# preamble, Methods 3, 2025

#### 2025-09-11

REMEMBER: This preamble is **NOT** part of your portfolio, but is a prerequisite for doing the portfolio

# Preamble - GitHub, Python, Conda

The goals of the preamble are:

- 1) create a *Conda* environment that contains the *Python* packages that we need. Note that we are not creating an R environment I expect you to maintain your own
- 2) install your R-packages
- 3) connect your GitHub profile to the GitHub classroom such that you can hand in assignments and access course materials

# 1) What is Conda?

Conda is a package management and environment system.

## What is a package?

(from ChatGPT)

- Python: Packages are collections of Python modules. They are often distributed as .whl or .tar.gz files. Examples include numpy, pandas, and requests.
- R: Packages are collections of R functions, data, and compiled code. They are distributed as .tar.gz files. Examples include *qqplot2*, *dplyr*, and *shiny*.

# Why environments?

We use environments to create **isolated**, **shareable** and **conflict-free spaces** for our projects, e.g. the *methods3\_2025* that we are going to create now won't interfere with other projects that we may be running

#### Installling Conda: miniconda vs. anaconda

Section can be skipped if you already have miniconda installed

I recommend installing *Conda* using the **miniconda** distribution. The **anaconda** distribution comes with many pre-installed packages, which may lead to conflicting packages when creating environments.

Command line install (preferred method)

**Link**: https://docs.anaconda.com/miniconda/#quick-command-line-install

(If you already have anaconda installed, you may prefer keeping it to not create any conflicts )

## Create method3\_2025 environment

```
cd <path_that_contains methods_environment.yml> # go to folder with methods3_environment.yml file conda env create -f methods3_environment.yml # create environment conda activate methods3_2024 # activating your newly created environment
```

## First goal achieved

You are now in an environment that is **isolated** from all *Python* that you may have installed at earlier date.

The **shareable** methods3\_environment.yml file contains the recipe for the environment and Conda makes sure that the installation is **conflict-free** in terms of dependencies.

#### Installed packages:

```
cat methods3_environment.yml
```

name: methods3\_2025
channels:
 - defaults
 - conda-forge
dependencies:
## Python version
 - python>=3.10

#### ## Python packages

- pip
- scikit-learn
- matplotlib
- numpy
- scipy
- pandas
- seaborn

# 2) R packages to install

We are going to be dependent on the packages lme4 and fields. Please install as you usually do

# 3) Connect your \_GitHub account to the \_GitHub Classroom:

The GitHub Classroom is where assignments will be shared and where answers to them can be uploaded (If you don't have a GitHub account sign up at www.github.com)

## Accepting an assignment

Click the shared assignment link: https://classroom.github.com/a/ELwibfE5

The assignments will be handed in the by the study groups that you have been assigned:

**IMPORTANT**: Create your study group using the appropriate name below: (I do know some of you go by other names, but using the names you are registered by makes it easier to cross-check with the official rosters)

#### Team names

- DávidViktorChristianIda
- AmalieMaikenMikkelCarina
- ArinaNajaJohanaTeréziaHannah
- AsgerDominikJesperKamila
- $\bullet \quad Aya Signe Jens Katrine Nanna$

- CasperSørenMadalinaAsgerSofie
- CiljaSarahLineaMelanie
- DomonkosYoavMadsDóra
- EmaVictoriaAlessandraMartinJosefine
- EmmaMariusMilleAndrea
- GreteJakobNeleHelenKamila
- AnnePaulineAlexAsta-MarieSophia
- JuleJuliaAurelijaSilas
- KatarinaNoraCamillaSørenAdriána
- KatrineVicthoriaEmilVivi
- MagnusNannaEmiliaWilliam
- NoemiRitaWilliamSara
- PetraSineHansAgnes

Methods-3-2025

# Accept the group assignment — Preamble\_test

Before you can accept this assignment, you must create or join a team. Be sure to select the correct team as you won't be able to change this later.

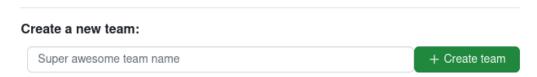


Figure 1: Enter team name

## Doing the assignment

When you have accepted the assignment, you will see the .Rmd or .ipynb file with the assignment:

You can then clone the repository, by clicking the green code button and copying the URL

```
git clone <URL to repository>
```

Then provide the solution in the Rmd and knit a pdf

Then add, commit and push to your repository (from your cloned folder)

```
git add assignment_0_test.Rmd
git add assignment_0_test.html
git commit -m "solution <group_name>"
git push
```

Now your assignment repository contains the solution, where I (Lau) can access them

The URL's to the six assignments that go into the portfolio will be shared through Brightspace

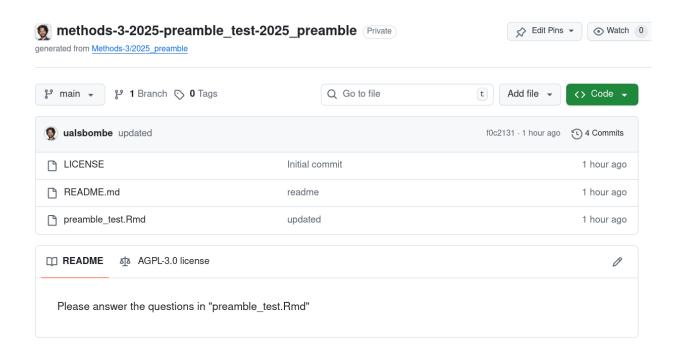


Figure 2: The assignment repository

```
title: "preamble_test"

date: "2024-09-11"

output: html_document

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# Exercises and objectives

The objectives of today's exercises are:\
1) Check that your environment works by running single level ...
2) ... and multilevel models in R
3) Run a single level model in Python
```

Figure 3: The assignment

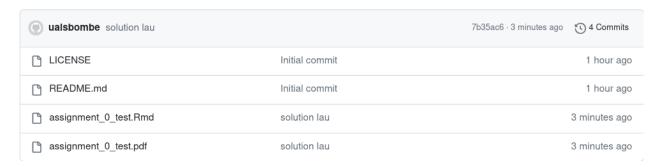


Figure 4: repository with solution

# Conclusion

Third goal achieved: you can now access the portfolio assignments as they are uploaded, and you will be able to upload your answers to your group repositories

This concludes the preamble test