

BioCreative

2016

ICBO

ICBO Workshop #W03



Organizer: Dave Clements

www.icbo-conference.org

Integrating Ontological and Genomic Analysis with Galaxy

ICBO 2016

August 1, 2016 Corvallis, Oregon

Dave Clements Galaxy Team Johns Hopkins University http://galaxyproject.org/



#usegalaxy @galaxyproject

Agenda

A quick stroll through the Galaxy

Demonstrate Galaxy by addressing a specific question

Galaxy Ecosystem

What is Galaxy?

Keith Bradnam's definition:

"A web-based platform that provides a simplified interface to many popular bioinformormatics tools."

From

"13 Questions You May Have About Galaxy"

http://bit.ly/13questions

Galaxy is available several ways ...

As a free for everyone service on the web: usegalaxy.org







Welcome to the Metabiome Portal @ GMU
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Cistrome



A Galaxy Server dedicated to ChIP-* analysis



Public Galaxy Servers and still counting



The Genomic HyperBrowser

Powered by Galaxy





Whale Shark Galaxy! ×G



Genomic analysis tools for southern and Mediterranean plants

bit.ly/gxyServers

Galaxy is available as Open Source Software

Galaxy is installed in locations around the world.

http://getgalaxy.org

Galaxy is available on the Cloud







The Open Source Toolkit for Cloud Computing



We are using this today

http://aws.amazon.com/education http://globus.org/ http://wiki.galaxyproject.org/Cloud

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Galaxy Ecosystem Update

Quick Poll: Are you ...

1. A bioinformatics novice

2. A bioinformatics apprentice

3. A bioinformatics guru

Yes, those are your only choices.

Tutorial

http://bit.ly/ICBO_GXY_PDF

cloud1.galaxyproject.org
cloud2.galaxyproject.org
cloud3.galaxyproject.org

Demonstrate Galaxy by addressing a specific question

Why?

Goal is to demonstrate how to use Galaxy in concert with ontological tools.

Warning: No meaningful science will happen in this workshop.

http://bit.ly/ICBO_GXY_PDF

Specific Question: Repeats in coding regions

Which genes physically overlap with large numbers of repeats?

Once we have that we'll look for GO term enrichment and then use Galaxy to analyze the reported enrichment

http://bit.ly/ICBO_GXY_PDF

Repeats in Coding Regions: A General Plan

- Get some data
 - Get Data → UCSC Table Browser
- Identify which genes have overlapping repeats
- Count repeats per gene
- Look for enrichment

http://bit.ly/ICBO_GXY_PDF

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Other Features and Galaxy Ecosystem Update

2016 Galaxy Community Conference (GCC2016)

June 25-29, 2016 Bloomington, Indiana galaxyproject.org/GCC2016

Slides & posters are now online. Video will be shortly



LE CORUN

Le Corum Conference centre

gcc2017.sciencesconf.org

November 7-11



Salt Lake City, Utah



#GAMe2017 @EMBL_ABR

GAMe 2017

Galaxy Australasia Meeting

3 - 9 February, Melbourne, Australia

www.embl-abr.org.au/GAMe2017

Galaxy Community Resources: Galaxy Biostar Tens of thousands of users leads to a lot of questions. Absolutely have to encourage community support. Project traditionally used mailing list Moved the user support list to Galaxy Biostar, an online forum, that uses the Biostar platform



https://biostar.usegalaxy.org/

Scaling Training



Galaxy Training Network bit.ly/gxygtn

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ENCE-IT is supported by the NEE cyberSBES program award No. 13233762

The Galaxy project a supported in part by MiR, NHOR, and the Huck matitudes of the Life Sciences.

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CpG-Island Analysis Illumina Guanosine Cytosine BeadChip

LimiT

ADMIRE

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Proteomics

Metabolomics

Natural Language

Image Analysis

Climate Change

Social Science

Cosmology







Analyses and Bioinformatics for Marine Science



The Galaxy Team



Enis Afgan

Dannon Baker Dan

Dan Blankenberg

Dave Bouvier

Marten Cech

John Chilton



Dave Clements

Nate Coraor

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Jeremy Goecks

ks Sam Guerler

Jen Jackson



Ross Lazarus

Anton Nekrutenko

Nick Stoler

James Taylor

Nitesh Turaga

http://wiki.galaxyproject.org/GalaxyTeam

Acknowledgements

You

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Thanks