Running a Local Galaxy

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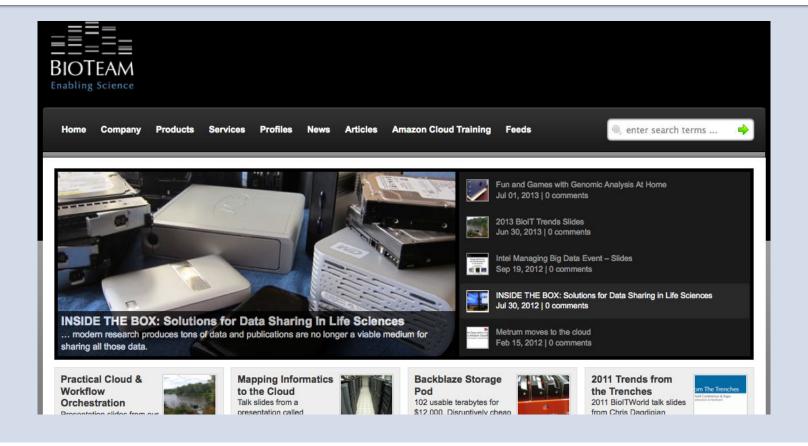


Agenda



- Who is BioTeam
- Why Local Galaxy
- Running Galaxy Locally
- Local Galaxy Best Practices

Who is BioTeam





Who We Are



Over a decade of Life Sciences IT consulting

- We are scientists forced to learn IT to get research done
- Served over 400 organizations
 - Academic, Non-profit
 - Government, Military
 - Pharm, AgBio, Biotech
 - Cloud & Datacenter Providers





Who We Are



Active contributors to open-source projects

















Bridging the IT Gap



·BioTeam

Encapsulate IT best-practices expertise to eliminate redundant effort spent designing and building infrastructure

Galaxy Project

Decrease the barrier to entry into data analysis by improving accessibility of the Galaxy platform



Bridging the IT Gap



 BioTeam offers a all-in-one solution to help run a local instance Galaxy





- BioTeam is the the official appliance provider for Galaxy
 - Exclusive partnership with the Galaxy Team
 - Donations back to the Galaxy Project

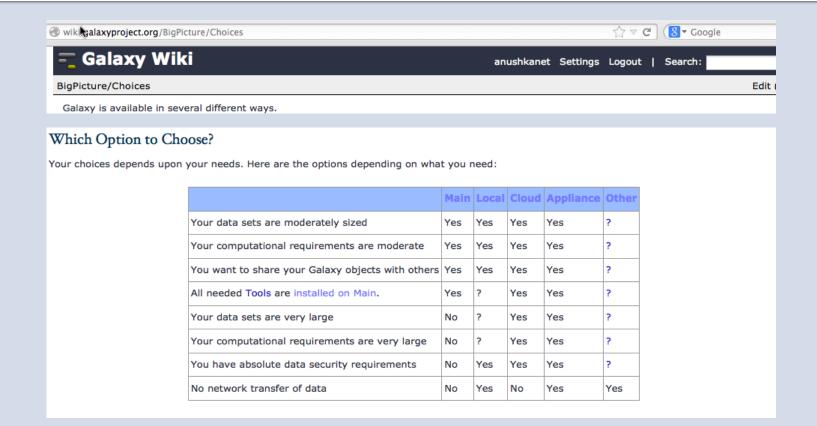


SlipStream Galaxy



FEATURES		
Powerful Server	Complete analysis tasks quickly	
Optimized Galaxy	Configured for stable production use	
Analysis Tools	Provides comprehensive set of open-source tools	
Automated Updates	Software can be updated automatically	
Preinstalled Datasets	Includes 5 model organisms (additional upon request)	
Open Platform	Install other software for your own use	
Price: \$19,995 (USD)		

Why a Local Installation





Current Challenges



- Galaxy Main is a fantastic public resource!!
- Limitations due to popularity
 - Wait times
 - Job and storage quotas
 - Data transfer bottlenecks
 - Pre-defined set of tools and datasets



Current Challenges



- Cloud is a scalable option but has its challenges
 - Understand how to manage pricing
 - Data transfer bottlenecks
 - Familiarity with cloud (e.g. Amazon Web Services)
 - Dealing with sensitive data

Running Galaxy locally solves these issues



Be Prepared



- IT/informatics expertise
 - Acquire and set up infrastructure
 - Install Galaxy, tools, necessary dependencies
 - Optimize/customize for your use cases
- Define policies
 - Managing usage
 - Data back-up
 - Software updates/upgrades



Be Prepared



- Informatics support
 - Handle user questions/requests
 - Gather user feedback
- Ongoing dedicated resource
 - Manage updates
 - Facilitate user support
 - Maintain infrastructure



Benefits of a Local Galaxy



It's all about control

YOU CONTROL	BENEFIT
Amount of storage	Handle large datasets
Type of hardware	Run compute intensive jobs
What tools to install	Customize to your research
Data access	Granular control of security
Networking architecture	No data transfer bottleneck
Software behavior	Optimize how jobs are run

Local Instance of Galaxy

Galaxy Wiki

anushkanet Settings Logout | Search:

Admin/Get Galaxy

Edit ((

Get Galaxy: Galaxy Download and Installation

In addition to using the public Galaxy server (a.k.a. Main), you can also install your own instance of Galaxy (what this page is about), or create a cloud-based instance of Galaxy. Another option is to use one of the ever-increasing number of Public Galaxy Servers hosted by other organizations.

See Big Picture: Choices for help on deciding which of these options may be best for your situation.

Reasons to Install Your Own Galaxy

You only need to download Galaxy if you plan to:

- 1. Develop it further
- 2. Add new tools
- 3. Plug-in new datasources, or
- 4. Run a local production server for your site because you have
 - 1. Sensitive data (e.g., clinical)
 - 2. Large datasets or processing requirements that are too big to be processed on Main

Installation Procedure

The installation procedure is simple and is nearly identical for UNIX/Linux and Mac OS X. We are no longer supporting the Windows platform with

Contents

- 1. Reasons to Install Your Own Galaxy
- 2. Installation Procedure
 - 1. Check your Python version
 - 2. Get the latest copy from the repository
 - 3. Start it up
 - 4. Join the Mailing List
 - 5. Keep your instance backed up
 - 6. Keep your code up to date
- 3. Advanced Configuration
- 4. Other Help



Customizing Galaxy



- Customizing a local installation
 - Customize galaxy itself
 - Install 3rd party/commercial tools
 - Develop your own tools
 - Add shared genome builds
 - Integrate with instruments
 - Manage sensitive or proprietary data



Base Software



Software	Comment
Python	Version 2.6 or 2.7
Galaxy	Most recent version
Web Server	To enable web hosting
Database	To store histories, users, track datasets, etc.
Analysis Tools	Need to follow install directions for each individual tools

http://wiki.galaxyproject.org/Admin/GetGalaxy



Simple Start-up



Basic install of Galaxy

%

hg clone https://bitbucket.org/galaxy/galaxy-dist

% sh run.sh

In your browser type: http://localhost:8080



Additional Configuration



- Install tools from toolshed
- Global config file: universe_wsgi.ini
 - Over 100 configurable parameters
 - Add admin users
 - Configure all galaxy settings
- Tool behavior
 - tool_conf.xml
 - Individual tool wrappers (also xml files)



Galaxy for NGS



- · Galaxy for NGS requires additional tools
- http://wiki.galaxyproject.org/Admin/Tools/Tool %20Dependencies
- Set up reference genomes or fetch indexes



Production Instance of Galaxy



Scalability

- Handle more users (>5)
- Run more jobs (>8 concurrent)
- Support large datasets (>3TB)

Efficiency

- Schedule jobs
- Optimize runs
- Manage data



Production Instance of Galaxy



Hardware	Recommendation
# of Cores	32-64 processors
RAM	256-512 GB
Storage Amount	>10TB

Best Practices



Login | Search:

Develop/BestPractices

page was renamed from Develop/Best Practices

Galaxy Software Development Best Practices

Lists software development best practices for the Galaxy Project. These are works in progress and practices vary in how broadly they have been applied.

Contents

- 1. Metastandards
- 2. Python Standards
- 3. JavaScript Standards
 - 1. Backbone
- 4. Email Threads

Develop

Mercurial
Source Doc
Best Practices
Bitbucket
Issues & Requests
Data Model
Search



Galaxy Best Practices



- Turn off developer settings
 - Speed up the galaxy server
- Switch databases (e.g. PostgreSQL)
 - handle multiple database requests at once
- Switch web servers (e.g. Apache,nginx)
 - Handle file transfers, external authentication



Galaxy Best Practices



- Minimize redundant storage
 - Galaxy stores everything!
 - Filesystem compression
 - Deleting datasets and histories
- Improve Data Transfer
 - FTP, HTTP, SCP are slow
 - Consider data transfer products (e.g. Aspera, Globus)



Galaxy Best Practices



- Dev, Test, Production environment
- Automated testing
- Leverage toolshed
- DEDICATED RESOURCE!!!
 - ½ FTE to support a large production installation



Take Away



- Galaxy is a fantastic public resource
- Local Galaxy offers a high level of control over your analysis environment
- There is tremendous value in being aware of the underlying components of Galaxy
- The flexibility makes it complex to manage so plan accordingly



SlipStream Galaxy



Thank You

Contact BioTeam to discuss your Galaxy needs

•Website: <u>www.bioteam.net</u>

·Email: anushka@bioteam.net



EXTRA SLIDES

SlipStream Galaxy Appliance



Lowering the barriers







Powerful dedicated desktop server pre-configured with a fully operational production instance of Galaxy



SlipStream Galaxy



	Hardware Specifications
CPU	2x Intel® Xeon® Processor E5-2690, 8-core (16 cores total)
Memory	24x 16 GB RDIMM (384 GB)
Storage	7x 3TB SAS 6 Gbps HDD (16 TB usable) 1x 100GB SSD
Network	Dual Gigabit network adaptor
Power	Dual redundant power supplies



SlipStream Galaxy



FEATURES		
Optimized Galaxy	Production configuration, optimized data transfer	
Analysis Tools	Comprehensive set of open-source tools	
Automated Updates	All software can be updated automatically	
Preinstalled Datasets	5 model organisms (additional upon request)	
Grid Compute	Grid Engine -based job management	
Hardware Maintenance	Warranty includes maintenance	
Open Platform	Install other software for your own use	
Price: \$19,995 (USD)		



Partnering with Thought Leaders



•EARLY ACCESS PROGRAM (Limited Availability)

- Seamless Adoption
- Dedicated Support
- Workflow Generation
- Early Development Partner Feedback
- "A device that centralizes functions with respect to data archives, storage, and analysis is a tremendous aid." - Ed DeLong, MIT



SlipStream Galaxy



•Become an Early Access Partner Today!!

•Web:

www.bioteam.net/slipstream/galaxy-edition