

open science
open access
open research data
open source
→ **collaboration and**
reproducibility

GitHub

platform / server to store, share and collaborate
on projects

Git

a version control system to store different versions of your work and which connects RStudio with GitHub

RStudio

software using **R** programming language for
statistical analyses (and much more:)

let's put it to practice!

- create a new repository on GitHub

in github.com



→ Repositories

→ New repository

→ Repository name *mykosym*

→ Choose visibility *public/private*

→ Create repository

- use URL to open it as a new project in RStudio

in github.com



→ Repositories

→ *mykosym*

Quick setup — if you've done this kind of thing before

 Set up in Desktop

or

HTTPS

SSH

<https://github.com/mo0org/mykosym.git>



Get started by [creating a new file](#) or [uploading an existing file](#). We recommend every repository include a [README](#), [LICENSE](#), and [.gitignore](#).

copy link to your repo

in rstudio

→  Project: (None) ☐

→ New Project...

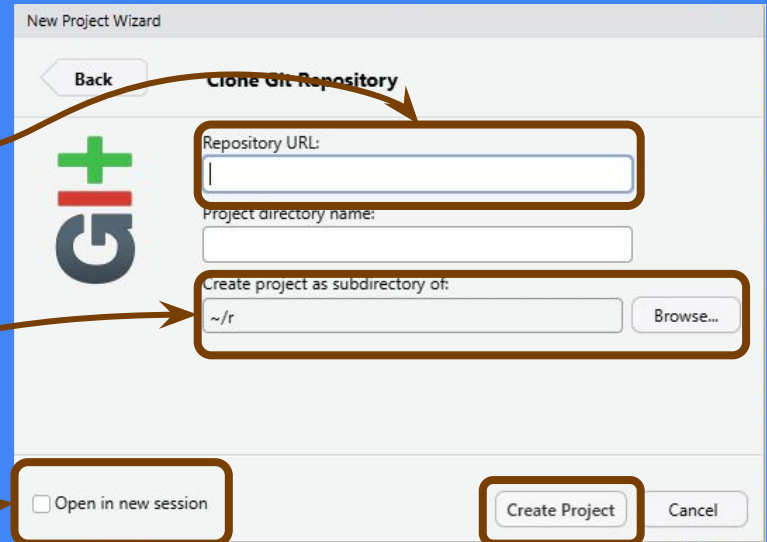
→  Version Control

→ Git

insert URL from clipboard


choose where to store the
project on your local repo

one open session= one project



New Project Wizard

Back Clone Git Repository



Repository URL:


Project directory name:

Create project as subdirectory of:
 Browse...

☐ Open in new session

Create Project Cancel

- create files and scripts in the R.project

in your file explorer 
→ open the project file



→ add files (and your datasets, existing r.scripts, etc.)

```
fs::dir_tree()
```

```
data
├── raw_data_csv
│   ├── cidlo_DN_data.csv
│   ├── OTU_test_14_10_25.csv
│   ├── root_biomass.csv
│   ├── soil_chem.csv
│   ├── vegetace_cs.csv
│   ├── vegetace_dn.csv
│   ├── vegetace_sv.csv
│   └── vegetation_raw.csv
├── raw_data_xls
│   ├── colonisation.xlsx
│   ├── dna_root.xlsx
│   ├── dna_soil.xlsx
│   ├── info_sites.xlsx
│   ├── root_biomass.xlsx
│   ├── soil_chem.xlsx
│   └── teplota_moist.xlsx
├── outputs
│   ├── figures
│   ├── tables
│   └── writing
├── README.html
├── README.md
├── r_scripts
│   ├── analyses
│   │   └── vegetation.R
│   └── cleaning_data
│       ├── prep_otu_tbl.R
│       ├── prep_teplota_moist_tbl.R
│       └── prep_vegetation_tbl.R
└── seasonality_project.Rproj
```

- write a line of code to open a .csv dataset

in rstudio

- ↳ open a new script (Ctrl+Shift+N)
- ↳ save it to the script file (Ctrl+S)
- ↳ use the here() and read.csv() functions to open a dataset

```
data<-read.csv(here::here("path/dataset_name.csv"))
```

```
sep = ""  
header = TRUE  
dec = ""  
na.strings = ""
```

```
install.packages("here")
```

`data <-` store the following line in the object "data"

`read.csv()` R function to read .csv files (for help `?read.csv`)

`sep = ""` argument to the `read.csv()` function to specify which type of column separator is used in the .csv file ; , / .

`header = TRUE` argument to the `read.csv()` function to specify if the .csv file has a header or not **TRUE FALSE**

`dec = ""` argument to the `read.csv()` function to specify type of decimal is used in the .csv file , .

`na.strings = ""` argument to the `read.csv()` function to specify what to replace by NA space 0

`here::here("path/dataset_name.csv")` R function to find the path to the file we want to open (for help `?here::here`)

in the path we do not specify the whole path as in `setwd()` but only what is after the name of the project:

C:/Users/Laura/OneDrive -umontpellier.fr/Dokumenty/r/mykosym/data/raw_data

- write a line of code to save a dataset to .csv

in rstudio (opened script where we have uploaded an initial dataset and we have created a subset of it)

```
data<-read.csv(here::here("path/dataset_name.csv"))  
subset<-data[,2:10]
```

↪ use the `here()` and `write.csv()` functions to create a new dataset in your repo

```
write.csv(subset,here::here("path"),subset_name.csv)
```

- write a line of code to save a figure to .pdf

in rstudio

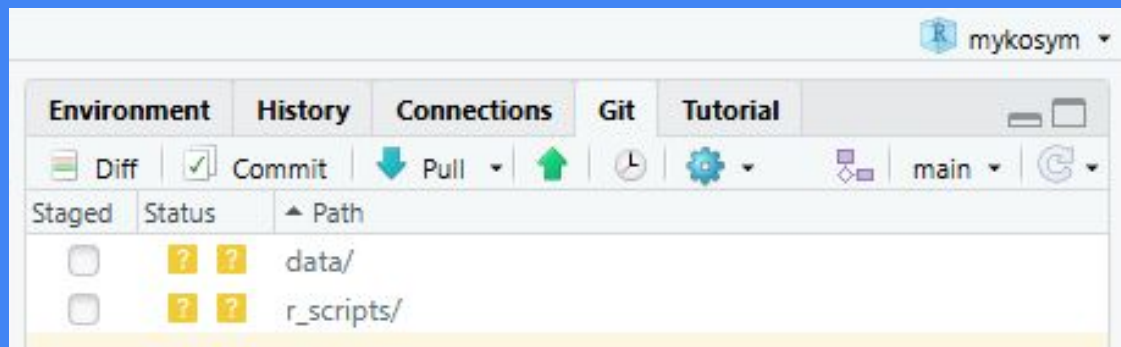
→ use the `here()` and `pdf()` functions to save a figure to your repo

```
pdf(here::here("path", "plot1.pdf"))  
plot(1,1)  
dev.off()
```

- stage, commit and push changes to remote repo

in rstudio

⇒ go to the git space (in the environment window)



⇒ stage all the changes you want to be seen in the remote repo (✓ tick the white boxes)

⇒ click on commit

⇒ write a commit message (describe the changes you made)

⇒ click on commit

⇒ click on push