The Galaxy Visual Analysis Framework

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Vision

Galaxy is an open, Web-based platform for accessible, reproducible, and transparent computational biomedical research

What is Galaxy?

GUI for genomics

for complete analyses: analyze, visualize, share, publish

A free and open Website (http://usegalaxy.org)

integrating a wealth of tools, compute resources, terabytes of reference data and permanent storage

Open source software that makes it simple to

- integrate your own tools and data
- customize and run on your own resources

An open, Web-based approach for

making highly interactive visual analysis

distributed, extendable, sharable, fast

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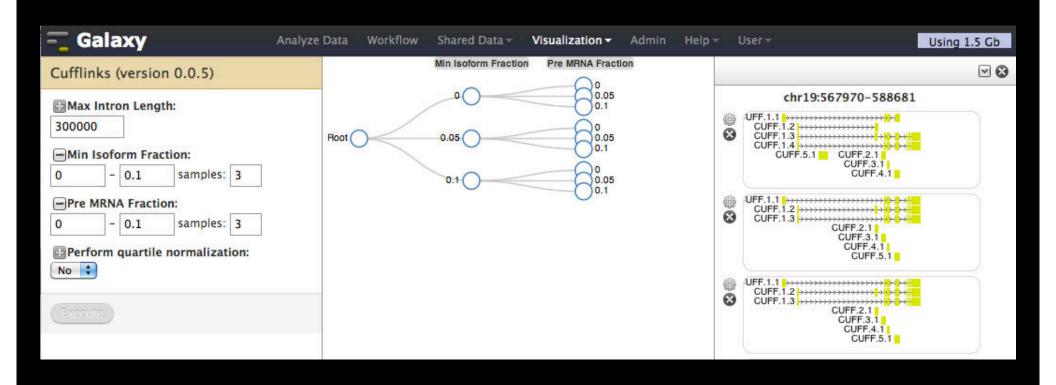
your own huge datasets

Let's go!

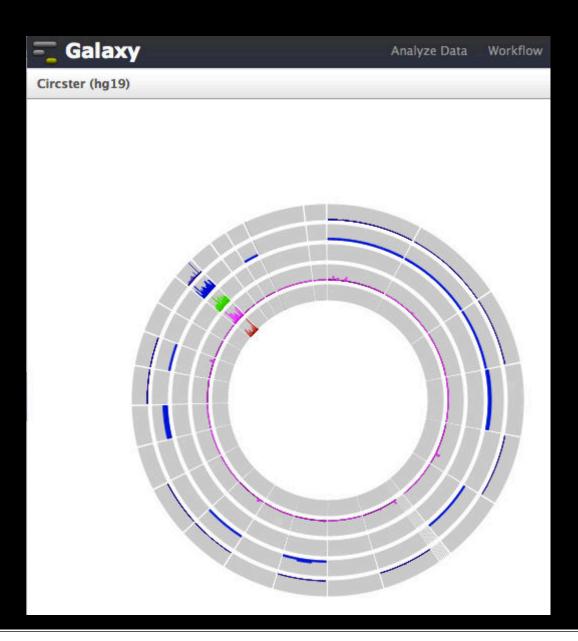
Trackster



Paramamonster



Circster



Trackster

Completely Web-based

only a modern Web browser is needed

Supports arbitrarily large NGS datasets

SAM/BAM, BED, GFF/GTF, VCF, WIG

Highly flexible

e.g. custom rainbow tracks

Integrated with Galaxy tool framework

- dynamic filtering
- re-running tools

"Paramamonster"

Visualization for

- tool parameter space
- outputs from different settings

Can easily find good settings by visual inspection

for many settings, across multiple regions

Can explore parameter space systematically or ad-hoc

Circster

Circos-like visualization that provides genome-wide views

Complements Trackster

Very much a work in progress

Summary

Galaxy is a Web-based platform for doing all facets of genomics analysis

Galaxy visualizations are

- Web-based
- highly interactive and dynamic
- integrated with Galaxy tools
- sharable and publishable

Only a Web browser is required for using any and all Galaxy features and visualizations







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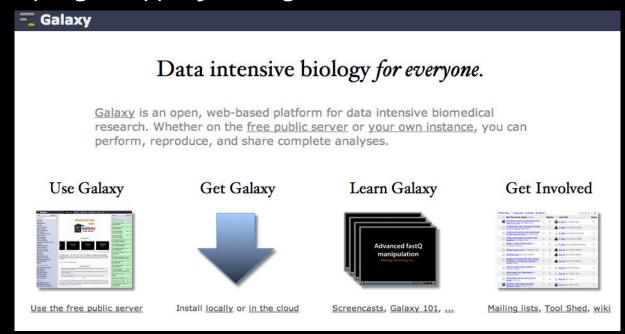
Greg von Kuster



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Thanks! Questions?

http://galaxyproject.org



Galaxy publications: http://galaxyproject.org/wiki/Citing

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