

# caIntegrator2

## *A translational research tool for 21<sup>st</sup> Century Biomedicine*

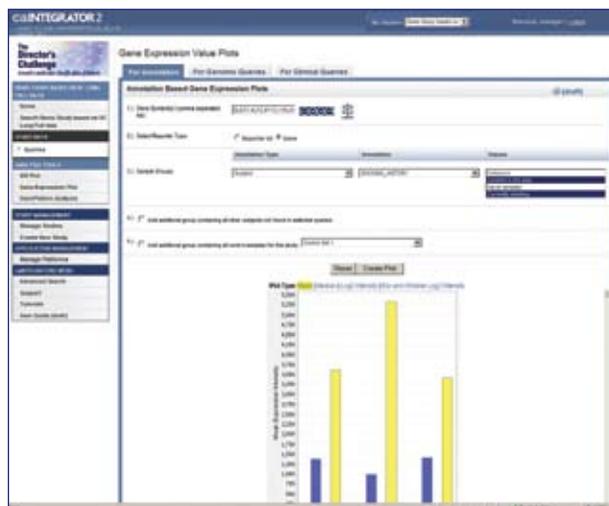
caIntegrator2 is a Web-based software package that allows researchers to set up custom caBIG®-compatible Web portals to conduct integrative research without requiring programming experience. These portals bring together heterogeneous clinical, microarray and medical imaging data to enrich multidisciplinary research.

Using caIntegrator2, researchers can execute, save and share queries to identify and collect many types of data, combining clinical information with genetic and genomic data to enable multidimensional analysis. caIntegrator2 uses caGrid analytical services such as GenePattern and BioConductor to perform analysis on the integrated study data, including clinical survival data.

caIntegrator2 leverages the Cancer Data Standards Registry and Repository (caDSR) to map experimental data to well-defined datatypes and utilizes caGrid and Java client APIs to access data from caBIG® applications such as caArray and the National Biomedical Imaging Archive (NBIA). caIntegrator is also integrated with cancer Biomedical Infrastructure Objects (caBIO) to perform queries on genes and pathways.

### Capabilities

- Web-based deployment and study management
- Remotely access multiple caGrid-accessible datasets including integration with caArray and NBIA
- Map study annotations to caDSR datatypes
- Run complex queries across clinical, microarray and imaging datasets
- Create Kaplan-Meier survival curves based on clinical or gene expression data
- Leverage existing grid capabilities for data analysis through integration with GenePattern and BioConductor



caIntegrator2 interface

### Categories of Use

- |  |  |  |   |
|--|--|--|---|
| <input type="checkbox"/> Biospecimens  | <input type="checkbox"/> Data Sharing                        | <input checked="" type="checkbox"/> <b>Imaging</b>     | <input type="checkbox"/> Proteomics                               |
| <input type="checkbox"/> Clinical Trials Management                              | <input checked="" type="checkbox"/> <b>Genome Annotation</b> | <input checked="" type="checkbox"/> <b>Microarrays</b> | <input checked="" type="checkbox"/> <b>Translational Research</b> |
| <input checked="" type="checkbox"/> <b>Data Analysis &amp; Statistical Tools</b> | <input type="checkbox"/> Infrastructure                      | <input checked="" type="checkbox"/> <b>Pathways</b>    | <input type="checkbox"/> Vocabularies                             |



## Architecture Overview

- **Application type:** Open-source, Web-based J2EE Java application employing Spring, Struts 2 and Hibernate technologies.
- **System requirements:** Full installation requires installation of JBoss; MySQL; ANT (Apache); JDK.

## Installation and Administration

- **Skill sets needed:** Linux/Windows server administrator, MySQL database administrator, software engineer with experience in Java desirable.
- **Infrastructure needed:** Windows/Red Hat Enterprise Linux-based server (32- or 64-bit).
- **Long-term administration needs:** calIntegrator2 requires ongoing system and database administration such as disk space, database management and database and file system backup. Application maintenance is required for user account management.

## Key Contributors

- NCI Center for Biomedical Informatics and Information Technology (NCI CBIIT)
- 5AM Solutions
- Science Applications International Corporation (SAIC)
- ScenPro

## Resources

<b>Tool Overview Page</b>	<a href="https://cabig.nci.nih.gov/tools/calIntegrator">https://cabig.nci.nih.gov/tools/calIntegrator</a>
<b>Primary Workspace</b>	Integrative Cancer Research (ICR) <a href="https://list.nih.gov/cgi-bin/wa?SUBED1=cabig_ICR-I&amp;A=1">https://list.nih.gov/cgi-bin/wa?SUBED1=cabig_ICR-I&amp;A=1</a>
<b>Molecular Analysis Tools Knowledge Center</b>	<a href="https://cabig-kc.nci.nih.gov/Molecular/KC/index.php/Main_Page">https://cabig-kc.nci.nih.gov/Molecular/KC/index.php/Main_Page</a>
<b>caBIG<sup>®</sup> Tool Inventory</b>	<a href="https://cabig.nci.nih.gov/inventory">https://cabig.nci.nih.gov/inventory</a>
<b>NCI Center for Bioinformatics Application Support</b>	<a href="mailto:ncicb@pop.nci.nih.gov">ncicb@pop.nci.nih.gov</a>

