

Building Excellence in Genomics and Computational Bioscience

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TGAC Galaxy server and training

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Outline



- Introduction
- Status of the new Galaxy server
- Ongoing projects
- Future work

My role @ TGAC



- Installation and maintenance of a new Galaxy platform, hosted at TGAC, for internal bioinformaticians and external researchers
- Exploit TGAC's hardware and software expertise and resources, promoting the Centre as a National Capability for UK bioscience
- Training on Galaxy for BBSRC institutes
- Network with the Galaxy community

What is Galaxy?



- Web-based platform for data-intensive biomedical research
 - Open source (Academic Free License)
 - Developed by the A. Nekrutenko lab at Penn State and the J. Taylor lab at Johns Hopkins University
- >2500 citations http://www.citeulike.org/group/16008/
- >70 public Galaxy servers
 - None in the UK!

Galaxy features

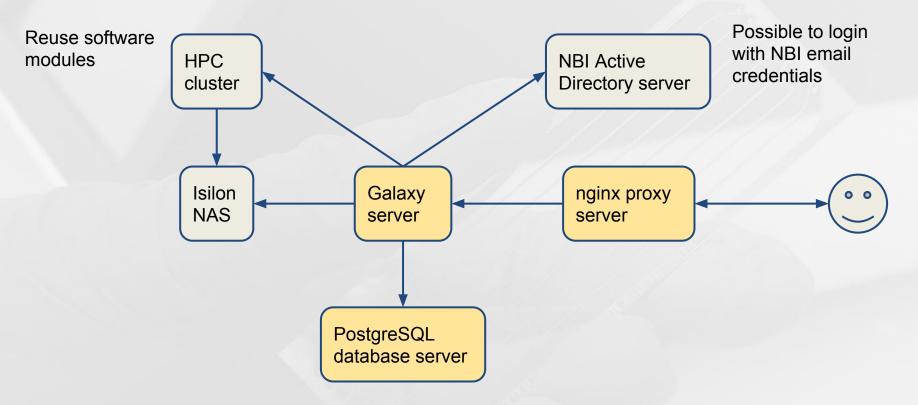


- Uniform access to bioinformatics tools
 - every software can be embedded
- Tool Shed: free tool store https://toolshed.g2.bx.psu.edu/
- Workflow: editing / extracting; reuse on different inputs; automatic scheduling of jobs
- Data visualisations
- Reproducibility: every analysis step is tracked
- Sharing of datasets, workflows and histories
- Scalability: can use cluster or cloud resources

New TGAC server architecture



https://galaxy.tgac.ac.uk/



TGAC server tools



- NGS mapping: BWA, Bowtie, Bowtie2
- RNA-Seq: TopHat, Cufflinks suite,
 SortMeRNA, StringTie
- De novo assembly: ABySS, SPAdes
- Phylogenetics: ClustalW, T-Coffee, MAFFT
- Variant analysis: FreeBayes
- FastQC, Samtools, bedtools, Picard, NCBI BLAST+, EMBOSS

Galaxy server plans



- Presently accessible only inside the NRP
 - We will open the server to the other BBSRC institutes...
 - ...and then to the rest of the world
 - Sustainability
- More tools and reference genomes/indexes
- Training!

Training plans



- Presently one-to-one
- Survey to be sent to NRP to investigate the training needs in Galaxy
 - expectations, personal projects/data
- First TGAC course to be held early 2016
 - Second NRP event mid 2016
 - Focused events: RNA-Seq, metagenomics...
- Train the trainer for Galaxy?

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Galaxy Team and community