

# Session de formation 2023





















bioinformatics platform dedicated to the genetics and genomics of tropical and Mediterranean plants and their pathogens





comparative genomics transcriptome assembly SNP detection phylogeny structural variation differential expression GWASpangenomics

population genetics polyploidy

Mutualisation



Cacao Banana Coffee Rice Palm Cassava Pseudocercospora Magnaporthe









4 institutes





3 research units





Storage and computing resources

Tools



Trainings



### Storage and computing resources





Meso@LR au CINES

1090 threads:

35 standard nodes

2 bigmem nodes

1 GPU node

500 To of replicated storage





CINES

1130 threads:

30 standard node

1 supermem node

1 GPU node

150 To on 3 NAS + 210 To scratch





Resources mutualised at Meso@LR through the Mudis4Ls project (purchase/storage/data)





## Collaborative development of tools

Pangenomic

Gene families

Comparative

Genomics

Phylogeny

**Assemblies** 

Annotation

Data mining

Diversity exploration

genotype manipulation

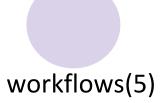
mosaic manipulation

web applications (16)



tools

visualisation (8)



packages (4)

Metagenomic



https://github.com/SouthGreenPlatform/

## I-Trop

Plant & Health Bioinformatics Platform

\*\*Select= 1
\*\*Select=1
\*\*Select=1
\*\*Scene.objects.
\*\*Scen



#### https://bioinfo.ird.fr/















AURORE COMTE



JACQUES DAINAT



ALEXIS DEREEPER



BRUNO GRANOUILLAC



JULIE ORJUELA-



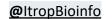
NDOMASSI TANDO



CHRISTINE

bioinfo@ird.fr









Florian Charriat Antoni Exbrayat



Guilhem Sempere







Bruno Granouillac Jacques Dainat

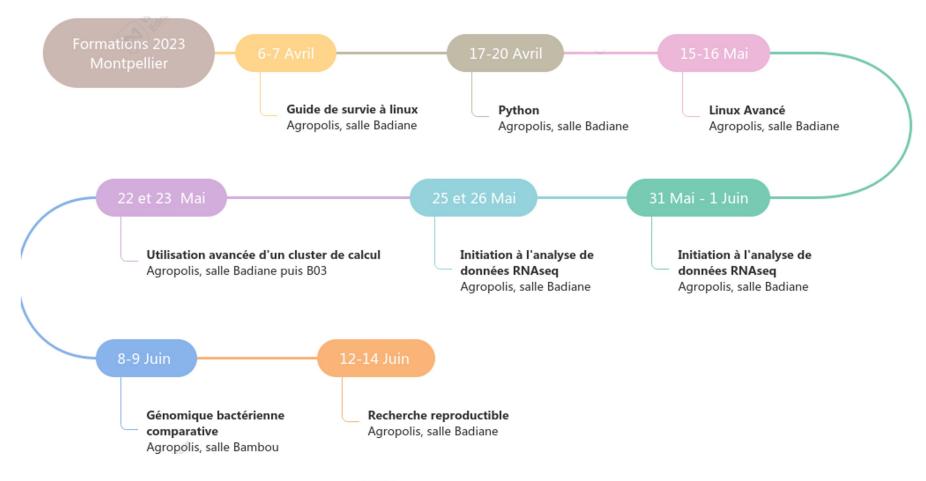
Nicolas Fernandez



**Thomas Denecker** 

And more collaborators!

























# **Formation Python**



Adapté du cours HLIN404 de Anne-Muriel Chiffoleau (UM)









