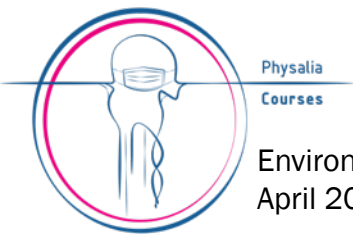


# Environmental metagenomics

Course outline and practical info



Physalia  
Courses

Environmental metagenomics  
April 2021

Igor S. Pessi & Antti Karkman, University of Helsinki

# About us

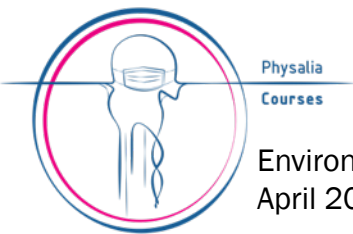
## Organizer:

- Carlo Pecoraro, Physalia Courses
  - [info@physalia-courses.org](mailto:info@physalia-courses.org)



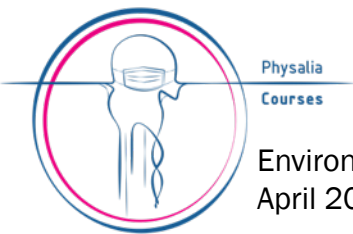
## Instructors:

- Antti Karkman, University of Helsinki
  - [antti.karkman@helsinki.fi](mailto:antti.karkman@helsinki.fi)
- Igor Pessi, University of Helsinki
  - [igor.pessi@helsinki.fi](mailto:igor.pessi@helsinki.fi)



# About you

- Name
- University/Institute/Company
- Research interest(s)
- Previous experience(s) with microbial ecology, metagenomics, bioinformatics, etc.
- General hopes for this course



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# Course outline

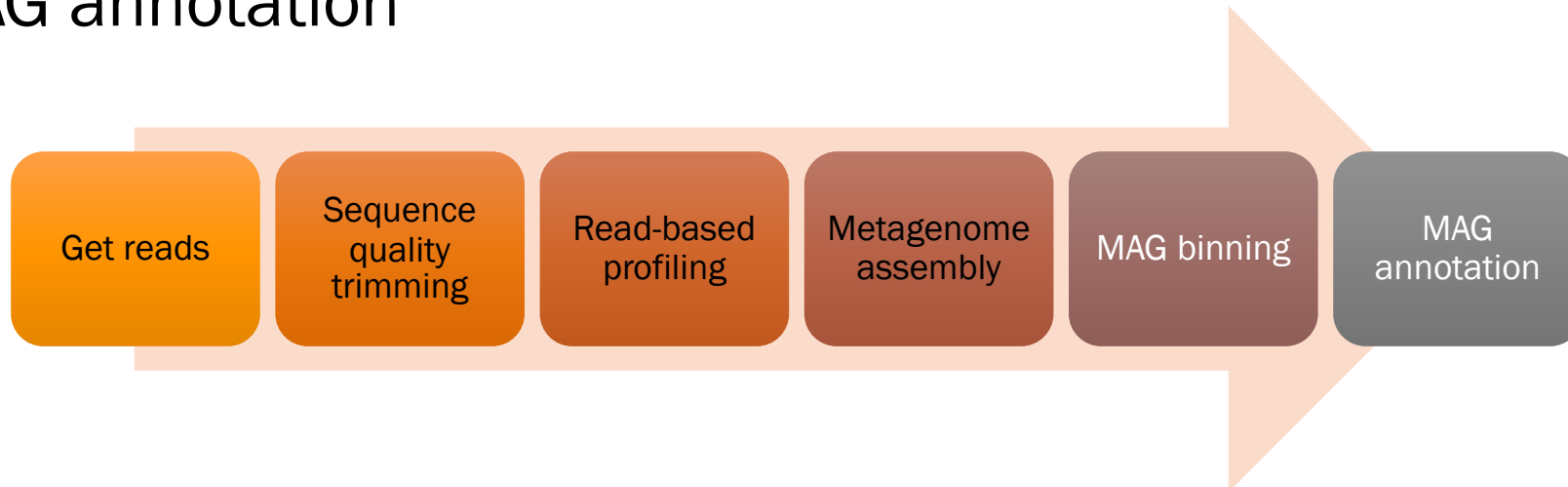
**Day 1:** Working with the command line and QC

**Day 2:** Read-based analyses

**Day 3:** Metagenome assembly

**Day 4:** MAG binning

**Day 5:** MAG annotation



# Practical information: Zoom & GitHub

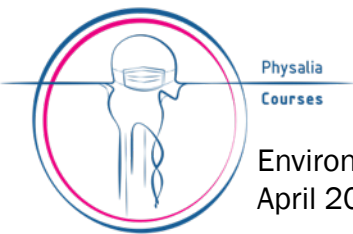
The course will take place in Zoom from 9 AM to 4 PM (CET)

- Link to the Zoom room in Slack ([#general](#))

The course page containing exercises and presentations is:

- [https://github.com/karkman/physalia\\_metagenomics](https://github.com/karkman/physalia_metagenomics)

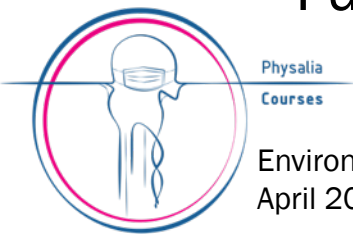
**Please bookmark this address!**



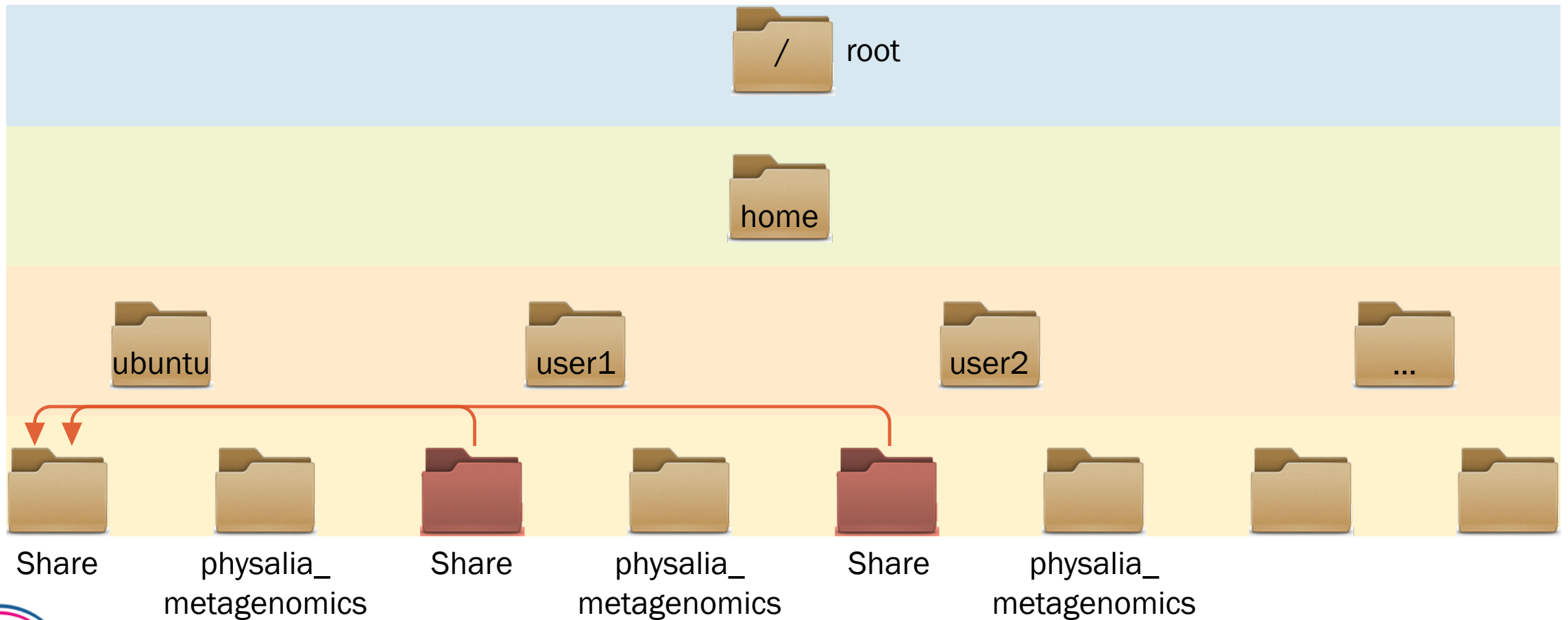
# Practical information: Amazon Cloud (AWS EC2)



- See Slack ([#before-start](#)) for information on how to connect
  - Remember, the IP address will change every day
- Everyone has a user, a home and a shared folder
  - E.g. Lucie Malard is [user2](#)
  - Their home folder is [/home/user2](#)
  - Their shared folder is [/home/user2/Share](#)
  - List of usernames can be found in Slack ([#before-start](#)).
- We will mostly use conda for managing the software environments
  - The environments are already set up for everyone
  - Further instructions on the GitHub page



# Practical information: folders and files

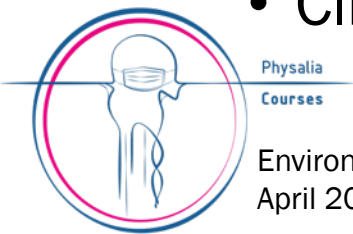


# Practical information: FileZilla



## Set-up instructions:

- Open FileZilla
- Click File > Site Manager
- Click New site
- Change Protocol to SFTP– SSH File Transfer Protocol
- In Host, type the IP address (has to be changed every day)
- In Port, type 22
- Change Logon type to Key file
- In User, type your username
- In Key file, select your pem file
- Click connect





# Kilpisjärvi, Finnish Arctic (69° N)



# Kilpisjärvi, Finnish Arctic (69° N)

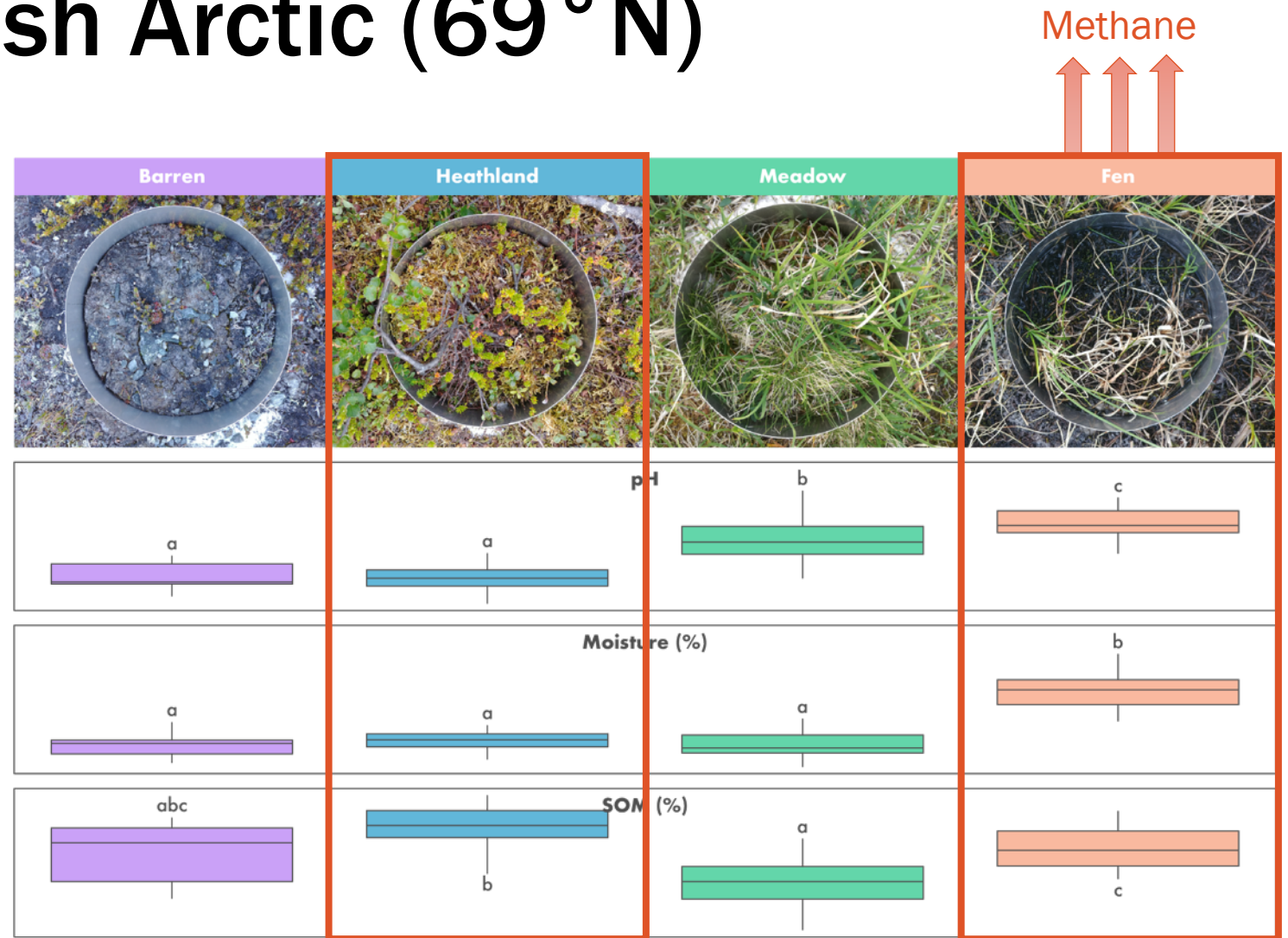
## 4 samples

- Sample01/Sample04: heathland soils
- Sample02/Sample03: fen soils

## Sequencing:

- Illumina NovaSeq: all samples
- Nanopore MinION: Sample03 and Sample04

Real (not toy) data!



Pessi et al., 2020: <https://www.biorxiv.org/content/10.1101/2020.12.21.419267v1>