

Core Facility Computational Bioanalytics

Introduction into Galaxy for Life Science Researchers

Galaxy is an open source, web-based platform for data intensive biomedical research. This workshop will teach participants the basics of Galaxy, introducing the Galaxy interface including a working example (Preprocessing and QC, Microbiome, ChIP-Seq or RNA-Seq analysis) .

The course participants will take the first steps with Galaxy. Upon successful completion of this course, they will be familiar with:

- Galaxy interface,
- Working with histories
- Data upload, data analysis and visualization
- Share data with others

Teaching and learning method: Interactive hands-on training

Languages of instruction: English

Target audience: PhD students, technicians and researchers

Entrance qualifications: No

Hardware and Software: Please take a laptop with you. You can log into Galaxy with your MUG-account.

Costs: for free

Registration: zmf-sekretariat@medunigraz.at (deadline: September, 18th 2019 or November 27th 2019)

As the number of participants is limited (min. 3 and max. 10), please register early to confirm your seat!

October 2nd 2019 (2 pm - 4 pm)
December 11th 2019 (2 pm - 4 pm)
ZMF seminar room ground floor EG-086

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