InterMine

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Open source data warehouse and query interface

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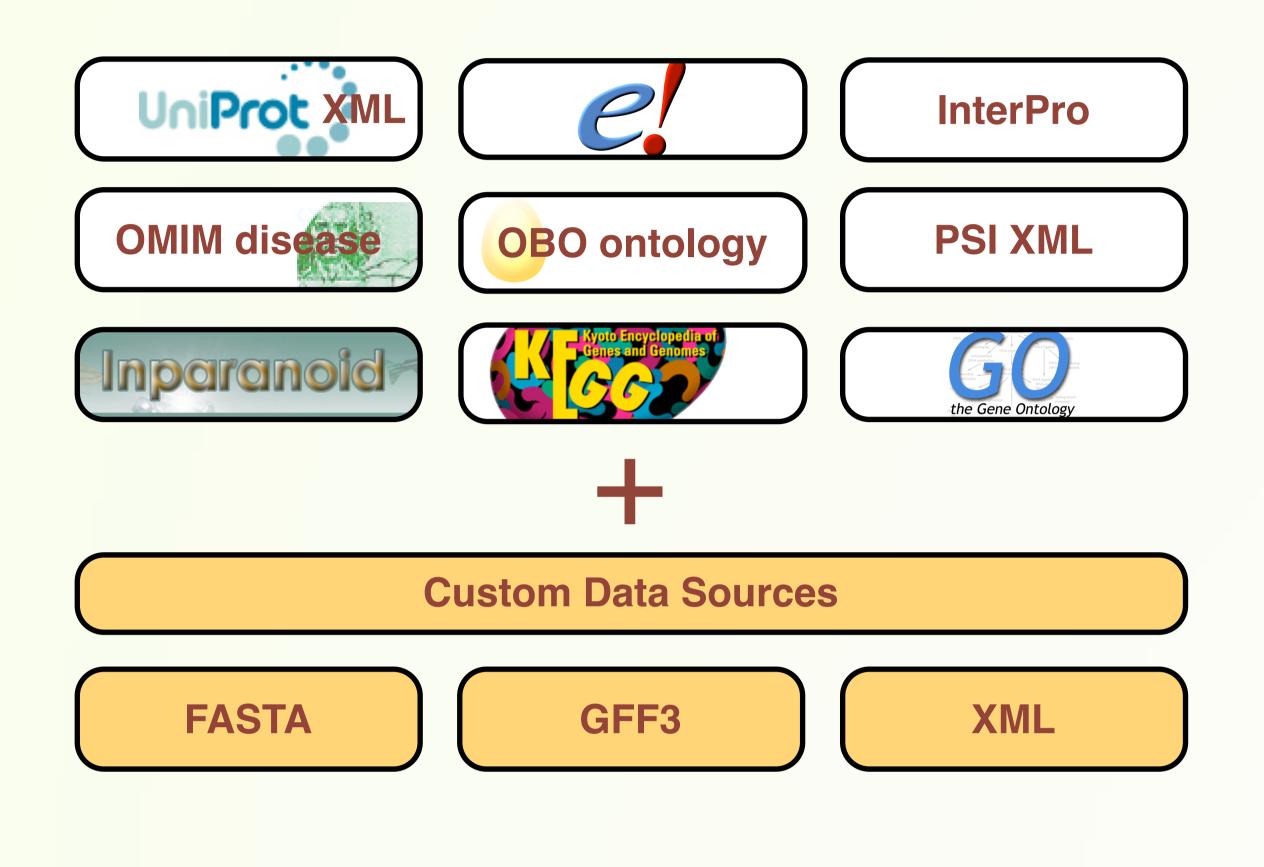
Introduction

InterMine (www.intermine.org) is an open source system for building query-optimised data warehouses. It supports data integration from standard biological formats and makes it easy to add your own data. A sophisticated web application provides flexible query access for any data model. Queries can also be built with an API and executed via web services.

InterMine was developed to enable FlyMine (www.flymine.org) and is now used in several other projects worldwide, including the \$57m modENCODE project (www.modencode.org).

InterMine is written in Java, all code is freely available under the LGPL license.

Software Overview InterMine is used in several projects worldwide. modENCODE **FlyMine** MitoMiner other Drosophila/ Drosophila/ *Mines mitochondria Anopheles C. elegans other data models biology specific genomic/proteomic data model integrate biological data (e.g. UniProt, GFF3) web application configuration, tools, widgets InterMine Java object-based data warehouse configurable data integration



Configurable data integration



InterMine Data Warehouse

flexible web interface, query API

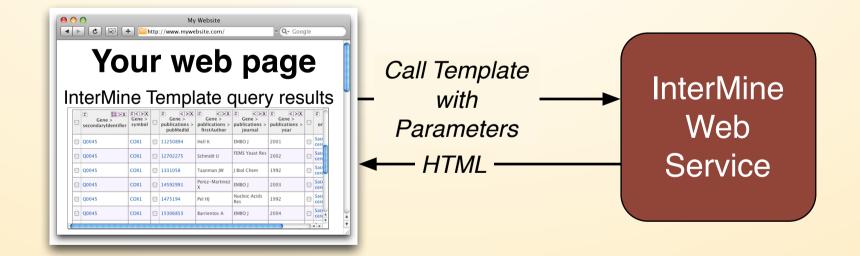
InterMine makes it easy to integrate multiple data sources into a central data warehouse. It has a core data model based on the sequence ontology and supports many biological data formats. The object-based data model is defined as XML from which a database schema and Java classes are automatically generated. It is simple to add custom sources to extend the data model and integrate your own data. Java and Perl APIs are provided for converting data.

Supported formats include GFF3, FASTA, Chado, GO gene association, UniProt XML, PSI XML, PDB XML and Ensembl. A central configuration file defines the sources to load into an InterMine instance and specific organisms and data files to include.

Web Services

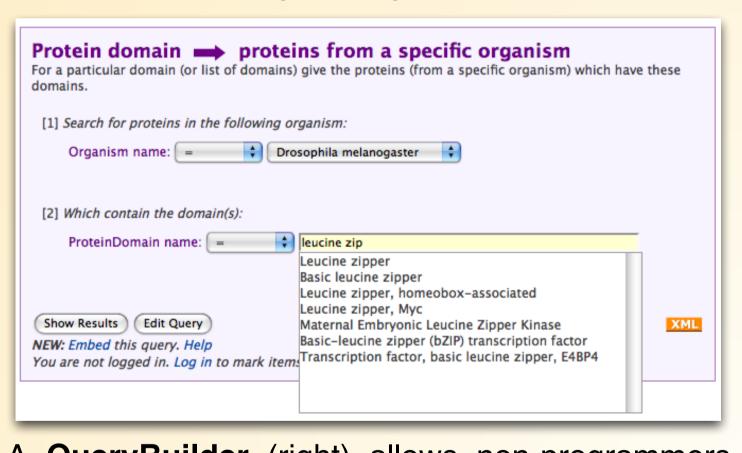
Custom queries, template queries and lists are available via RESTful web services. Results can be returned as XML, tab or comma separated formats. Queries can exported as XML from the web application for use with the web service or created using client APIs in Java or Perl (Summer 2008).

It is also possible to fetch HTML to include query or widget results directly in other web pages. All template queries have an 'embed this query' link which provides a URL to include results in an existing report page, this allows external sites to present data without the need for local data integration.

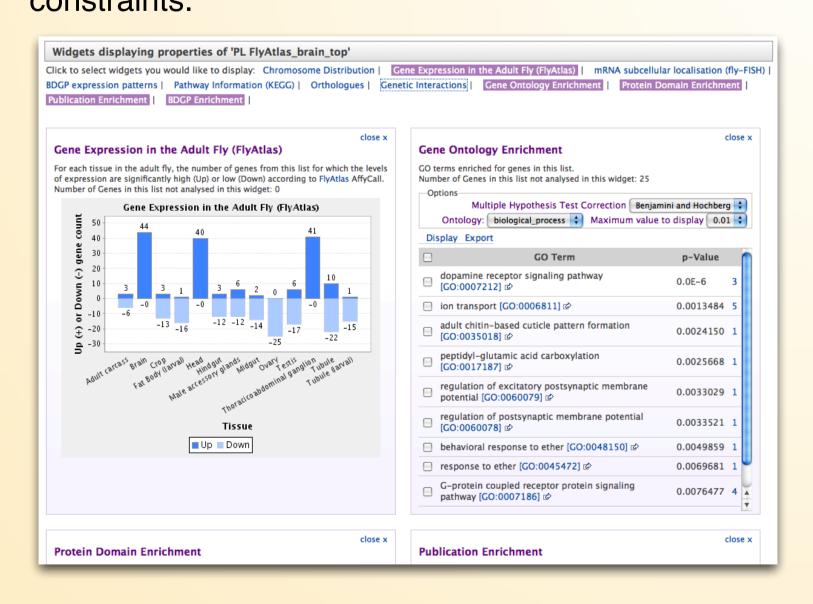


Web Application

A web application works 'out of the box' for any data model to provide flexible query access to the data warehouse, designed for functionality beyond looking up an identifier and viewing a report. The application is highly customisable through configuration and development of new data displayers or integration of tools.



A QueryBuilder (right) allows non-programmers to create custom queries. Any query can be turned into a re-usable template query (above) providing a simple form with a description and editable constraints.



Query summary Model browser Constraint editor Order output columns Set sort order

Lists of any type (e.g. genes, proteins) can be uploaded or created from query results. Lists can be used in any query or template in place of a single identifier. A list analysis page (left) summarises properties of a list through interactive widgets, for example graphs of gene expression or statistical enrichment of GO terms or protein domains. A framework is provided for creating new widgets.

A MyMine account allows users to save lists and queries between sessions.

An important feature is that much of the presentation can be controlled by non-programmers: an admin user can create and publish new template queries and lists at any time and can adapt report pages by applying tags.

Architecture

InterMine is written in Java and built around a custom object/relational mapping tool optimised for read-only performance. The data model is object-based and defined by an XML file from which a database schema and Java classes are automatically generated.

A cost-based query optimiser attempts to rewrite incoming queries to make use of precomputed tables to join fewer tables and run faster. New pre-computed tables can be added at any time to adapt to actual usage. In contrast to traditional data warehouses performance optimisation is thus separated from schema design.

