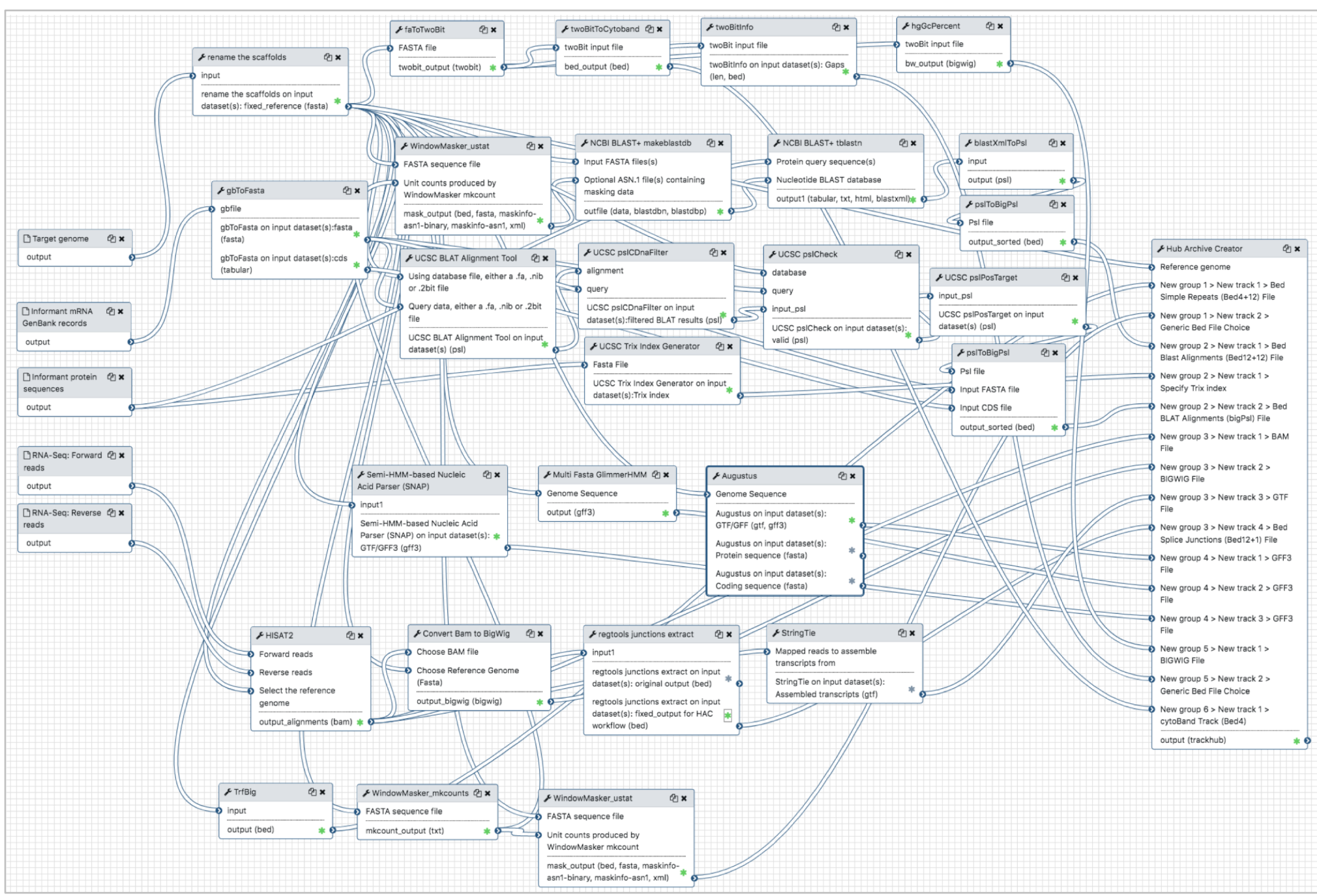


Galaxy: A Global Platform & Community for Life Science Data Integration & Analysis

Dave Clements¹ and the Galaxy community²

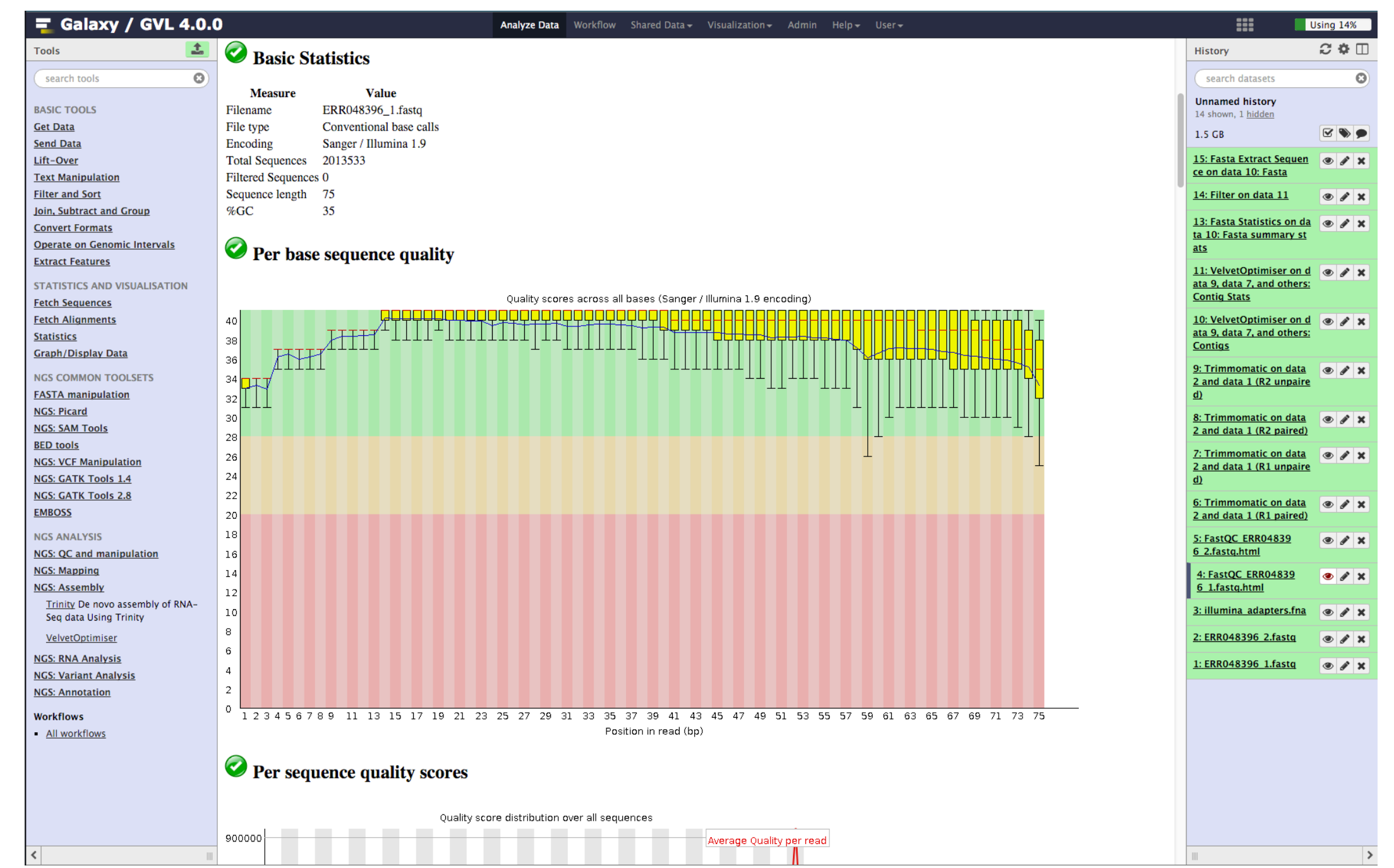
1. Galaxy Project, Johns Hopkins University; 2. galaxyproject.org



G-OnRamp, Wilson Leung



GCC2019, Bérénice Batut



Microbial de novo Assembly for Illumina Data, Melbourne Bioinformatics

Use Researchers

On the web, right now



120+ Public Servers



On cloud services



galaxyproject.org/use

Docker & Virtual Machines



Over 35 Docker and VM images with fully functional pre-populated Galaxy instances on them, that you can setup and run on your laptop in minutes

Visualization, Epigenomics, metagenomics, RNA-Seq, BLAST+, ChIP-Seq, Exome-Seq, Image Analysis, Mass Spec ...

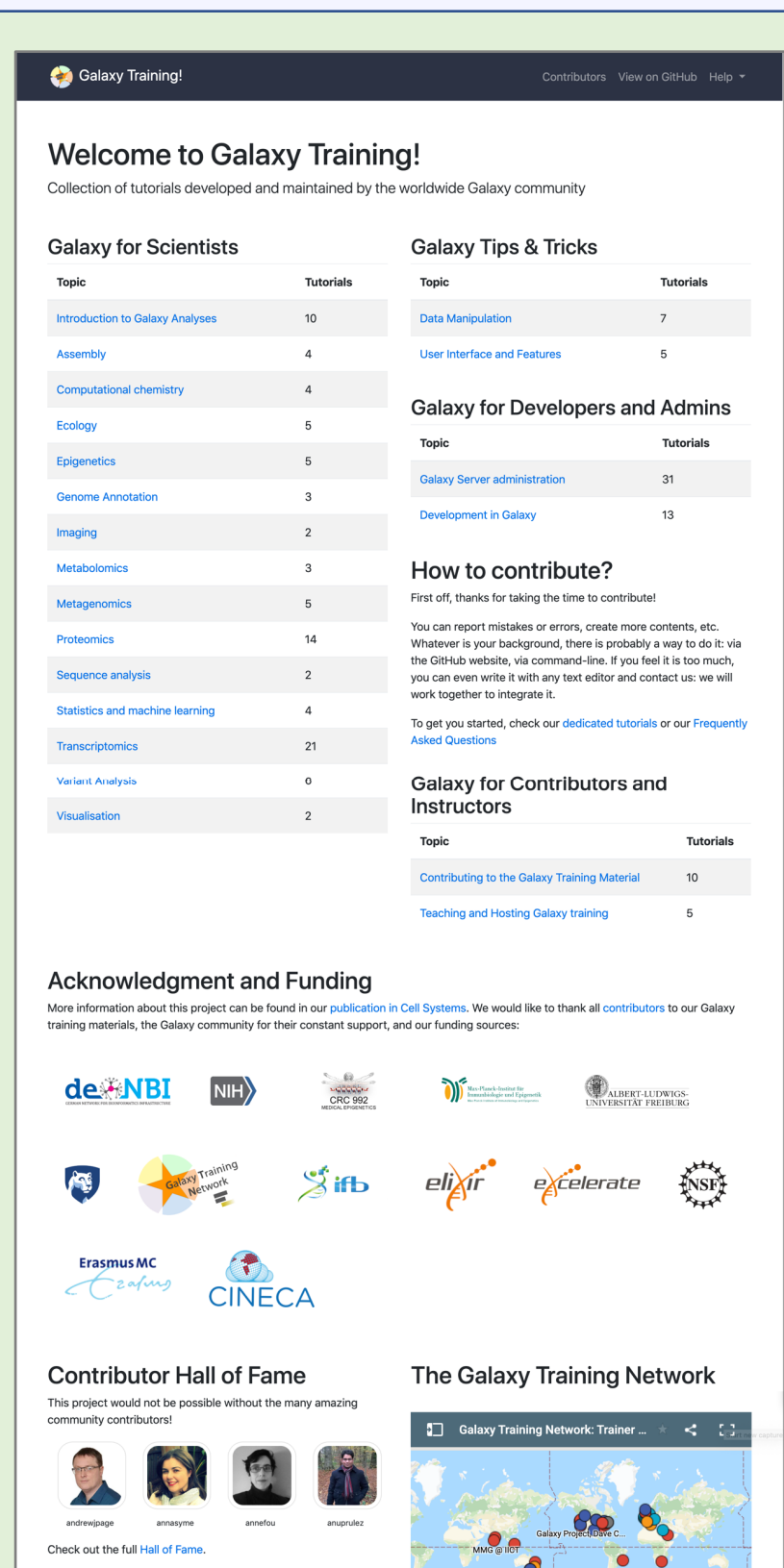
Learn: Everyone



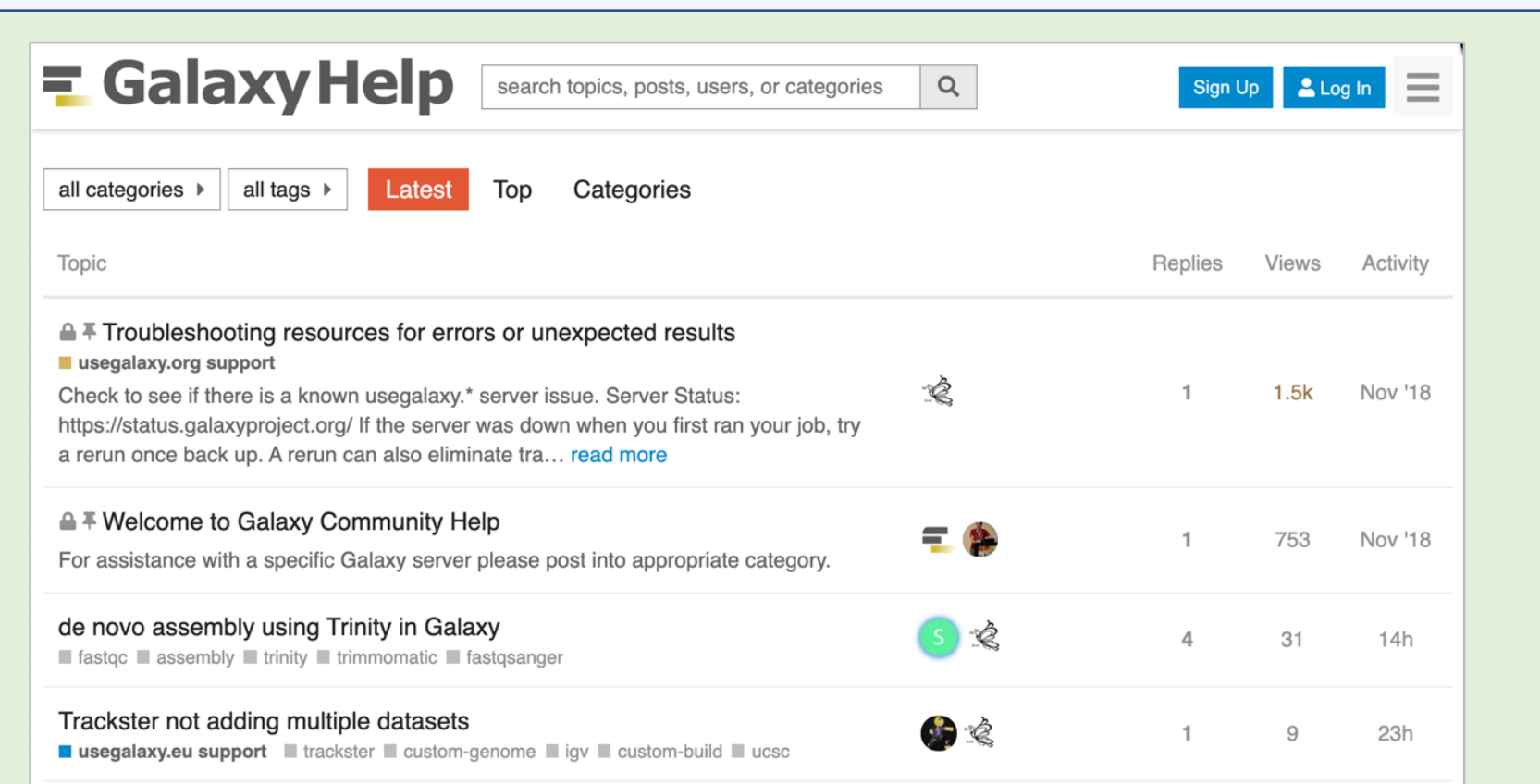
Hands-on tutorials, slides, training datasets, exercises, ...

See Poster P141: The Galaxy Training Network: A Community Based Training Resource

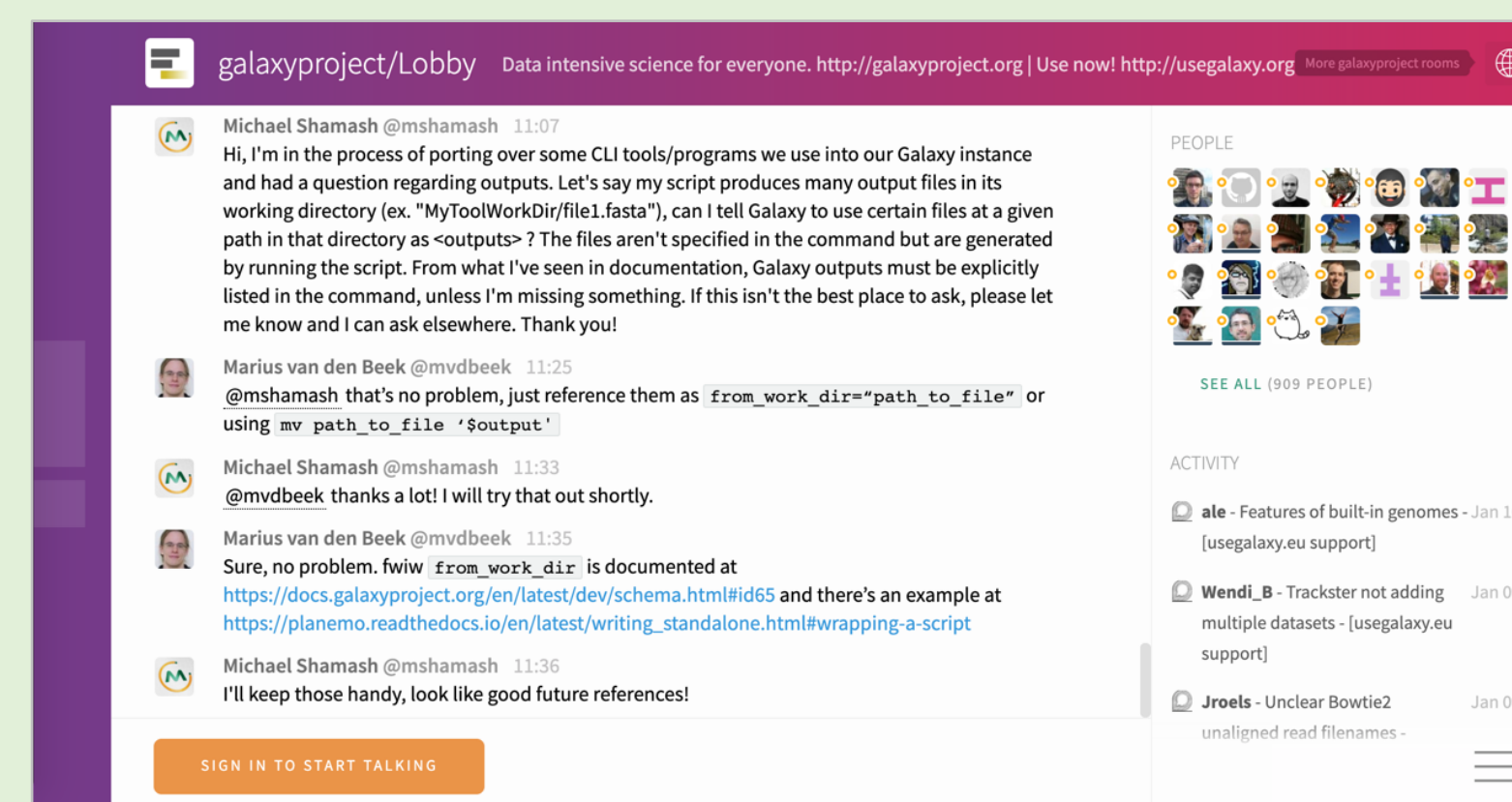
training.galaxyproject.org



Ask: Support



help.galaxyproject.org

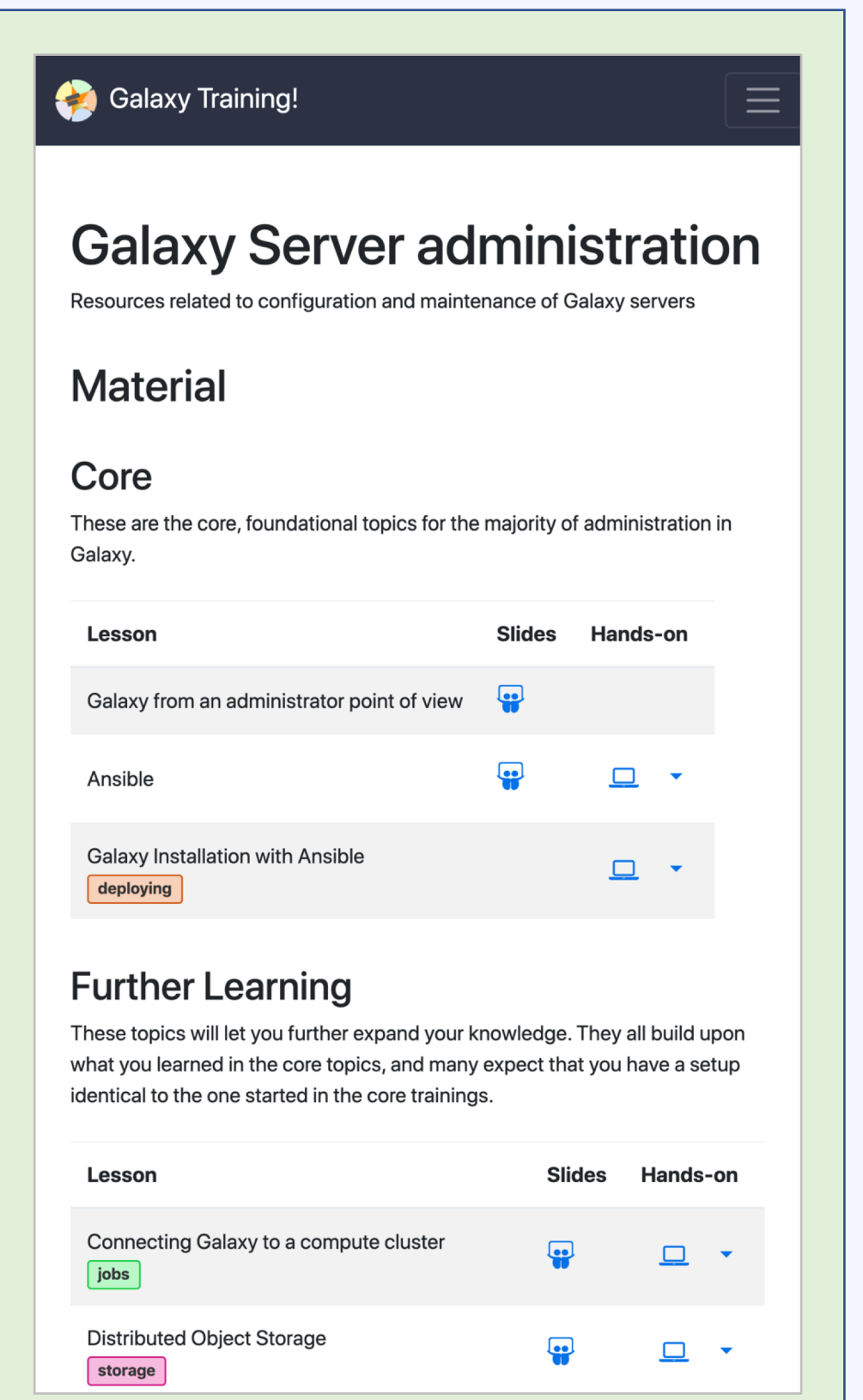


gitter.im/galaxyproject

Enable: Providers

Galaxy is deployed locally, from lab servers to institutional and consortium clusters and clouds.

getgalaxy.org



Publish: Tool Devs

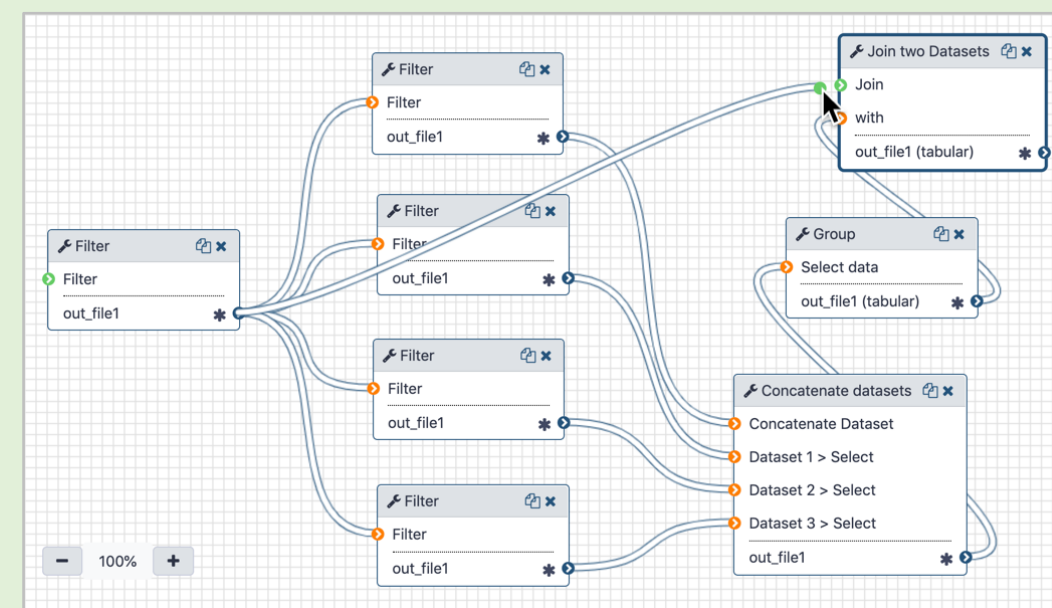


The Galaxy ToolShed contains over 7,000 tools, datatypes, & workflows, all defined for Galaxy, and easy to install. Wrapping your tools for Galaxy makes them easy to deploy and use.



Repeat, Share, Publish

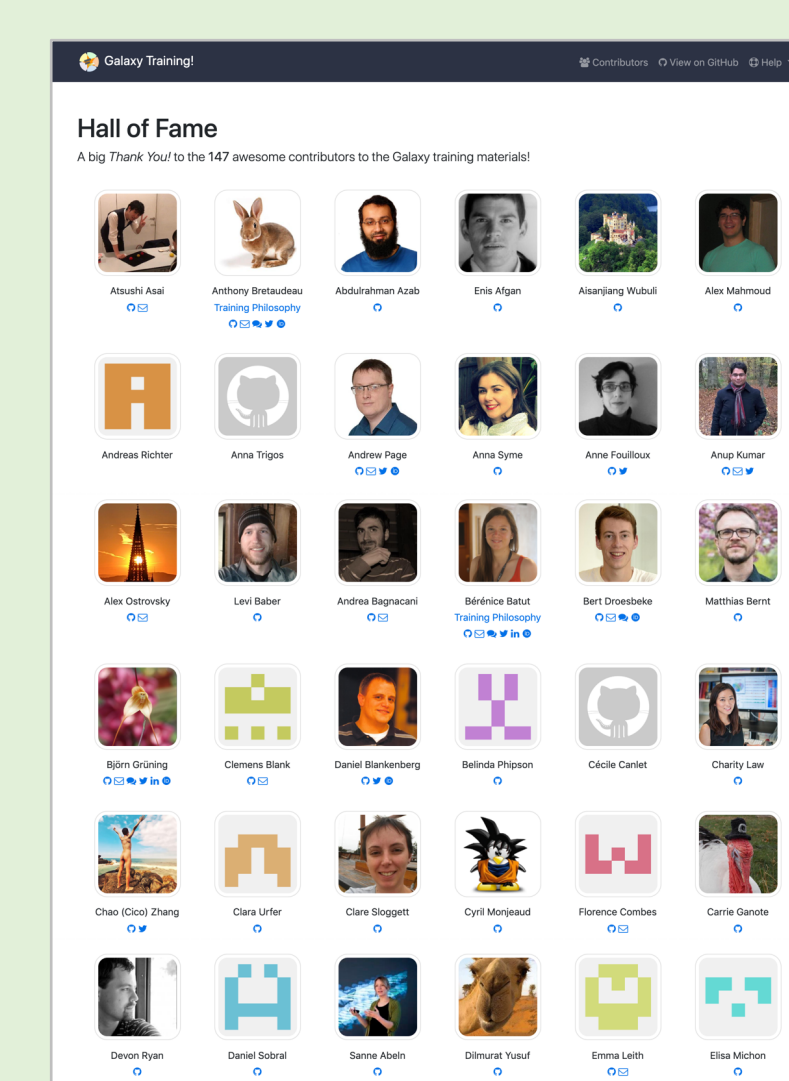
Workflows are created from scratch, or extracted from past analyses.



Individual tools through entire analyses can be rerun. Galaxy saves every dataset, tool, and tool parameter.

Galaxy supports sharing and publishing analyses.

Participate: Everyone



BlackDuck Open Hub:

Over the past twelve months, 157 developers contributed new code to Galaxy ... This is one of the largest open-source teams in the world, ...

openhub.net/p/galaxybx/factoids

galaxyproject.org