Microbial data analysis in Galaxy

Whether you are analysing microbiome samples or bacterial isolates, long reads or short, shotgun or 16S, genomics transcriptomics or proteomics, this is the place to be!

microGalaxy is a community of practice in the Galaxy community for anything regarding microbial data analysis in Galaxy!

What is available?

Tools

Many tools for microbial data analysis are freely available in the ToolShed which can be installed on any Galaxy server. To keep an overview, the microGalaxy community maintains a <u>curated list</u> of the **150+ tools** available.

If tools are missing or information is not up-to-date in the list, please help us! Contact Saskia or Bérénice about it.

Workflows and tutorials

Several curated Galaxy workflows are publicly available for different kinds of microbial data analysis. Many of these are accompanied by comprehensive <u>GTN Tutorials</u> that will guide you through the analysis step by step.

Microbiome

Target	Technique	Sequencing	Analysis	GTN Tutorial	Workflow	Notes
Bacteria	16S	Short reads	Taxonomic Profiling	<u>View</u> <u>Tutorial</u>	<u>View</u> <u>Workflow</u>	mothur SOP
Bacteria	16S	Long reads	Taxonomic Profiling	<u>View</u> <u>Tutorial</u>	<u>View</u> <u>Workflow</u>	
All	Shotgun metagenomics	Short reads	Taxonomic Profiling, Functional Analysis		<u>View</u> <u>Workflows</u>	ASaiM (missing referenc e)
All	Shotgun metagenomics	Long reads	Taxonomic profiling	View Tutorial		

Plasmid	Shotgun	Long reads	AMR detection	<u>View</u> <u>Tutorial</u>		
All	Shotgun metatranscript omics	Short reads	Functional Analysis	<u>View</u> <u>Tutorial</u>	<u>View</u> <u>Workflow</u>	ASaiM- MT (missing referenc e)

Bacterial Isolates

Target	Sequencing	Analysis	GTN Tutorial	Workflow	Notes
M. tuberculosis	Short reads	Variant Detection	<u>View</u> <u>Tutorial</u>	<u>View</u> <u>Workflow</u>	
M. tuberculosis	Short reads	Transmission of Mtb	<u>View</u> <u>tutorial</u>		
M. tuberculosis	Short reads	Phylogenetics applied to Mtb evolution & epidemiology	<u>View</u> tutorial		
Plasmids	Long reads	AMR detection		View Workflows	

Want to include your workflow here? Let us know!

A dedicated interface

Training material and events

How is Galaxy used for microbial data analysis?

Projects / Showcases

Below is a list of projects involving members of this community:

Project (name, link)	Descriptio n	Techniqu es	Seque ncing	Analyses	People involve d	Fundin g	Status
IRIDA (Website)	The Integrated Rapid Infectious Disease Analysis				Aaron Petkau		
Foodborn e pathogen detection		Metageno mics	Long reads		Bérénice Batut, Engy Nasr	EOSC- Life	Ongoin g
BeerDEco ded,		Metageno mics	Long reads		Bérénice Batut, Street Science Commu nity	Mozilla, de.NBI,	Ongoin g
DNAnalyz er		Metageno mics	Long reads		Bérénice Batut, Street Science Commu nity	Universi ty of Freibur g	Ongoin g
Cloud data		Metageno mics, metatrans criptomics	Short reads	Taxonomy profiling, Functiona I profiling, Assembly, MAGs builiding	Bérénice Batut, Engy Nasr		Ongoin g
ASaiM		Metageno mics, metatrans criptomics	Short reads	Taxonomy profiling, Functiona I profiling	Bérénice Batut, Saskia Hilteman , Pratik Jagtap, Subina Mehta, etc		Finishe d?

Mycobacter ium tuberculosi s NGS made easy: data	Series of modules containing recorded videos by experts and tutorials		Illumin a (wanti ng to expan d to nanop ore)	Mapping & Variant calling Molecular epidemiol ogy of TB using NGS Phylogen etics for studying TB evolution and epidemiol ogy	Daniela Brites; Peter van Heusde n; Galo A. Goig Christop h Stritt	Initially funded trou gh TB projects of capacity building; currentl y no funding	Ready to be used but also Working on improve ments
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Papers

Reference	Short summary, including how Galaxy was used

Partners

Partners (name, link)	Description	People involved
Seq4AMR Seq4AMR (Website)	JPIAMR Network for Integrating Microbial Sequencing and Platforms for Antimicrobial Resistance	Saskia Hilteman
ELIXIR Emerging Microbiome Community		Bérénice Batut
µbioinfo Slack		

microGalaxy Community

microGalaxy is a community of practice to

- **Develop** and **Sustain** microbial data analysis in Galaxy
- Implement standardised "best practices"
- Expand documentation and training
- Coordinate efforts to avoid redundancy in tools, workflows and training
- Share the amazing things we are all doing together and outside Galaxy
- Support each other

Meetings

The microGalaxy community usually meets **4 times per year** (including 1 BoF at GCC and other community events) for a **2-hour meeting**. The meeting **time alternates** between 5-7pm Central European time and 9-11am Central European time in order to support members leaving in different timezones. The main aims of the meetings are to

- · Get news and updates from the community
- Have 1 or more "short" talks about a project or a topic of interest
- Discuss and coordinate our efforts.

Details about the next events and minutes of previous meetings can be found here.

Join us

Anybody interested in microbial data analysis in the Galaxy is welcome to join microGalaxy! **Everybody is Welcome!**

- Join our quarterly meetings!
- Join the Gitter Chat (also available via Matrix)
- Join the mailing list microgalaxy@lists.galaxyproject.org

Resources

Resource	Description	Links
jpiamr	JPIAMR: Global Coordination of Antimicrobial Resistance Research	Website

