

# **The Genome Portal of the Department of Energy Joint Genome Institute**

**Henrik Nordberg\***, Michael Cantor, Serge Dusheyko, Susan Hua, Alexander Poliakov, Tatyana Smirnova, Inna Dubchak

Department of Energy Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA, 94598, USA;

\*Presenting author: [hnordberg@lbl.gov](mailto:hnordberg@lbl.gov) Henrik Nordberg

March 2014

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

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# PORTAL 7.0 RELEASED!!!

# Expanded Genome Search

Powerful new search tools support the growing volume of sequence data at the JGI. Users can locate and view sequencing projects using GenBank accession number, taxonomic information, Principal Investigator contact, and many other search terms.



*Home page with search*

Community Science Program JGI Proposals, that include multiple genomes, transcriptomes, or other data sets, have "group" portals aggregating individual portals within the proposal and providing a single access point for download of all data related to the proposal.

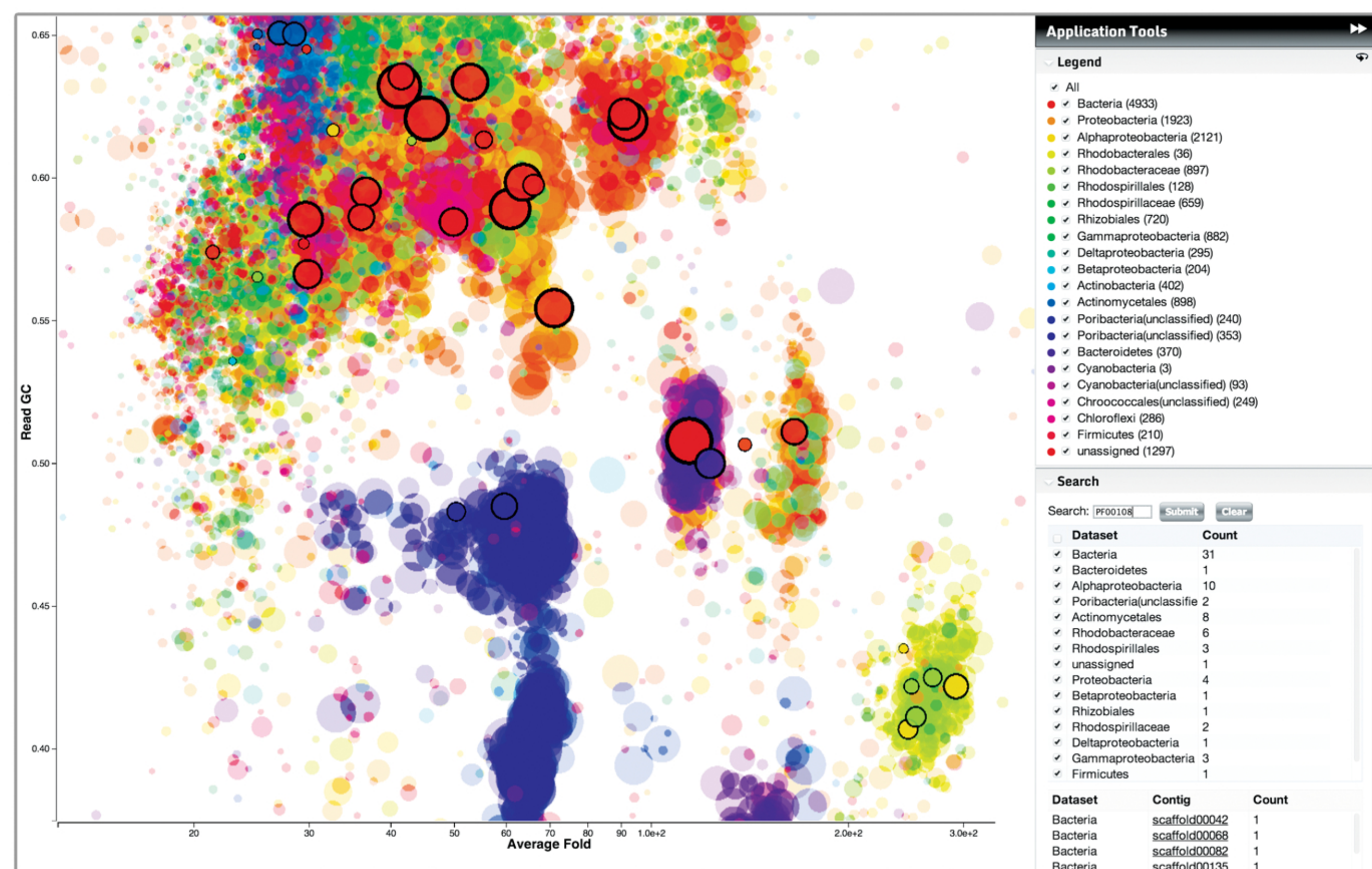
[illegible]

*Search results table*

## Cite Us

The genome portal of the  
Department of Energy Joint  
Genome Institute: 2014 updates.  
Nordberg H, Cantor M, Dusheyko  
S, Hua S, Poliakov A, Shabalov I,  
Smirnova T, Grigoriev IV,  
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2014,42(1):D26-31.  
hnordberg@lbl.gov  
ildubchak@lbl.gov

**genome.jgi.doe.gov**



### Metagenome visualization

# Metagenome Visualization

## Coming soon to a browser near you:

Hundreds of thousands of metagenome contigs in a dance of color and sound (okay, not sound) are displayed before your very eyes. Cluster them by GC content, coverage, read depth and other metadata to reveal their secrets. Search for annotations and drill in to individual contigs to browse their predicted coding regions. Compare samples. It's the JGI metagenomics visualization tool!

## The Download Area

## I. Download data with the web UI

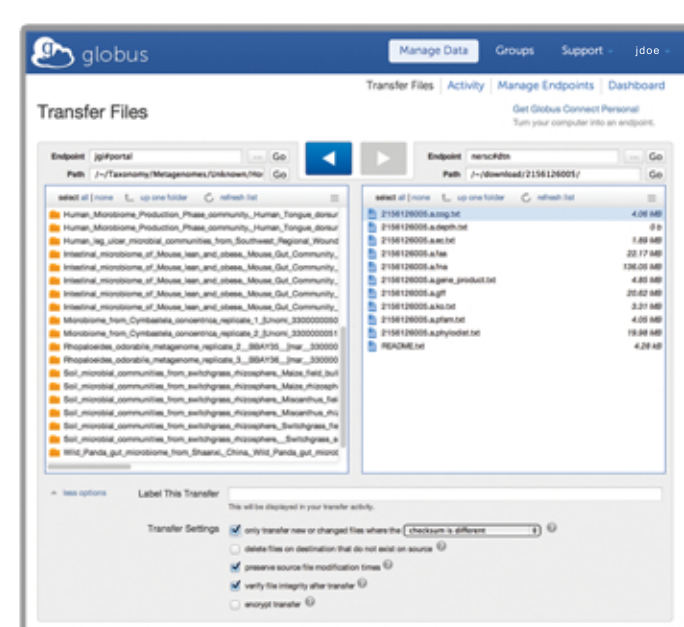
Find your portal of interest and click on the Download tab. Downloads are available in a tree structure that divides the files into logical groups for RAW files, assemblies, and annotation data.



portal web UI

## II. Download with Globus service (fast, reliable, convenient)

Need to download large files (up to several hundred gigabytes) or a large number of files? Data is now available via the public JGI endpoint on Globus (<https://www.globus.org>). Globus provides a user friendly interface to facilitate big data transfers. Use the `jgi#portal` endpoint with your JGI credentials.



*globus.org UI*

### III. Download using the API

Retrieve data programmatically using the expanded Download API.



*API script*

# More Projects! More Metadata!

The Genome Portal now provides unified access to nearly every sequencing project at the JGI. More than 15 000 DOE JGI projects of different types (standard and improved draft, finished genomes, gene expression profiling, resequencing, metagenome projects, single cell projects, transcriptomes, metatranscriptomes, exomes, and others) are publicly available through the Genome Portal.

The Genome Portal also provides access to over 17 000 annotated genomes and metagenomes, along with specialized analytical tools to navigate these data sets and compare genomes.



*Individual portal page*