



**GENOMICS
RESEARCH
CENTER**



Introduction to Galaxy

July 19-20, 2016

9:00 AM – 5:00 PM

Nesbit Computer Classroom (G-9576),
University of Rochester Medical Center

Instructor: Dave Clements, M.S., Galaxy Training Network

What is Galaxy?

- Web-based platform for data intensive life science research
- Enables non-bioinformaticians to create, run, tune, and share their own bioinformatic analyses
- Limited programming or Linux command line experience is required.

What will the workshop cover?

- How to integrate data and perform simple and complex analysis
- Data visualization and visual analytics
- How to share and reuse your bioinformatic analyses
- All from within Galaxy
- See the [Agenda](#) for more detailed information

Attendance is capped at 25 participants, so please [register](#) to reserve your space.

Contact miner_bioinformatics@urmc.rochester.edu with questions.

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