



#### The HUB as a cloud based service

Riccardo Soldan March 2023



### Content

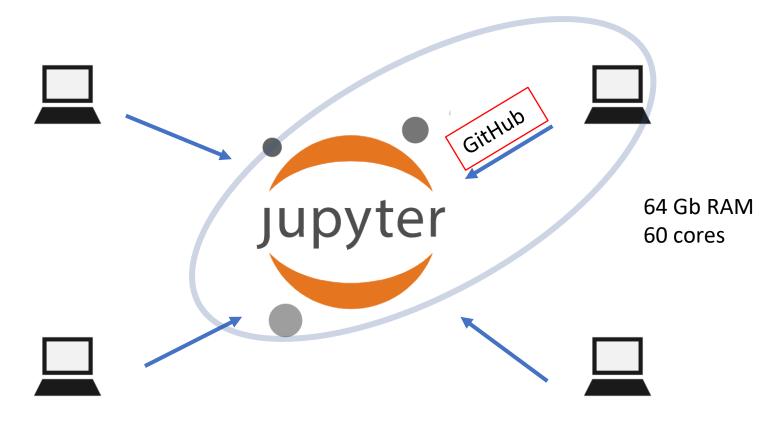
- 1. Introduction to the HUB
- 2. What can be done and what cannot be done

All the necessary information and material used in this training is available in our GitHub repository



### The HUB

The HUB is a cloud based environment, powered by University of Cantabria, that allow <u>registered users</u> to access a JupyterLab. Here, users can work with R or python, install libraries, perform computational work







#### How to use the HUB

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### Content

- 1. Login
- 2. Customize your environment (notebooks, folders, etc)
- 3. Open a notebook
- 4. Installing packages
- 5. Code of conduct



## Login

- Log-in to the HUB (You need to login to GitHub first). It is as simple as clicking the link
- You will be able to access the HUB if you were registered among allowed users
- If you are not, fear not. This workshop is useful regardless, so stay focus

Sign in with GitHub



# Customize your environment

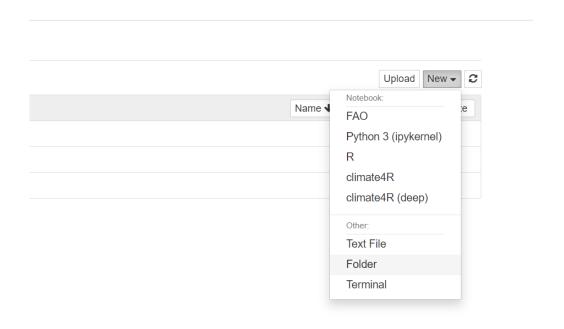
• Once you log in, you would see the page below. You are now using a virtual machine

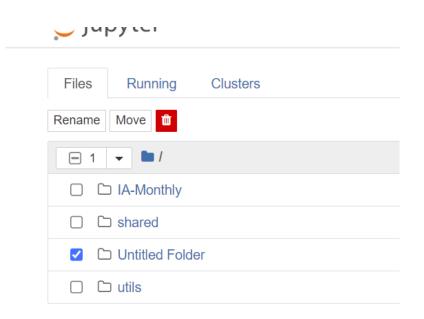




## Customize your environment

You can now customize your environment (e.g create a folder with your name)







## Customize your environment

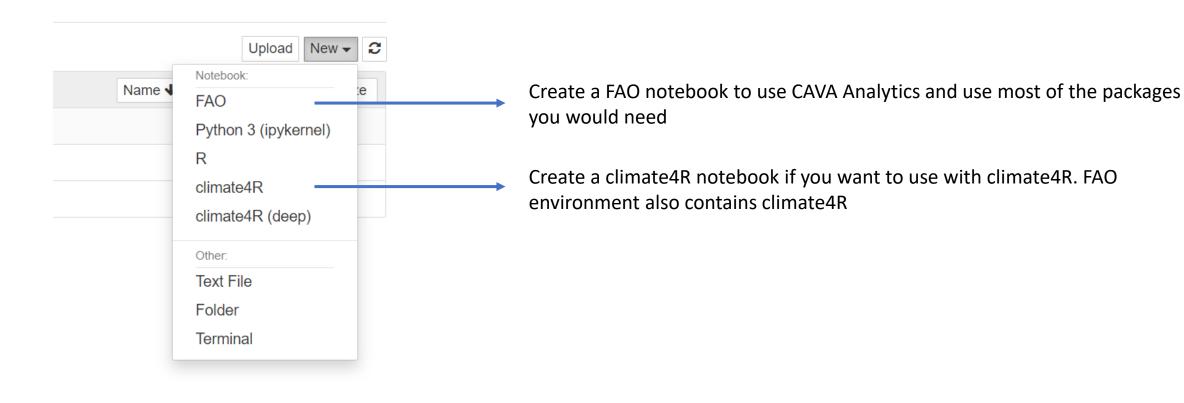
• You can also upload some data (e.g an excel file) (space is limited, do not upload Gbs of data)



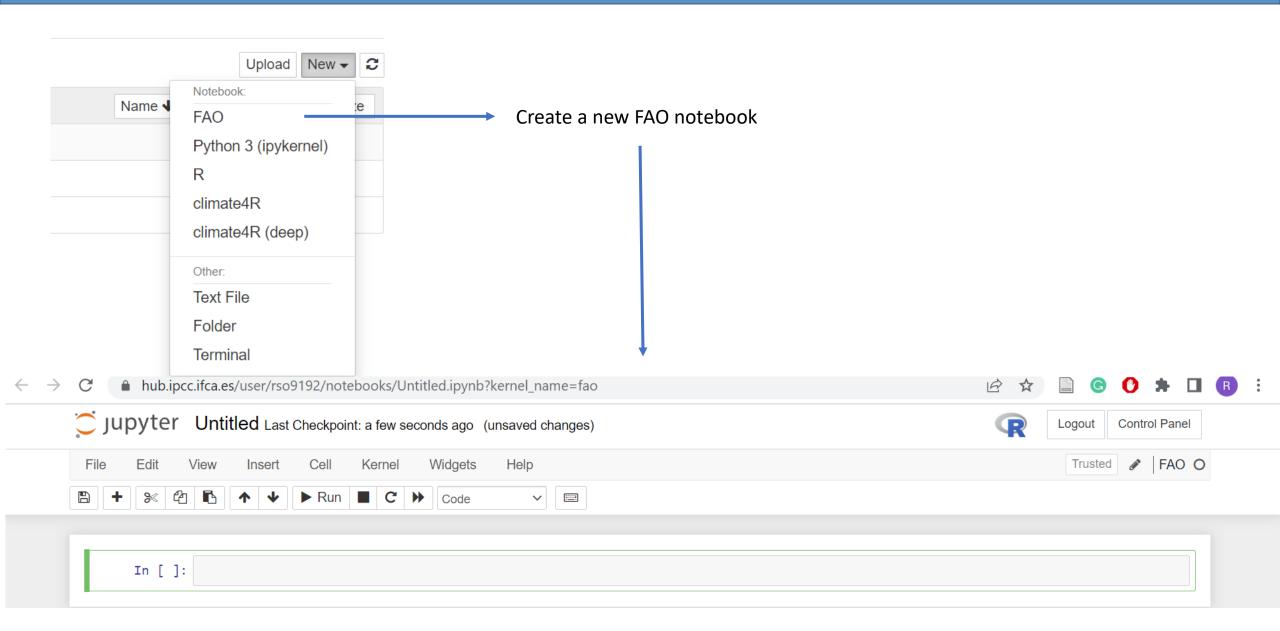


### Understanding environments

You can choose the environment that you want to use. Environments are conda environments that contains
 R installation and a pre-set of installed packages. Read why it is useful to work with conda environments <a href="here">here</a>



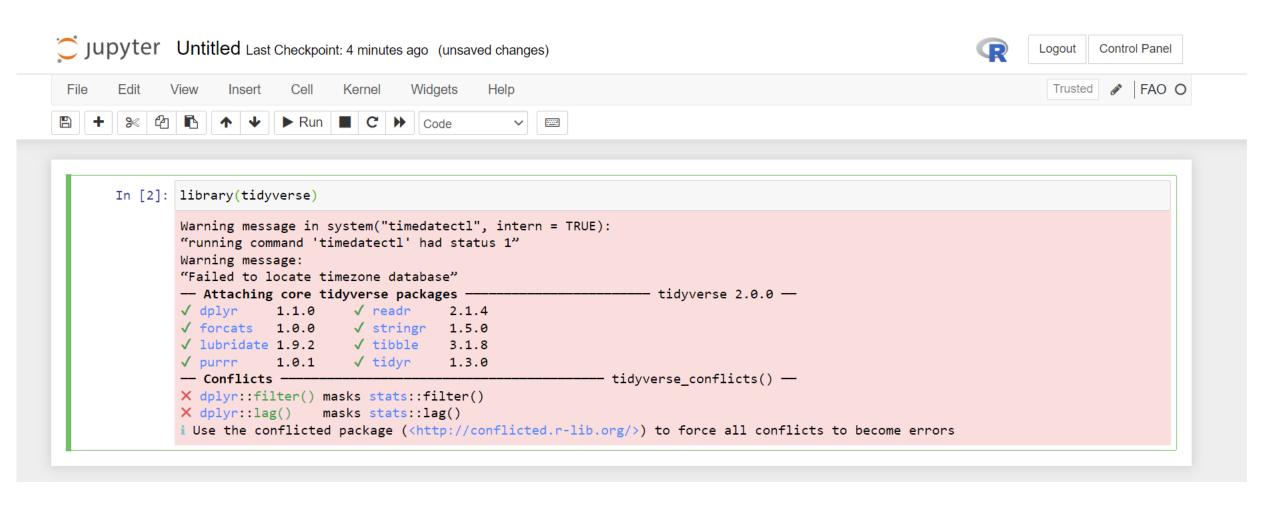
### Opening a FAO notebook





### Opening a FAO notebook

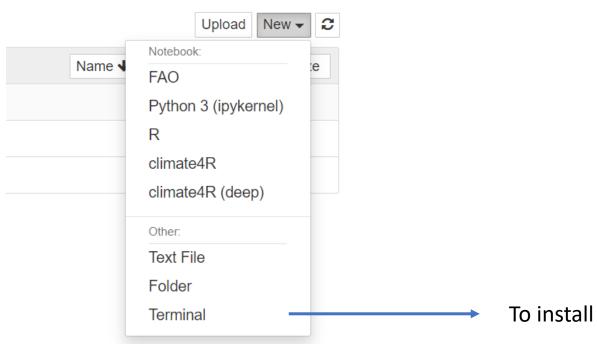
Now you can work as you would normally do in Rstudio





### Installing packages

• Let's imagine you need a library that it is not installed in FAO or climate4R environments. To install a new package, you need to open a **terminal** 



To install a new package open a terminal



### Installing packages

You will now see this

```
jovyan@630fd26c9279:~$
```

- Run conda init bash, close the terminal and open it again
- Check the available envs with conda info --envs

### Installing packages

Now activate the environment in which you want to install a package, for example conda activate fao

```
(base) jovyan@630fd26c9279:~$ conda activate fao
(fao) jovyan@630fd26c9279:~$
```

Open R

```
(base) jovyan@630fd26c9279:~$ conda activate fao

(fao) jovyan@630fd26c9279:~$ R

R version 4.2.2 (2022-10-31) -- "Innocent and Trusting"

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Platform: x86_64-conda-linux-gnu (64-bit)

R is free software and comes with ABSOLUTELY NO WARRANTY.

You are welcome to redistribute it under certain conditions.

Type 'license()' or 'licence()' for distribution details.
```

• Install the required package with install.packages("ggplot2")



#### Code of conduct

• In the tab running, you will see what is currently running. Remember to click on shutdown when done with your work. If you don't, you will be occupying memory





### Code of conduct

- Be mindful on the usage of computational resources as these are shared among several users
- Do not upload excessive amount of data (several Gbs) from your local computer
- Always close your notebooks when done with your analyses



# Exercise (10 minutes)

- Create a new folder with your name
- Open a FAO notebook and rename it
- Close the notebook

## Thank you!

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