Reproducible science workshop

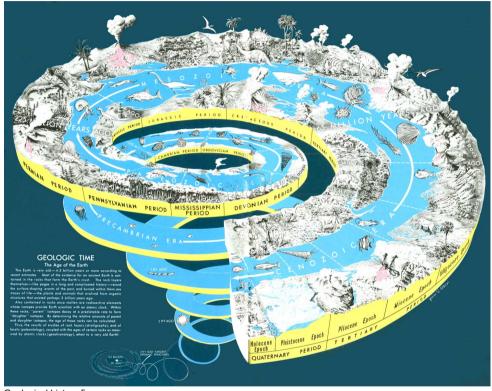
Jingxin Liu

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Reproducible science workshop

Jingxin Liu (https://www.researchgate.net/profile/Jing-Xin-Liu-2)

Play with embedding figures from my own computer



Geological history figure

Play with embedding figures from my own computer



Jinaxin

Play with embedding figures from internet and change the size of the figure



Jingxin

Play with embedding figures by knitr from my own computer



Geological history figure

The following is the tentative outline of a four day workshop that will be held in Xishuangbanna Tropical Botanical Garden (XTBG (http://english.xtbg.cas.cn/)) from October 19-22nd 2021. The purpose of this workshop is to get participants familiar with the concept of reproducible science that is surprisingly not commonly taught but which is becoming a requirement for better collaboration (not only with others but with oneself through time) and for credible science. At the end of this workshop, attendees should be familiar with tools use to achieve reproducible science, will be more confident with themselves, and be inclined to adopt these tools which will ease their collaboration with themselves as well as others. This course consists of four main parts which aim to putting everything from data to final report including r code and some interpretations and also exercises as step by step work me through method. This course will be mainly taught by Dossa G.G.O. click here to email me (mailto:%20dossa@xtbg.org.cn). Dossa has been teaching part of this workshop as a short module at the yearly postgraduate Advanced Fieldcourse in Ecology and Conservation ??? XTBG AFEC-X (https://www.pfs-tropasia.org/) for the past two years.

Who should attend this course?

Anyone who carries out research either as student, or research staff and seeks guidance to manage data efficiently, to improve peers credibility in their research output, and is keen to make science solid and credible. Consider thinking about the following questions:

- Have you ever worked collaboratively with others, and received multiple comments that you needed to deal with?
- Have you been ever asked by your collaborators/supervisor to go back to previous version of your manuscripts/analysis/ results?

- Have you ever thought that you finalized an analysis but to realize a mistake was in the data file?
- Have you ever spent endless time to explain your past / current projects to your collaborators (including yourself) but realize you even do not remember what you did at some point or why you did certain things?
- Have you ever copied a table from R/SPSS/Minitab/SAS to excel, and subsequently from Excel to your word
 processing document? And most annoyingly, you need to repeats these steps whenever anything has
 changed in the analysis or code.

If you answer **Yes** to any of the above questions then, this workshop is designed for you. Above all, there are five selfish reasons for why you would love to take this workshop click here *Five selfish reasons* (https://genomebiology.biomedcentral.com/articles/10.1186/s13059-015-0850-7)

hand make a table a table with your name, sex where you fill in your name your along the name(s) of your supervisor and their sex.

my first table made in Rmarkdown

Supervisor name	sex	Name	Sex
Jin Chen	Male	Jingxin Liu	Male
my first table made in Rmarkdown	Positi	on.	Sex
Jingxin Liu	Student		Male
Jin Chen	PI		Male

play with git bash

set up git etc in Rstudio

then open git bash, then input the codes below

git config –global user.name "xtbgtraining" git config –global user.email "liujingxin@msn.com (mailto:liujingxin@msn.com)" git config –global –list

then the result is like the figure below

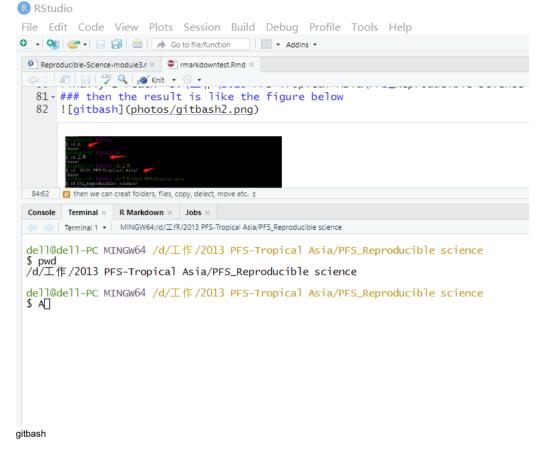
gitbash

then imput commonds to go to the directory that I want to work in cd d: # goes to drive D cd 工作 # goes into 工作 ... finally I reach "D:\2013 PFS-Tropical Asia_Reproducible science"

```
### then the result is like the figure below
   cd d:
 (base)
   cd 工作
 (base)
                   -PC MINGW64 /d/工作
   cd 2013\ PFS-Tropical\ Asia/
   cd PFS_Reproducible\ science/
           dell-PC MINGW64 /d/工作/2013 PFS-Tropical Asia/PFS_Reproducible science
 1 Workshop outline.html'
2 Markowetz 2015 Five selfish reasons to work reproducibly.pdf'
3 BES-Guide-Reproducible-Code-2019.pdf'
3 中国科学院西双版纳热带植物园会议审批表.docx'
                                                                                                                                          'R Markdown- The Definitive
'Reproducible science works
                                                                                                                                         'Reproducible science works
Reproducible-Science-modul
                                                                                                                                           Reproducible science works
  4 Reproducible Research in Computational Science-2011.pdf'
  fig5.jpg
How R Helps Airbnb Make the Most of Its Data-2018.pdf'
 pracl_Tidyverse Data manipulation.html'
pracl_Tidyverse Data manipulation.Rmd'
 pracl_Tidyverse Data manipulation_solutionfromDossa.Rmd'
 prac2_datavisualisationwithggplot2.Rmd
  prac2_datavisualisationwithggplot2_solution.html
 pracz_datavisualisationwithggplotz_solution.html
pracz_datavisualisationwithggplotz_solution.Rmd
R Graphics GGplot Cookbook.pdf'
R Markdown Cookbook.pdf'
R Markdown Cookbook-2021_Yihui Xie.pdf'
R Markdown- The Definitive Guide.pdf'
                                                                                                                                           rmarkdowntest.html
                                                                                                                                           rmarkdowntest.Rmd
 (base)
```

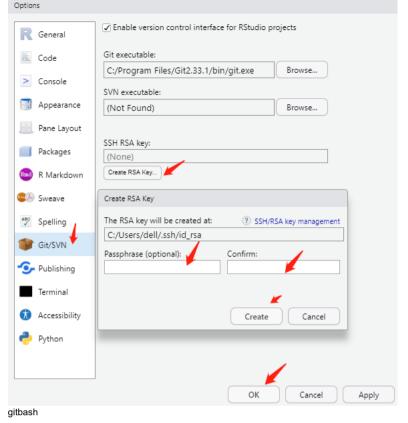
then we can creat folders, files, copy, delect, move etc.

actually as we seted up in rstudio that git bash can be opened in rstudio, see the figure



##How to creat ssh key, and use it to clone repositories from github account

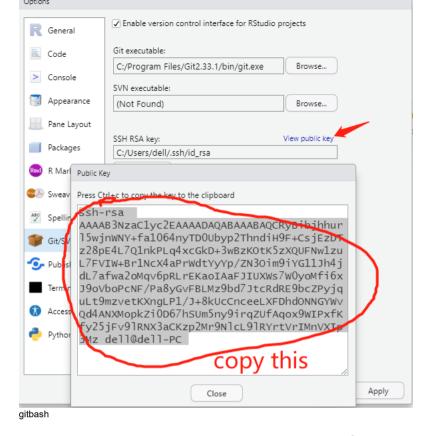
step 1 rstudio-tools-global option



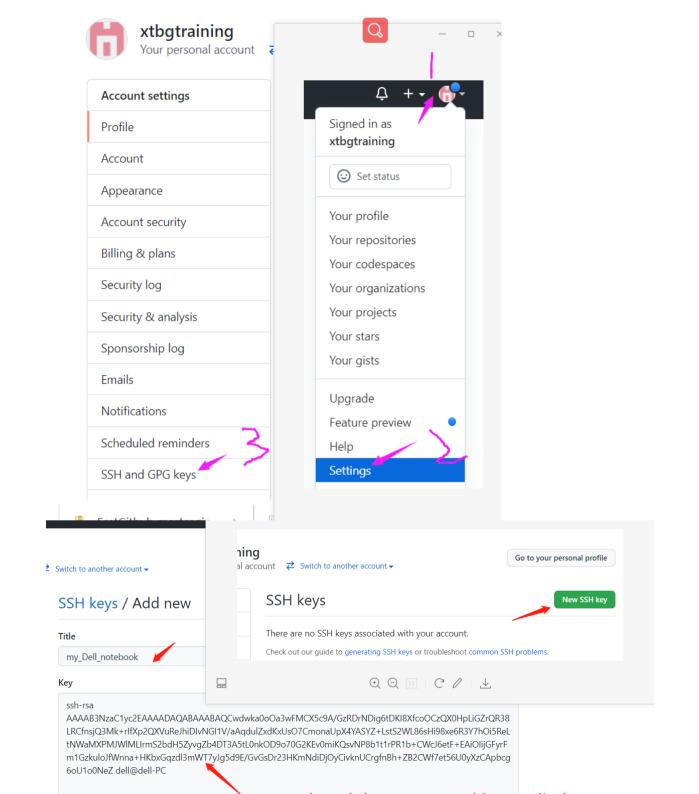
step 2, copy the ssh key in rstudio

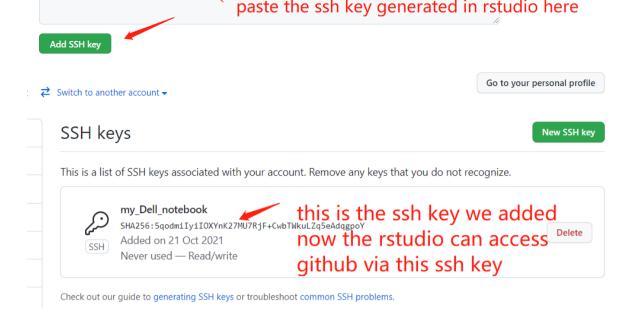
eeh_re

AAAAB3NzaC1yc2EAAAADAQABAAABAQCRyBibjhhurl5wjnWNY+fa1064nyTD0Ubyp2ThndiH9F+CsjEzDTz28pE4L7Q1nkPLq4xcGkD+3wBzKOtK5zXQUFNw1zuL7FVIW+Br1NcX4aPrWdtYyYp/ZN3Oim9iYGdell@dell-PC (mailto:dell@dell-PC)

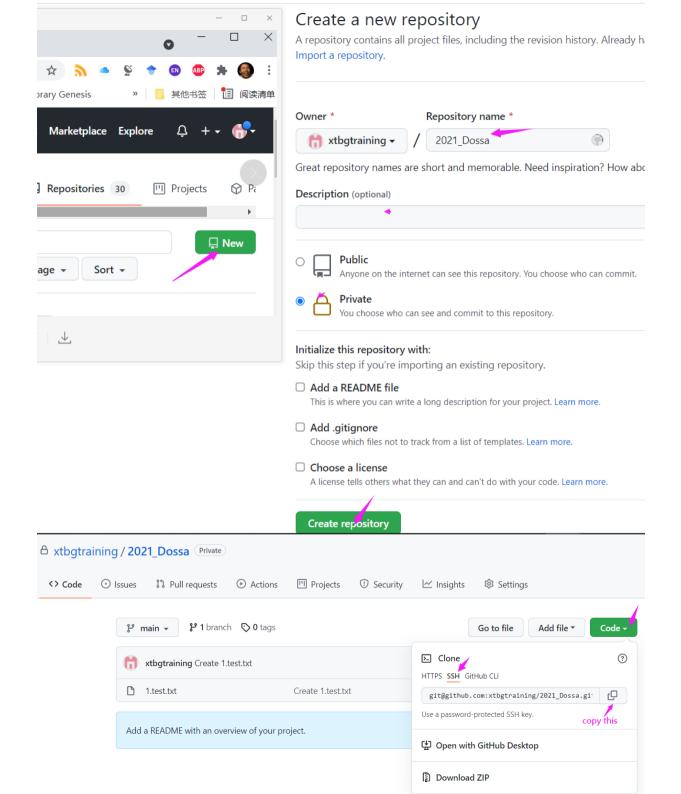


step 3 log into github and go to , there you will find a ssh key zone to creat ssh key past the copied ssh key there, this step links rstudio with your github account

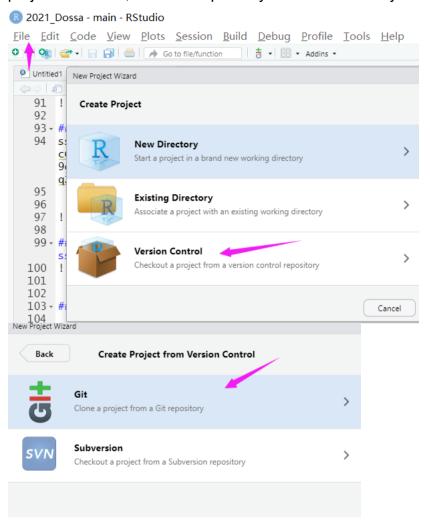


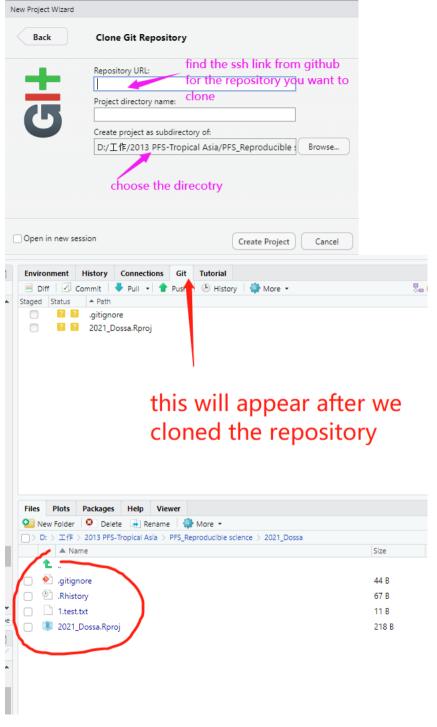


step 4 creat a new repository in your github account and get its ssh link



step 5 in rstudio, creat new project choose "version control", then "git", then copy the ssh link from the repository you want to clone from your repository, then press creat project in rstudio, then the repository will be cloned to your computer





software for using github in China

https://github.com/dotnetcore/FastGithub (https://github.com/dotnetcore/FastGithub)