

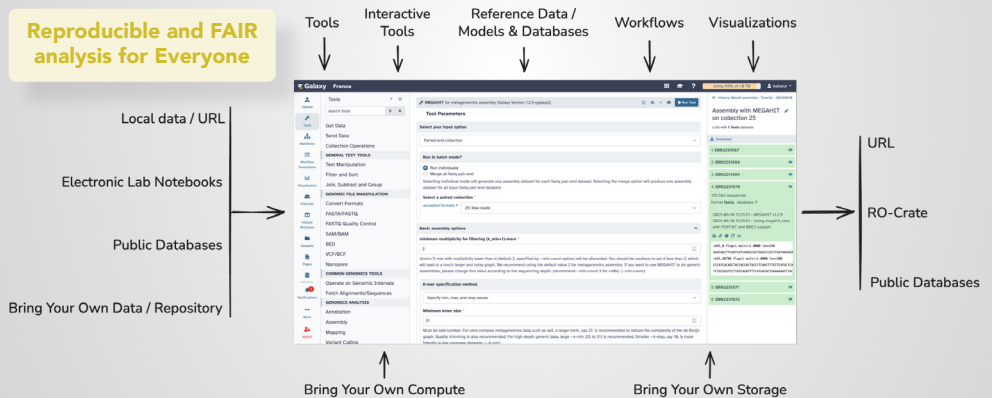
Galaxy

Galaxy is an open-source platform for FAIR data analysis.

It enables users to:

- use tools from various domains (that can be plugged into workflows) through its graphical web interface.
- run code in interactive environments (RStudio, Jupyter...) along with other tools or workflows.
- manage data by sharing and publishing results, workflows, and visualizations.
- ensure reproducibility by capturing the necessary information to repeat and understand data analyses.

The Galaxy Community is actively involved in helping the ecosystem improve and sharing scientific discoveries.



Galaxy Training: A powerful framework for teaching!

400+ tutorials

25+ learning pathways

150+ hours of recording

35+ topics including

proteomics, metabolomics, climate, NLP, etc

Training Infrastructure as a Service



Many major project use Galaxy

 **ABRomics**
ANTIBIORESISTANCE

 **BRC Analytics**

 **EARTH BIOGENOME PROJECT**

 **eosc**



Objectives:

1. **Meeting** every 2 months
2. **Connect** galaxy users, developers, admins across France
3. **Share** knowledge (ELIXIR, Galaxy Community, etc)
4. **Support** [UseGalaxy.fr](https://usegalaxy.fr)
5. **Combat** misconceptions about Galaxy within the French scientific community

Mailing list: bit.ly/galaxy-gt



useGalaxy.fr

Communities - Galaxy subdomains :

Microbiology, Single-cell, Workflow4Metabolomics,
Plants-Pathogens, etc.

ELIXIR



Launch: 2019



10.4k Users



6.2M Jobs



1.7k+ Tools
sorted by
thematic



25+
Interactive
Tools

TiaaS: Since February, 2022



110+ Events



2.9k+ taught
Students

On the last year



3k Active
Users



2.5M+ Jobs



4.7k CPU
Cores



61 TB
of RAM



11 Virtual
GPU Cards



Support online by
the community

Update June 2023



UseGalaxy.fr,
a service hosted on
the IFB-core cluster



INSTITUT FRANÇAIS DE BIOINFORMATIQUE

