paths are relative to the project directory (not hard-coded full paths!)
 - **Daily work habit:** Restarting R often and re-running your script under development from the top will help you catch issues early on
 - **Using git for version control**
- Practising these habits together will give you the biggest pay-off
 - Reproducing your analyses will be easy
 - Organising your projects will help you make sense of them in 6/12/etc months
 - Can move your project anywhere or share it with anyone without changing paths

Final thoughts / disclaimers

- Project-oriented workflow in R is not suitable/applicable to every scenario
 - Sometimes data is stored externally, and can't moved/copied (so you can't use within-project paths, but potentially can use symbolic links)
- Not all work is done interactively in Rstudio
 - Some people use R from the terminal on the server - again, because of data access/size
 - Some analyses are computation-heavy and require to be submitted as scripts / run in parallel on server
- If your current workflow with `setwd()` works for you and your colleagues, that's totally fine, but consider future-proofing! ;)

Recommended and used resources

<https://www.tidyverse.org/blog/2017/12/workflow-vs-script/>

<https://richpauloo.github.io/2018-10-17-How-to-keep-your-R-projects-organized/>

<https://www.rforecology.com/post/organizing-your-r-studio-projects/>

<https://kkulma.github.io/2018-03-18-Prime-Hints-for-Running-a-data-project-in-R/>

<https://rstats.wtf/project-oriented-workflow.html>

<https://appsilon.com/rstudio-shortcuts-and-tips/>

<https://datacornering.com/my-favorite-rstudio-tips-and-tricks/>

<https://happygitwithr.com/>

<https://rstudio.github.io/renv/articles/renv.html>

<https://rstats.wtf/>

Thank you!