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MyJGI: Tools for Collaborators

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Authors

Fazo, Joni Pletcher, David Taylor, Kristen <u>et al.</u>

Publication Date

2008-03-27



MyJGI: Tools for Collaborators

Joni Fazo¹, Annette Greiner², Anthony Kosky², Rene Perrier², David Pletcher¹, Kristen Taylor¹, Arkady Voloshin²

The JGI Informatics department launched <u>http://my.jgi.doe.gov</u>, aka MyJGI, in October of 2007. This new web site provides information and tools useful to collaborators with sequencing projects at JGI.



General Info

Step-by-step instructions for collaborators initiating a project with the JGI:

- Sample user agreement for download.
- Recommended protocols offering successful examples for preparing samples or libraries to be sent to the JGI.
- Links to JGI's policies on project scheduling, finishing, publication, and data release



¹DOE Joint Genome Institute - Lawrence Livermore National Laboratory ²DOE Joint Genome Institute - Lawrence Berkeley National Laboratory

| MyJGI | |
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| Please Log In | |
| Forgot your password? Need a login? | |
| User Name: | |
| | |
| Password: | |
| | |
| Log In | |
| Please make sure cookies are enabled in your browser. | |
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Logins

The Sample Info and Status Reporter areas require a user name and password to access. Collaborators may request a login from their JGI project manager. We have recently implemented single sign-on for MyJGI, so that a user who has logged in to use either the Sample Info area or the Status Reporter can use the other without logging in again.

Sample Info

Use CSI, the Collaborator Sample Information tool, to efficiently get your samples into the JGI pipeline:

- 1. Prior to shipment of samples to the JGI, use CSI to provide project managers and scientists with required sample information, such as gel image, volume and weight.
- 2. JGI project managers then use internal web forms to review and approve each sample for shipment.
- 3. The process integrates with downstream systems such as the JGI Global Project Management Tracking System (GPTS) and LIMS.



Status Reporter

Provides collaborators with real-time information about the status of each of their sequencing projects.

- For most projects, includes detailed information about the planned lifecycle of the project, release dates and key milestones.
- We have recently added full support for eukaryote projects, and all active projects now show at least basic sequencing status information.
- Once a collaborator logs in, they are presented with a list of their own proposals, from which they can select one to view status information. A single click brings up the status information for all projects that are part of that proposal.

| Evolutionary Genomics of Neurospora | Conception day |
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| | Carl and the Property Parents are |
| -2,07 -22 | Crenolitrix polyspora metagenome Prant Navajer <u>Satabary</u> Prant Solan Metagenarie - Pr Majner Mohael |
| | This project is in the Production phase. Its status is Phase Bequeroing Phase partial plus project namager if you have more detailed internation. |
| echnologies used: | |

dynamic use of CSS.