Lawrence Berkeley National Laboratory

Recent Work

Title

Automating the JGI's Community Sequencing Program

Permalink

https://escholarship.org/uc/item/9kn3p6qq

Authors

Patil, Jayant Fazo, Joni Kobayashi, Art et al.

Publication Date

2007-03-26

Automating the JGI's Community Sequencing Program

Jayant Patil¹, Joni Fazo², Art Kobayashi², Annette Greiner¹, Arkady Voloshin¹, Pat Kale², Marsha Fenner¹, David Bruce³, Jan-Feng Cheng¹ and Jim Bristow¹

¹Lawrence Berkeley National Laboratory / DOE Joint Genome Institute 2800 Mitchell Drive, Walnut Creek, CA 94598

²Lawrence Livermore National Laboratory / DOE Joint Genome Institute 2800 Mitchell Drive, Walnut Creek, CA 94598

²Los Alamos National Laboratory / DOE Joint Genome Institute 2800 Mitchell Drive, Walnut Creek, CA 94598

March 26, 2007

ACKNOWLEDGMENTS:

The work conducted by the U.S. Department of Energy Joint Genome Institute is supported by the Office of Science of the U.S. Department of Energy under Contract No. DE-AC02-05CH11231

DISCLAIMER:

[LBNL] This document was prepared as an account of work sponsored by the United States Government. While this document is believed to contain correct information, neither the United States Government nor any agency thereof, nor The Regents of the University of California, nor any of their employees, makes any warranty, express or implied, or assumes any legal responsibility for the accuracy, completeness, or usefulness of any information, apparatus, product, or process disclosed, or represents that its use would not infringe privately owned rights. Reference herein to any specific commercial product, process, or service by its trade name, trademark, manufacturer, or otherwise, does not necessarily constitute or imply its endorsement, recommendation, or favoring by the United States Government or any agency thereof, or The Regents of the University of California. The views and opinions of authors expressed herein do not necessarily state or reflect those of the United States Government or any agency thereof or The Regents of the University of California

[LLNL] This document was prepared as an account of work sponsored by an agency of the United States government. Neither the United States government nor Lawrence Livermore National Security, LLC, nor any of their employees makes any warranty, expressed or implied, or assumes any legal liability or responsibility for the accuracy, completeness, or usefulness of any information, apparatus, product, or process disclosed, or represents that its use would not infringe privately owned rights. Reference herein to any specific commercial product, process, or service by trade name, trademark, manufacturer, or otherwise does not necessarily constitute or imply its endorsement, recommendation, or favoring by the United States government or Lawrence Livermore National Security, LLC. The views and opinions of authors expressed herein do not necessarily state or reflect those of the United States government or Lawrence Livermore National Security, LLC, and shall not be used for advertising or product endorsement purposes.



Automating the JGI's Community Sequencing Program

Jayant Patil¹, Joni Fazo², Art Kobayashi², Annette Greiner¹, Arkady Voloshin¹, Pat Kale², Marsha Fenner¹, David Bruce³ Jan-Fang Cheng¹, and Jim Bristow¹

What is the Community Sequencing Program?

The Community Sequencing Program (CSP) provides the scientific community at large with access to high throughput sequencing at the Department of Energy's Joint Genome Institute (JGI) for projects of relevance to DOE missions. Through this program, the Department of Energy aims to advance sequence-based scientific research from a broad range of disciplines.

Proposal from the Scientific

How does the review process work?

All proposals undergo the following reviews:

- ➤ Technical review
- ➤ Proposal Study Panel (PSP) review
- ➤ Scientific Advisory Committee (SAC) review Proposals are evaluated on scientific merit and relevance to DOE mission.

DOE Mission

✓ Bioremediation ✓ Global carbon recycling ✓ Alternative energy production

What happens after proposals are accepted?

- Scope of Work, User Agreement and other required documents are approved.
- ➤ Proposals are converted to projects.
- Sequence traces are released to the Genbank Trace Archive on a regular basis.
- The JGI makes the assemblies, gene annotations, and analyses available to the community at large. Web User Interface



How to participate?

Established in 2004 as a manual process, we have since moved to an automated system that manages the proposal submission, review, and notification process. Scientists from around the world submit proposals through our website,

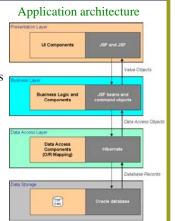
http://www.jgi.doe.gov/CSP/proposals.

- 1 DOE Joint Genome Institute BACE Division, Lawrence Livermore National Laboratory
- ² DOE Joint Genome Institute Genomics Division, Lawrence Berkeley National Laboratory
- 3 DOE Joint Genome Institute Los Alamos National Laboratory

Technology behind the CSP application

The automated CSP application is based on a J2EE multi-tier architecture:

- The *Presentation Layer* with web UI components built in Java Server Faces (JSF) and Java Server Pages (JSP).
- ➤ The *Business Laver* consists of JSF beans. servlets, and Java command classes.
- The Data Access Laver is an O/R mapping laver developed with the open-source technology Hibernate.
- ➤ Oracle database.





Iterative development

We are using an iterative software process with a focus on *Just In Time* software delivery to best support this exciting and important

2 to 3 weeks iterations that span multiple disciplines.

- ➤ Requirements ➤ Implementation
- > Testing
- ➤ Deployment

We are using Twiki for frequent team and user communication and documentation.

Support

- Development team provides daily support to worldwide CSP users.
- ➤ We are using the RequestTracker tool for management and tracking of requests and follow-up customer communications.

What's next?

- All JGI proposals will be accepted through the CSP application.
- ➤ Improve the user interface with AJAX technology.
- Enhance analytical reporting.