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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