

Galaxy for Genome Annotation

ER, BG, ND, AB

GGA

Wha

Projects

Docker Image

Real Life

Backgrour Current

Current Expansion

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Galaxy Genome Annotation: Galaxy and GMOD for Annotation, Teaching, and Genomic Databases

Eric Rasche, Björn Grüning, Nathan Dunn, Anthony Bretaudeau

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Galaxy for Genome Annotation

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Galaxy has room to expand into the Genome Annotation space. Needs community, we are that community. Needs tooling, we're building that. Needs integrations, we're building that.

Who are we?

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Who What

Projects

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Expansion

Futu

- Björn Grüning
- Eric Rasche
- Anthony Bretaudeau
- Peter van Heusen
- Nathan Dunn
- Eduardo de Paiva Alves
- Suzanna Lewis
- Torsten Seemann
- **■** (...you!)

GGA What are we building?

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Real Life
Background
Current
Expansion

■ Small components enhancing integration

- Standalone services augmenting the experience
- Which all build to: Completely pre-configured Galaxy+Apollo deployments with optional Tripal/Chado additions

Overview of Projects

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GGA Who What

Projects

Docker Images

Backgroun Current Expansion Future ■ Docker Image: Galaxy + Annotation Tools (Apollo Tools, Tripal Admin Tools, Circos, JBrowse, BLAST+, InterProScan, Glimmer, Augustus, FASTA manipulation tools, Spades, Mira, CD-Hit, ClustalW, AntiSmash, mummer, EMBOSS, BLAST, Diamond, Blast2GO)

- Dockerized GMOD Deployment (Most GMOD projects pre-configured to work together seamlessly)
- Apollo Client library (+parsec like tool, "Arrow")
- Various Apollo support projects (git-backup, experimental google docs integration)

Docker Images

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Projects

Docker Images

Galaxy/Apol Real Life

Current Expansion Future Docker images for major GMOD projects (Apollo, Chado, Tripal)

- Galaxy image with huge array of tools
- Docker-compose configuration for entire deployment

GGA Galaxy / Apollo Bridge

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Projects

Docker In

Docker Images Galaxy/Apollo

Background Current Expansion Future

O&A

- Initially simple tooling to move data in/out of Apollo
- Expanded to include annotation automation, tools for creating / removing "human-created" annotations.
- Jenkins scripts ran functional annotation workflows automatically, regularly.
- The Galaxy-Apollo bridge grew out of use in an undergraduate course
- We have improved upon them as students find new bugs and corner cases we did not expect

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Who What

Projects

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Real Life

Backgroun Current Expansion Future

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- Used in Course
- Current Course Applications
- Ongoing Expansions
- Future Work

Course Background

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Who

Projects

Docker Im

Galaxy/Apol

Real Life

Background

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- Undergraduate phage annotation course
- Genome sequence to publication
- (Parallel track for environmental sample to isolated phage)

Current Applications

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Galaxy/Ap

Real Life
Background
Current
Expansion
Future

■ Apollo for annotation

- But how to democratize annotation?
- Galaxy Tools can create genomes, update permissions
- (But we do end up having to check user's apollo permissions at every step)

Applications

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Real Life

Current Expansion ■ Using automated Chado export in Apollo

lacksquare Adding on Tripal + JBrowse-based public sequence display

■ Internal sample tracking database

■ More GMOD projects to come?

Future Work

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Who What

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Docker In

Galaxy/And

Real Life Backgroun

Expansion Future

Q&A

- More tools (IUC, GGA)
- More annotation community

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Q&A

Thank you

GGA GitHub galaxy-genome-annotation.github.io GGA Gitter gitter.im/galaxy-genome-annotation/Lobby

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