# Marine Ecological Genetics exercises

### **General** info

Fridays 14:00 – 15:30 (Apr 22 – Jul 22, except May 27)

Contact: <u>martin.helmkampf@leibniz-zmt.de</u>

Not graded, attendance only

### Required software

- R
- RStudio
- git

Install R first from <a href="https://cran.r-project.org">https://cran.r-project.org</a> (for Windows, choose the base version)

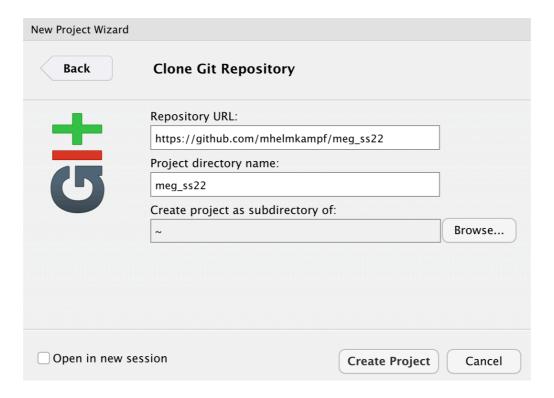
For RStudio, select and install the free Desktop version from <a href="https://www.rstudio.com/products/rstudio/download/">https://www.rstudio.com/products/rstudio/download/</a>

git comes already installed on computers running macOS. For Windows, install from <a href="https://git-scm.com/download">https://git-scm.com/download</a> (use default settings) and follow the instructions at <a href="https://www.computerhope.com/issues/ch001927.htm">https://www.computerhope.com/issues/ch001927.htm</a>

## **Course git repository**

https://github.com/mhelmkampf/meg\_ss22

Initial setup in RStudio: File > New Project > Version Control > Git



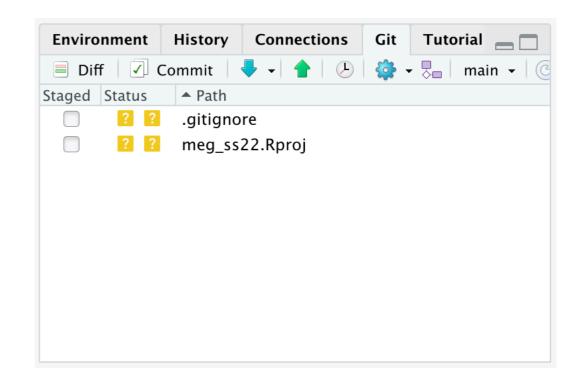
## **Course git repository**

Open existing project in RStudio:

File > Open File, select ~/meg\_ss22/meg\_ss22.Rproj

#### **Updating**

Go to "Git" tab, press blue arrow to "Pull"



Please save your own scripts (e.g. copies of the course scripts you would like to edit) to meg\_ss22/local and leave original scripts in meg\_ss22/code unchanged

## **Course outline**

Class	Date	Topics	Script
01	Apr 22	Introduction to R and RStudio	01_intro.R
02	Apr 29	Within population genetics (diversity, Hardy-Weinberg, inbreeding)	02_within.R