Today's Schedule

9am – 9:30am Introduction

9:30-10:30 RNA-Seq Demo

10:30 – 11am Workflows

11am – Noon Lunch

1pm – 2pm HPC overview

2pm – 5pm Unstructured Galaxy Help

National Center for Genome Analysis Support: http://ncgas.org

Introduction to Galaxy

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Summary

- Who is NCGAS?
- Galaxy what is it?
- Galaxy Anatomy

Who is NCGAS?

The National Center for Genome Analysis Support is based in Indiana but caters to a national audience with support from the NSF.

We provide computational resources and support for genomics, transcriptomics, and meta-projects.

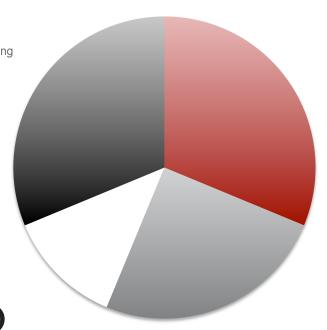












IU members

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What is Galaxy?

At its core, Galaxy is a web-based tool for analyzing data

- It provides a set of tools that one can apply to the data
- It provides a trail of analyses one has run
- It allows for sharing of data sets, methods, and workflows



What is Galaxy?

Galaxy was originally conceived as a tool to allow biologists to get work done without the need for training as an informatician

 Galaxy's purpose depends greatly on the tools made available to the user – these could be any command-line available software

There are many Galaxies in the Universe!

Galaxy Main – hosted at Texas Advanced Computing Center





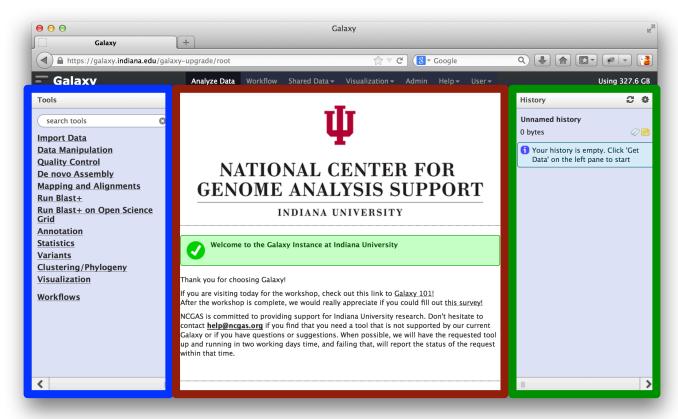
Galaxy-P – Proteomics-focused, hosted at U of Minnesota

IU Galaxy – just another Galaxy "instance" – each is autonomous and specialized



Galaxy Anatomy and Physiology

Tool bar – contains the available steps to apply to



History – shows steps previously taken to manipulate input data sets

Focus pane – shows options, parameters, and output for current item.

Galaxy – the perfect technology?

There is no such thing as a perfect technology, but Galaxy might be good for you if you –

- Don't like looking up parameters/reading man pages
- Don't like managing job submissions/scripts
- Want an automatic trail of data analysis steps
- Like pastel colored web interfaces