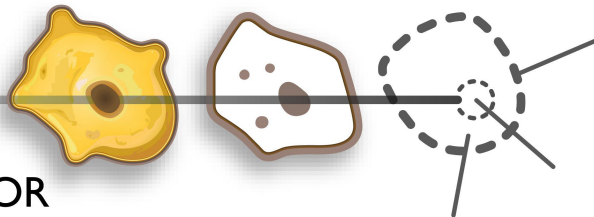


If we build it, will they come?

Social engineering of new technology to
disseminate biomedical ontologies

Mark A. Musen and the BioPortal Team
Stanford University



NATIONAL CENTER FOR

BIOMEDICAL ONTOLOGY

Thanks to a ton of people!

- Benjamin Dai
- Misha Dorf
- Nick Griffith
- Suzanna Lewis
- Dilvan Moreira
- Michael Montegut
- Chris Mungall
- Natasha Noy
- Kaustubh Supekar
- Nicole Washington
- Daniel Rubin
- Nigam Shah

A Small Portion of ICD9-CM

- 724 Unspecified disorders of the back
- 724.0 Spinal stenosis, other than cervical
- 724.00 Spinal stenosis, unspecified region
- 724.01 Spinal stenosis, thoracic region
- 724.02 Spinal stenosis, lumbar region
- 724.09 Spinal stenosis, other
- 724.1 Pain in thoracic spine
- 724.2 Lumbago
- 724.3 Sciatica
- 724.4 Thoracic or lumbosacral neuritis
- 724.5 Backache, unspecified
- 724.6 Disorders of sacrum
- 724.7 Disorders of coccyx
- 724.70 Unspecified disorder of coccyx
- 724.71 Hypermobility of coccyx
- 724.71 Coccygodynia
- 724.8 Other symptoms referable to back
- 724.9 Other unspecified back disorders

The NCI Thesaurus in Protégé-OWL

The screenshot displays the Protégé 3.0 beta interface for editing an OWL class. The main window is titled "Thesaurus Protégé 3.0 beta (file:K:\projects\owl\Thesaurus.pprj, OWL Files)". The interface is divided into several panes:

- Subclass Relationship:** Shows the asserted hierarchy for the project "Thesaurus". The class "Benign_Conditions_of_the_Mouse_Intestinal_Tract" is highlighted in the hierarchy under "Mouse_Cancer-Related_Conditions".
- Class Editor:** The selected class is "Benign_Conditions_of_the_Mouse_Intestinal_Tract". It shows the class name, an "rdfs:comment" field, and a list of annotations.
- Annotations:** A table listing properties and their values for the class.
- Properties and Restrictions:** A list of properties and restrictions associated with the class, including "rEO_Disease_Has_Associated_EO_Anatomy" and "rEO_Disease_Has_Associated_Cell_Type".
- Superclasses:** A list of superclasses for the class, including "Mouse_Noncancerous_Conditions" and "Mouse_Digestive_System_Disorder".
- Disjoints:** A section for defining disjoint classes, currently empty.

Annotations Table:

Property	Value	Lang
code	C22102	
DesignNote	Autonomous new grov...	
Display_Name	Benign Conditions of th...	
FULL_SYN	<term-name>Benign Co...	
FULL_SYN	<term-name>Benign Co...	
hasType	primitive	
Preferred_Name	Benign Conditions of th...	

Properties and Restrictions:

- rEO_Disease_Has_Associated_EO_Anatomy (someValuesFrom Gastrointestinal_Tract_MMHCC, someValuesFrom Gastrointestinal_Tract_MMHCC, someValuesFrom Digestive_System_MMHCC [from Mouse_Digestive_System_Disorder])
- rEO_Disease_Has_Associated_Cell_Type
- rEO_Disease_Has_Property_or_Attribute
- rEO_Disease_Maps_to_Human_Disease

Superclasses:

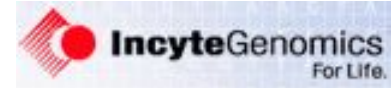
- Mouse_Noncancerous_Conditions
- Mouse_Digestive_System_Disorder

Disjoints:

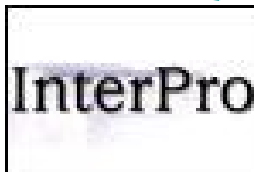
Logic View Properties View

WormBase

FlyBase



UniProt
the universal protein resource



[GO:0003673 : Gene Ontology \(92932\)](#)

[GO:0008150 : biological process \(56952\)](#)

[GO:0007610 : behavior \(566\)](#)

- [GO:0000004 : biological process unknown \(6152\)](#)

[GO:0007154 : cell communication \(11916\)](#)

[GO:0007155 : cell adhesion \(830\)](#)

- [GO:0030260 : cell invasion \(0\)](#)

[GO:0008037 : cell recognition \(210\)](#)

[GO:0007267 : cell-cell signaling \(1318\)](#)

- [GO:0045168 : cell-cell signaling involved in cell fate commitment \(0\)](#)

[GO:0030072 : peptide hormone secretion \(6\)](#)

- [GO:0030252 : growth hormone secretion \(2\)](#)
- [GO:0030073 : insulin secretion \(4\)](#)
- [GO:0030103 : vasopressin secretion \(2\)](#)

[GO:0019226 : transmission of nerve impulse \(688\)](#)

[GO:0030383 : host-pathogen interaction \(12\)](#)

####

Goals of Biomedical Ontologies

- To provide a classification of biomedical entities
- To annotate data to enable summarization and comparison across databases
- To provide for semantic data integration
- To drive NLP systems
- To simplify the engineering of complex software systems
- To provide a formal specification of biomedical knowledge

Open Biomedical Ontologies library

Domain	Prefix	Ontology	Defs file
Arabidopsis gross anatomy	TAIR	arabidopsis anatomy.ontology	arabidopsis anatomy.definitions
Arabidopsis development	TAIR	arabidopsis development.ontology	arabidopsis development.definitions
Cell type	CL	cell.obo	included in cell.obo
Cereal plant gross anatomy	GRO	anatomy gr ont	anatomy gr def
Cereal plant development	GRO	temporal gr ont	temporal gr def
Cereal plant trait ontology	TO	trait ontology	trait definitions
Chemical entities of biological interest	CHEBI	ontology.obo	included in ontology.obo
Protein covalent bond	CV	[none]	[none]
Protein-protein Interaction	MI	psi-mi.dag	psi-mi.def
Maize gross anatomy	ZEA	Zea mays anatomy ontology.txt	Zea mays anatomy ontology definitions.txt
Dictyostelium anatomy	DDANAT	anatomy.ontology	anatomy.definitions
Drosophila gross anatomy	FBbt	fly anatomy.ontology	fly anatomy.definitions
Habronattus courtship		protege source	included in protege source
Loggerhead nesting		protege source	included in protege source
Human anatomy and development	EV	ontologies	[none]
Microarray experimental conditions		MGEDOntology.daml	included in MGEDOntology.daml
Physical-chemical methods and properties	FIX	fix.ontology	[none]
Fungal gross anatomy	FAO	fungal anatomy.ontology	fungal anatomy.definitions
Molecular function	GO	gene_ontology.obo	included in gene_ontology.obo
Biological process	GO	gene_ontology.obo	included in gene_ontology.obo
Cellular component	GO	gene_ontology.obo	included in gene_ontology.obo



OLS

OLS - Ontology Lookup Service

Enter Ontology Term

Search Ontology:

Browse

Term Name: (Include obsolete terms)

Term ID:

Additional Information:

Enter a partial search term. As you are typing, you will see suggested terms that match what are entering in the form. If you select one from the pull-down list, its corresponding ID will be displayed in the form. If you see "... and more" in the list of suggested values, you can select this value to be redirected to a page where all possible values are listed. As an example, enter *mitoc* in the Term Name box while the *Gene Ontology* ontology is selected.

For better search results, do not type punctuation or symbols. For example, if you are looking for 4'-(L-tryptophan), try typing *4 L trypt*.

You can browse an ontology by clicking on the "browse" button next to the ontology selector. To view the complete ontology, do not select a term name. If a term name has been selected, it will be the root from which the ontology will be browsed.

Simple Term ID Search:

Term ID:

Search

Enter a complete term ID (example: GO:0008150) and click on the 'Search' button to quickly obtain all pertinent information for this term. Searches are case-sensitive, so ensure that the proper ontology prefix is used (GO:, rather than go: or Go:).

- OLS Home
- Documentation
 - Project
 - Publications
- Developer Resources
 - Download
 - Implementation Overview
 - Javadoc
 - Webservice documentation
- Contact Us
 - Acknowledgements

News



June 2007: Maintenance Release

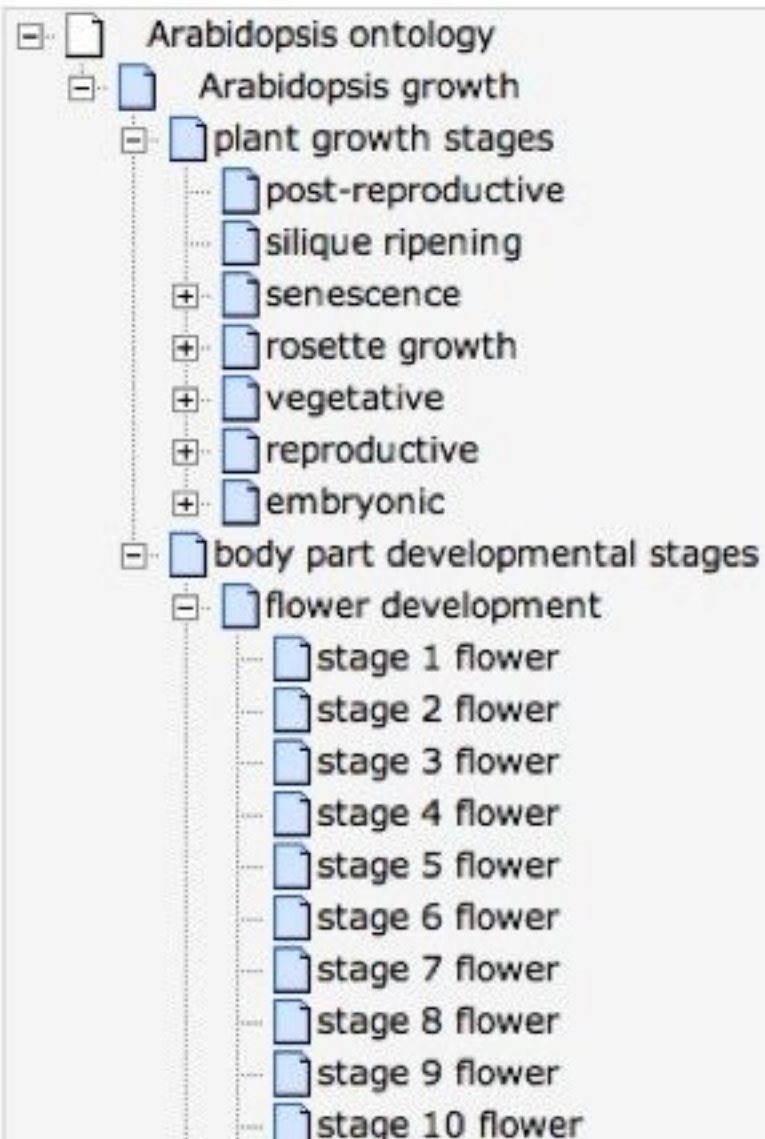
Implementation documentation has been updated to include more recent dependencies. Please note that unless stated otherwise in specific instances, newer versions of given dependencies should work without issue.



OLS - Ontology Lookup Service

TAIR Ontology Browser

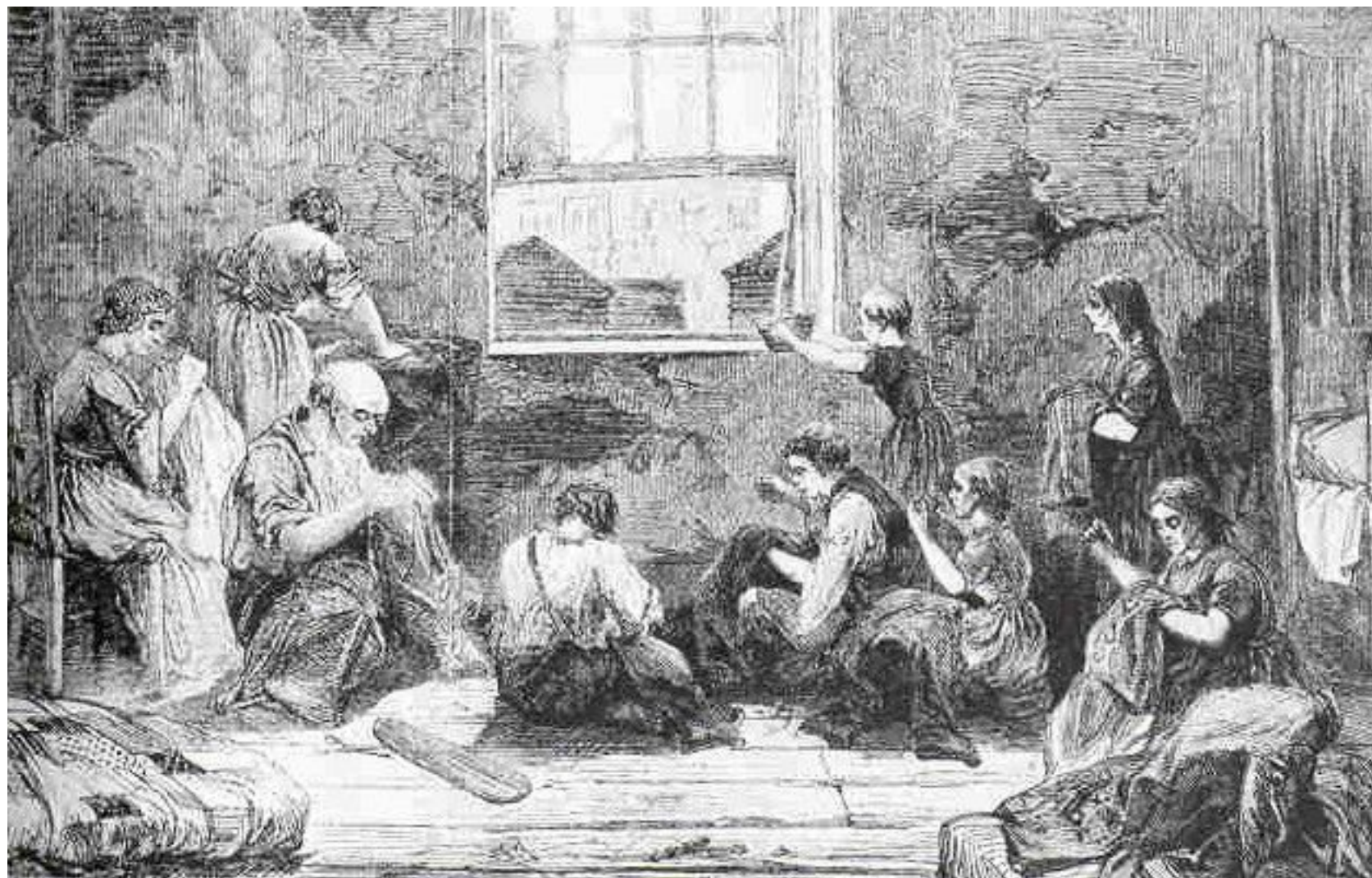
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- [Documentation](#)
 - Project
 - Publications
- [Developer Resources](#)
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 - Javadoc
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News

June 2007: Maintenance Release

Implementation documentation has been updated to include more recent dependencies. Please note that unless stated otherwise in specific instances, newer versions of dependencies should



In biology, lots of ontology developers are almost hobbyists

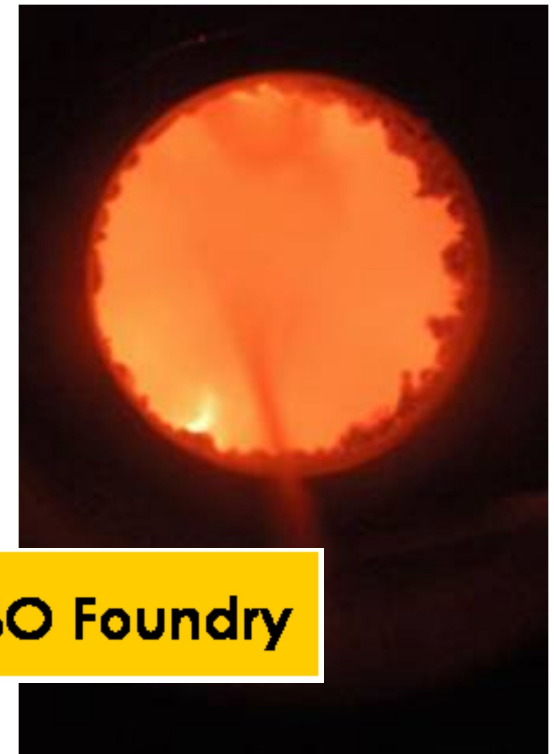
- Nearly always, ontologies are created to address pressing practical needs
- The people who have the most insight into professional knowledge of a given biomedical domain may have little appreciation for metaphysics, principles of knowledge representation, or computational logic
- There simply aren't enough good ontologists to go around

Issues in assuring ontology quality

- Unlike the case with journal submissions, it makes no sense for ontologies to be peer-reviewed by just a handful of experts
- Open, community-based review of ontologies may be haphazard and chaotic
- Top-down solutions may offer rigid review criteria at the expense of scalability
- There is a pressing need for empirical evaluation of methods for ontology evaluation

A Curated Approach for Quality Assurance

- A proposal to create a family of interoperable “gold standard” biomedical reference ontologies
- Formulated by Barry Smith and members of the GO Consortium
- *A Good Housekeeping* Seal of Approval for biomedical ontologies



The OBO Foundry

[Home](#)[Principles](#)[Ontology Table](#)[Ontology Index](#)[Mappings](#)[Project](#)[Repository](#)[Downloads](#)[Subscribe](#)[Contact](#)[Tools](#)

Please submit any updates or corrections to the **OBO** webmaster

Hosted by



For an ontology to be accepted as one of the Open Biomedical Ontologies in the Foundry, the following criteria must be met (further principles will be added over time):

Version as of 24 April 2006

- 1. The ontology must be *open* and available to be used by all without any constraint other than (a) its origin must be acknowledged and (b) it is not to be altered and subsequently redistributed under the original name or with the same identifiers.**

The OBO ontologies are for sharing and are resources for the entire community. For this reason, they must be available to all without any constraint or license on their use or redistribution. However, it is proper that their original source is always credited and that after any external alterations, they must never be redistributed under the same name or with the same identifiers.
- 2. The ontology is in, or can be expressed in, a *common shared syntax*. This may be either the OBO syntax, extensions of this syntax, or OWL.**

The reason for this is that the same tools can then be usefully applied. This facilitates shared software implementations. This criterion is not met in all of the ontologies currently listed, but we are working with the ontology developers to have them available in a common OBO syntax.
- 3. The ontologies possesses a *unique identifier space* within the OBO Foundry.**

The source of concepts from any ontology can be immediately identified by the prefix of the identifier of each concept. It is, therefore, important that this prefix be unique.
- 4. The ontology provider has procedures for identifying distinct successive *versions*.**
- 5. The ontology has a clearly specified and clearly *delineated content*.**

The ontology must be orthogonal to other ontologies already lodged within OBO.

The major reason for this principle is to allow two different ontologies, for example anatomy and process, to be combined through additional relationships. These relationships could then be used to constrain when terms could be jointly applied to describe complementary (but distinguishable) perspectives on the same biological or medical entity.

As a corollary to this, we would strive for community acceptance of a single ontology for one domain, rather than encouraging rivalry between ontologies.
- 6. The ontologies include textual *definitions* for all terms.**

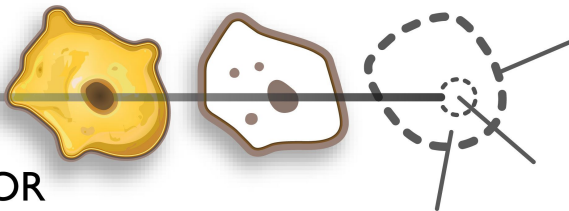
Many biological and medical terms may be ambiguous, so concepts should be defined so that their precise meaning within the context of a particular ontology is clear to a human reader.
- 7. The ontology uses relations which are unambiguously defined following the pattern of definitions laid down in the *OBO Relation Ontology*.**
- 8. The ontology is *well documented*.**
- 9. The ontology has a plurality of independent *users*.**
- 10. The ontology will be developed *collaboratively* with other OBO Foundry members.**

OBO Foundry must address lots of questions

- Can the top–down approach scale?
How many ontologies can be managed by a small panel of curators?
- Who gets to reject an ontology on the basis of form or content? What is the appeals process? How do we know whom to believe?
- Who will curate the curators?

The National Center for Biomedical Ontology

- One of three National Centers for Biomedical Computing launched by NIH in 2005
- Collaboration of Stanford, Berkeley, Mayo, Buffalo, Victoria, UCSF, Oregon, and Cambridge
- Primary goal is to make ontologies accessible and usable
- Research will develop technologies for ontology dissemination, indexing, alignment, and peer review





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NIH Roadmap

National Centers for Biomedical Computing

Home

NCBC Summary

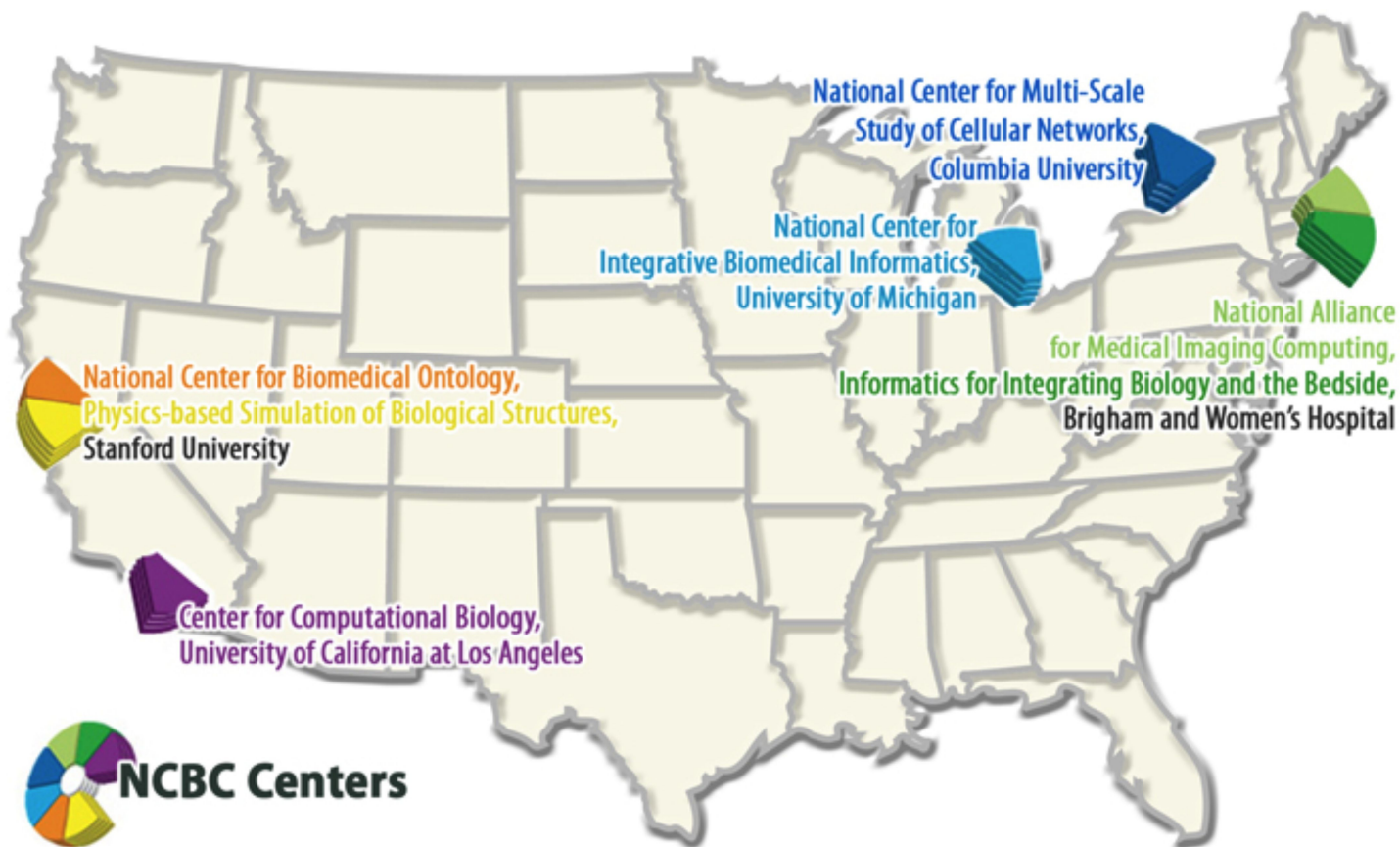
SDIWG

Tools and Applications

Ontology Working Group

DBP Interactions

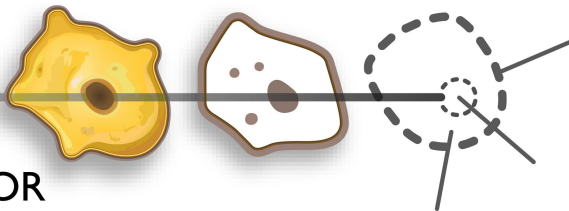
ATTENTION: [Final Report from National Centers for Biomedical Computing 2006 All Hands Meeting](#)



<http://www.ncbcs.org>

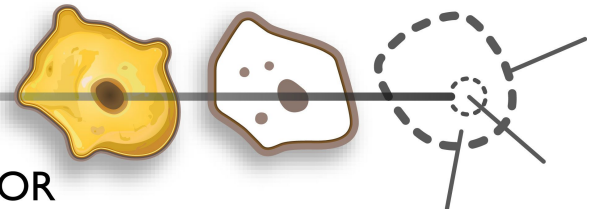
NCBO will offer

- Technology for uploading, browsing, and using biomedical ontologies
- Methods to make the online “publication” of ontologies more like that of journal articles
- Tools to enable the biomedical community to put ontologies to work on a daily basis



Goals for BioPortal

- Web accessible repository of ontologies for the biomedical community
 - Archived locally
 - Anywhere in cyberspace
- Support for ontology
 - Peer review
 - Annotation (marginalia)
 - Versioning
 - Alignment
 - Search



NATIONAL CENTER FOR

BIOMEDICAL ONTOLOGY

http://bioportal.bioontology.org



Version **Beta-RC2**

[Browse](#)

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Ontologies

[List View](#) [Category View](#)

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Name	Format	Current Version	Content Location	Action
Amino Acid	OWL Full	1.1	NCBO Library	  
Animal natural history and life history	Protégé	Unknown	Remote	  
Arabidopsis development	OBO	1.1	NCBO Library	  
Basic -Vertebrate	OWL Full	1.1	NCBO Library	  
Biological imaging methods	OBO	1.1	NCBO Library	  
BRENDA tissue / enzyme source	OBO	1.96	NCBO Library	  
C. elegans development	OBO	1.1	NCBO Library	  
C. elegans gross anatomy	OBO	Unknown	Remote	  
Protein modification	OBO	1.1/4	NCBO Library	  
Protein Ontology	OWL Full	2.0	NCBO Library	  
Protein-protein interaction	OBO	1.68	NCBO Library	  
Proteomics data and process provenance	OWL Full	1.1	NCBO Library	  
RadLex	Protégé	1.1	NCBO Library	  
Sample processing and separation techniques	OBO	Unknown	Remote	  
Sequence types and features	OBO	1.29	NCBO Library	  
Systems Biology	OBO	Unknown	Remote	  
Zebrafish anatomy and development	OBO	1.9	NCBO Library	  

Browsing/Visualizing Ontologies

Zebrafish anatomy and development

Tree View

Tree view constructed based on *is_a* hierarchy

- ⊕ Stages
- ⊖ zebrafish anatomical entity
 - ⊖ anatomical set
 - ⊖ anatomical structure
 - ⊖ acellular anatomical structure
 - ⊕ anatomical cluster
 - ⊕ cardinal organism part
 - ⊖ cell
 - ⊖ dopaminergic neuron
 - ⊖ epidermal cell
 - ⊖ granulocyte
 - ⊕ embryonic structure
 - ⊕ extraembryonic structure
 - ⊕ organ
 - ⊕ organ system
 - ⊖ portion of tissue
 - ⊕ whole organism
 - ⊖ unspecified

Class Details

General

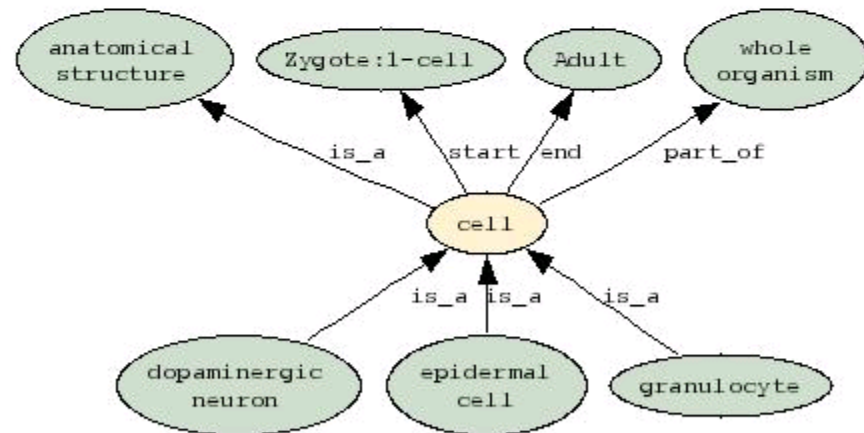
Class Name **cell**
Id **CL:0000000**

Attributes

Database_References **ZFIN:ZDB-ANAT-060816-76**

Graph View

Graph Type



Local Neighborhood view

Zebrafish anatomy and development

Tree View

Tree view constructed based on *is_a* hierarchy

- ⊕ Stages
- ⊖ zebrafish anatomical entity
 - ⊕ anatomical set
 - ⊖ anatomical structure
 - acellulare anatomical structure
 - ⊕ anatomical cluster
 - ⊕ cardinal organism part
 - ⊖ cell
 - dopaminergic neuron
 - epidermal cell
 - granulocyte
 - ⊕ embryonic structure
 - ⊕ extraembryonic structure
 - ⊕ organ
 - ⊕ organ system
 - portion of tissue
 - ⊕ whole organism
 - unspecified

Class Details

General

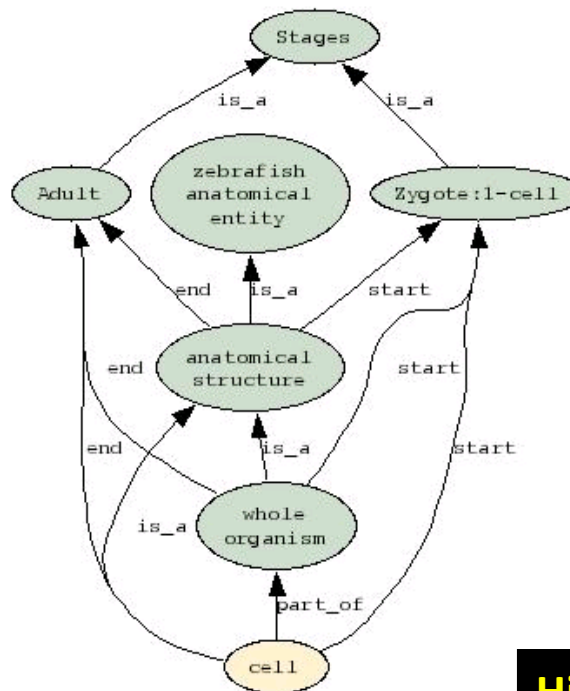
Class Name **cell**
 Id **CL:0000000**

Attributes

Database_References **ZFIN:ZDB-ANAT-060816-76**

Graph View

Graph Type ▾



Hierarchy-to-root view

[Browse](#)

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[Ontology Content](#) | [Ontology Metadata](#)

Search Ontology Content

Search for ontology content within a specific ontology, several ontologies or across all ontologies in the library.

* Ontology

* Search In Class Name or ID Attributes

* Search Type Contains Sounds Like

Include Obsolete Classes

* Search Text

Search text must be at least 3 characters in length.

Search Results

Ontology [Protein-protein interaction](#)

[Class Name \(11\)](#)

[Attributes \(83\)](#)

Class Name	Id	Attributes
3d repertoire	MI:0731	Definition: The aim of 3D Repertoire is to determine the structures of all amenable complexes in a cell at medium or toponomic and dynamic analyses of protein complexes in a cell . Complex models, EM pictures, expression and pur a database connected to the PDB repository. RELATED SYNONYM: "3D Repertoire"
agonist	MI:0625	Definition: Description of an activator that acts on an external cell receptor or other upstream molecule to stimulate or more of the interactors.
alliance for cellulare signaling	MI:0575	Definition: Alliance for Cell ular Signaling (AfCS -Nature) store yeast 2-hybrid Interaction data and expression data. Ir all.nhttp://www.signaling-gateway.org EXACT SYNONYM: "afcs" Database_References: search-url: "http://www.signaling-gateway.org/data/Y2H/cgi-bin/y2h_int.cgi?id=\${ac}", id-valid RELATED SYNONYM: "AfCS"
nucleic acid conjugation	MI:0715	Definition: Bacterial conjugation is the transfer of genetic material between bacteria through cell -to- cell contact. Bac bacterial equivalent of sexual reproduction or mating. It is not actually sexual, as it does not involve the fusing of gan a conjugative plasmid from a donor cell to a recipient EXACT SYNONYM: "nucl conjugation"

BioPortal's impact in the community

- National Cancer Institute
 - Deploying BioPortal locally to evaluate its use as a method for visualizing and navigating enterprise terminologies and ontologies
- Biomedical Informatics Research Network (BIRN)
 - Adopting BioPortal for disseminating and visualizing BIRN Lex terminology
- Radiological Society of North America
 - Using BioPortal for graphical visualization of RadLex

BioPortal will allow NCBO to experiment with new models for

- Dissemination of knowledge on the Web
- Integration and alignment of online content
- Knowledge visualization and cognitive support
- Peer review of online content

The NCI Thesaurus in Protégé-OWL

Thesaurus Protégé 3.0 beta (file:K:\projects\owl\Thesaurus.pprj, OWL Files)

File Edit Project OWL Code Window Help

protégé

OWLClasses Properties Forms Individuals Metadata

SUBCLASS RELATIONSHIP

For Project: Thesaurus

Asserted Hierarchy

- owl:Thing
 - Abnormal_Cell_Kind
 - Anatomy_Kind
 - Biological_Process_Kind
 - Chemicals_and_Drugs_Kind
 - Chemotherapy_Regimen_Kind
 - Clinical_or_Research_Activity_Kind
 - Diagnostic_and_Prognostic_Factors_Kind
 - Drug_Mechanism_of_Action_Kind
 - Drug_Physiologic_Effect_Kind
 - EO_Anatomy_Kind
 - EO_Findings_and_Disorders_Kind
 - Experimental_Organism_Diagnoses
 - Experimental_Allergic_Encephalomyelitis
 - Mouse_Pathologic_Diagnoses
 - Mouse_Cancer-Related_Conditions
 - Benign_Plasma_Cell_Proliferations_of...
 - Hyperplasia_of_the_Mouse_Intestinal...
 - Hyperplasia_of_the_Mouse_Pulmonar...
 - Melanocytic_Proliferative_Disorders_c...
 - Mouse_Noncancerous_Conditions
 - Benign_Conditions_of_the_Mouse...
 - Congestion_of_the_Mouse_In...

CLASS EDITOR

For Class: Benign_Conditions_of_the_Mouse_Intestinal_Tract (instance of owl:Class)

Name: Benign_Conditions_of_the_Mouse_Intestinal_Tract

SameAs: [] DifferentFrom: []

rdfs:comment: []

Annotations

Property	Value	Lang
code	C22102	
DesignNote	Autonomous new grov...	
Display_Name	Benign Conditions of th...	
FULL_SYN	<term-name>Benign Co...	
FULL_SYN	<term-name>Benign Co...	
hasType	primitive	
Preferred_Name	Benign Conditions of th...	

Properties and Restrictions

- rEO_Disease_Has_Associated_EO_Anatomy (someValuesFrom Gastrointestinal_Tract_MMHCC, someValuesFrom Gastrointestinal_Tract_MMHCC, someValuesFrom Digestive_System_MMHCC [from Mouse_Digestive_System_Disorder])
- rEO_Disease_Has_Associated_Cell_Type
- rEO_Disease_Has_Property_or_Attribute
- rEO_Disease_Maps_to_Human_Disease

Superclasses

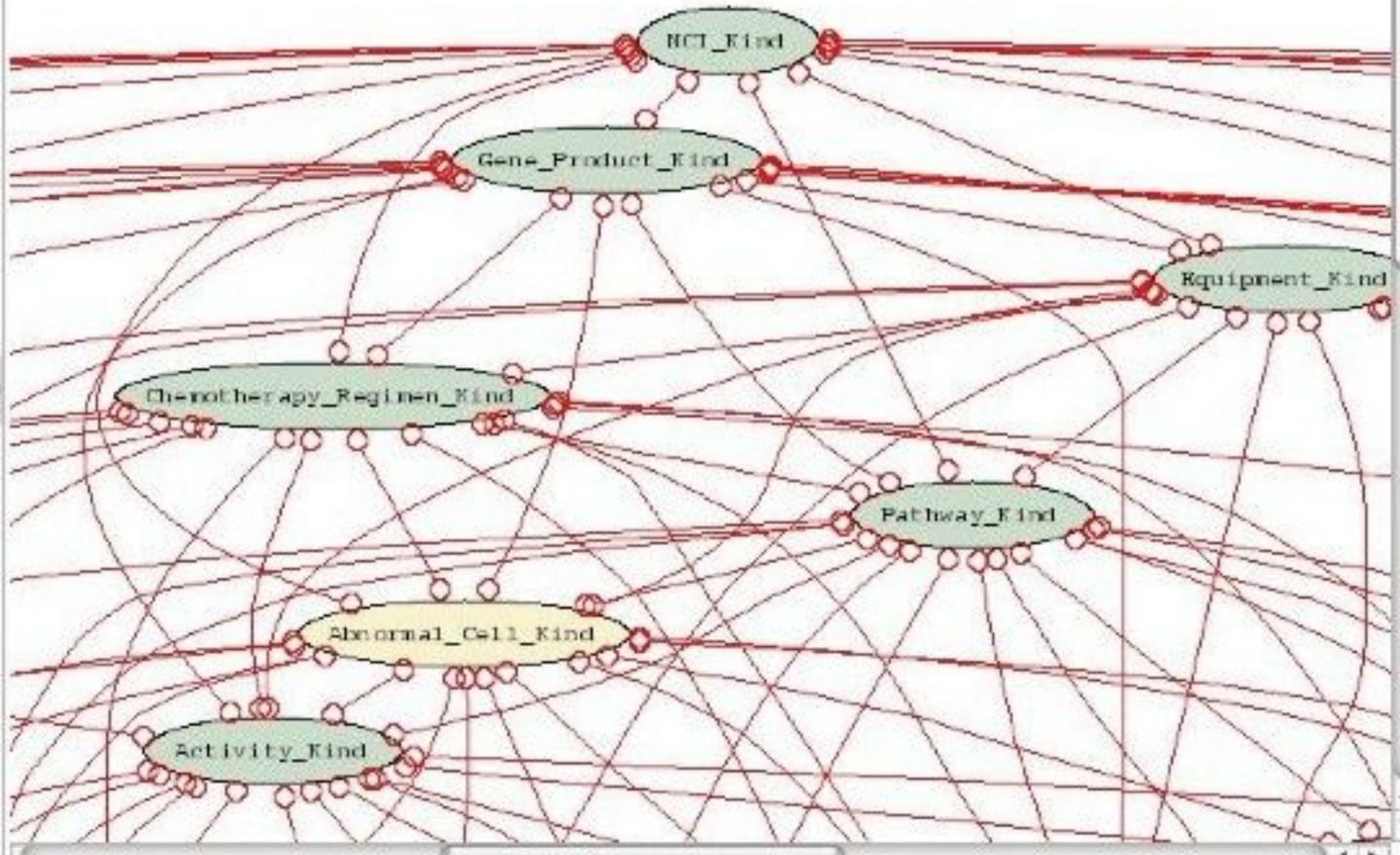
- Mouse_Noncancerous_Conditions
- Mouse_Digestive_System_Disorder

Disjoints

[]

NCI Thesaurus

- + Abnormal_Cell_Kind
- + Activity_Kind
- Anatomy_Kind
 - Anatomic Structure, Syst
 - + Body Fluid or Substan
 - Body Part
 - + Body Region
 - + Body Cavity
 - + Embryologic Structure
 - + Microanatomic Structu
 - + Organ
 - + Organ System
 - + Other Anatomic Conce
- + Biological_Process_Kind
- + Chemicals_and_Drugs_Kind
- + Chemotherapy_Regimen_Kir
- + Diagnostic_and_Prognostic_
- + EO_Anatomy_Kind
- + EO_Findings_and_Disorders
- + Equipment_Kind
- + Findings_and_Disorders_Kin



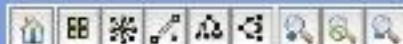
Home Browse Search

- + Abnormal_Cell_Kind
- + Activity_Kind
- + Anatomy_Kind
- + Biological_Process_Kind
- + Chemicals_and_Drugs_Kind
- + Chemotherapy_Regimen_Kind
- + Diagnostic_and_Prognostic_Fa
- + EO_Anatomy_Kind
- + EO_Findings_and_Disorders_K
- + Equipment_Kind
- + Findings_and_Disorders_Kind
- + Gene_Kind
- + Gene_Product_Kind
- + Molecular_Abnormality_Kind
- + NCI_Kind
- + Organism_Kind
- + Pathway_Kind
- + Properties_or_Attributes_Kind
- + Retired_Kind

Visualiz

Show M

NCI

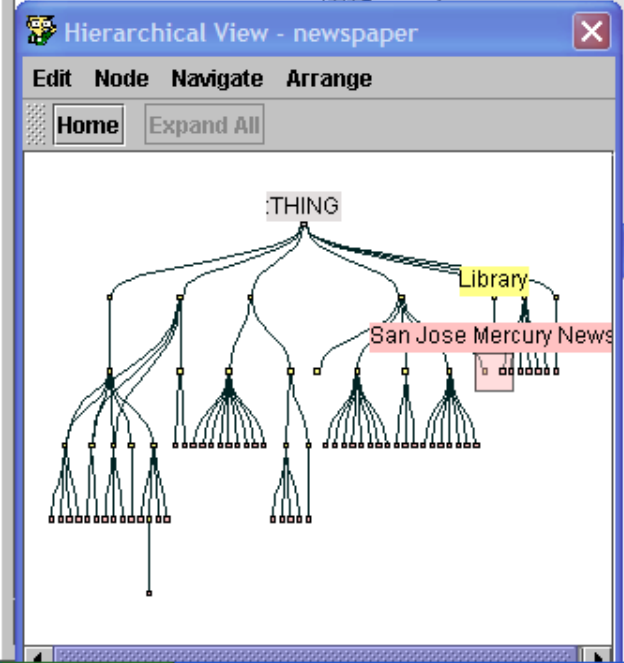
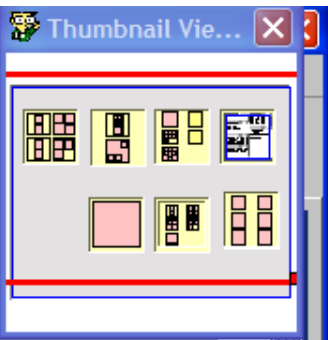
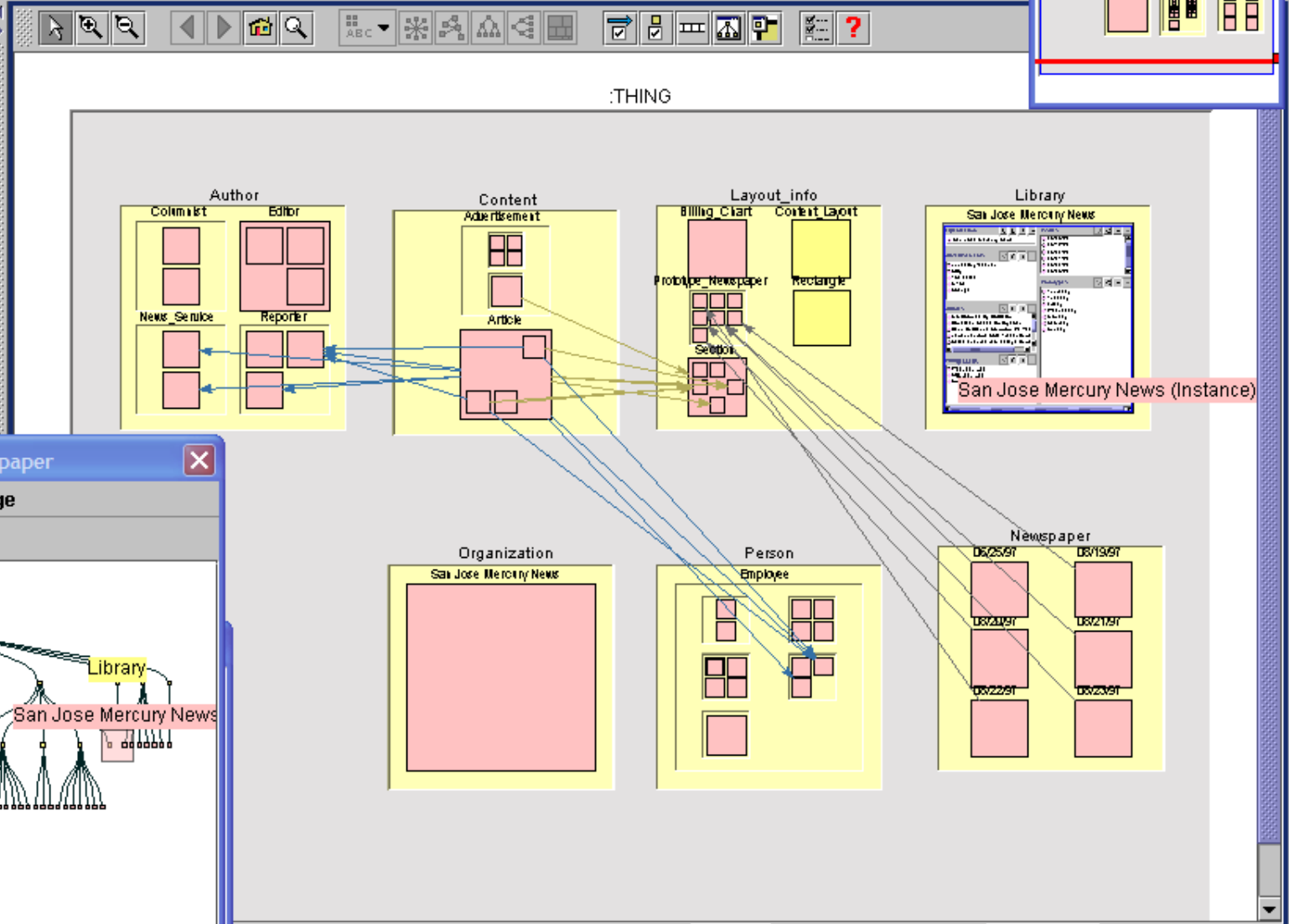


Done

Applet org.ncbo.uvic.shrimp.BioPortalApplet started



- Classes
- .THING A
 - .SYSTEM-CLASS A
 - Author A
 - Content A
 - Layout_info A
 - Library (1)
 - Newspaper (6)
 - Organization (1)
 - Person



Classes

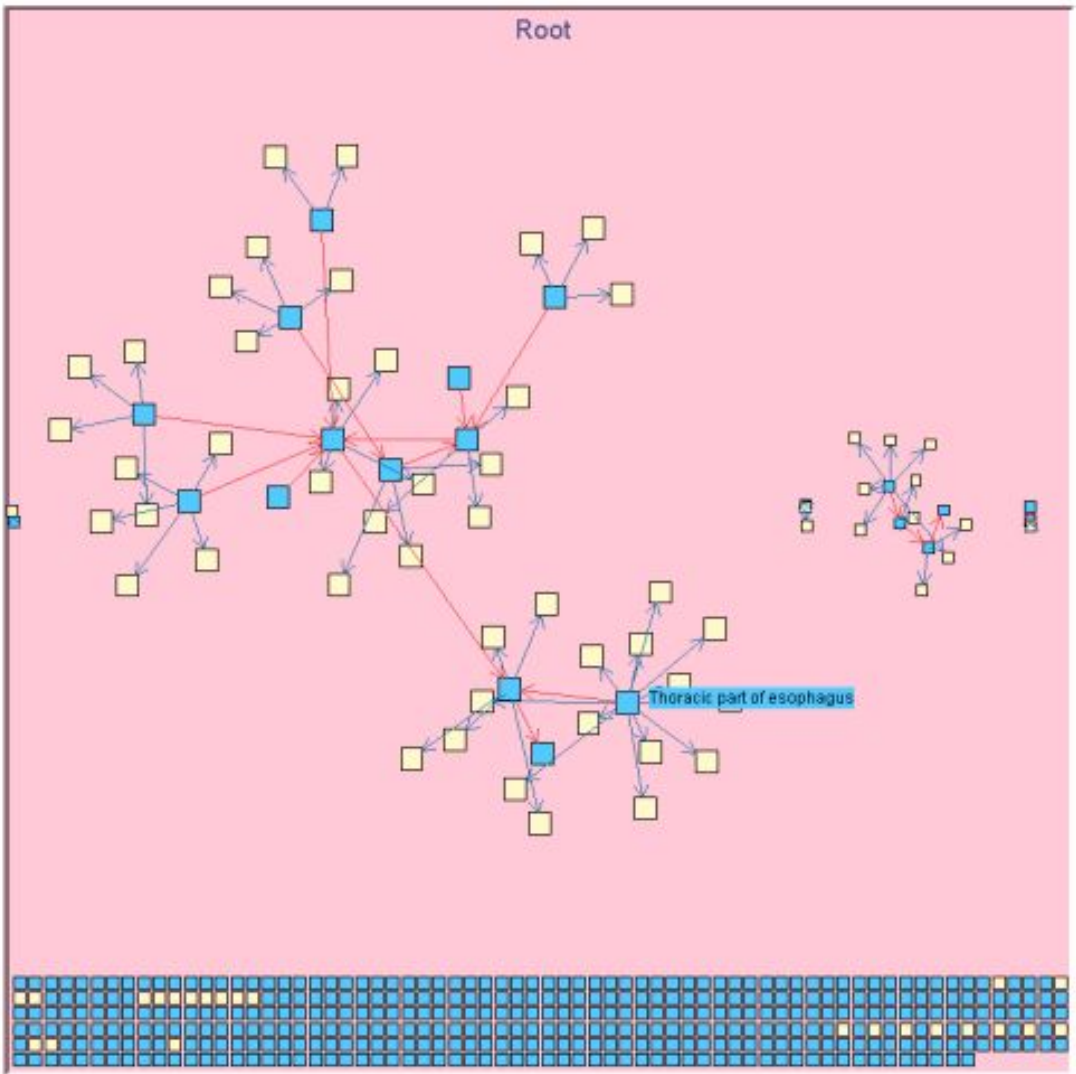
Hierarchy: -- none --

- THING
 - SYSTEM-CLASS
 - Anatomical entity
 - Physical anatom
 - Material phys
 - Non-material
 - Conceptual anal
 - Structural rel
 - Anatomical r
 - Dimensional entity
 - Volume

Arc Type Filter - Shrimp

Display All Hide All

- Has member
- Has morphological part
- Has skin
- inherent 3-D shape
- is boundary of
- is part of
- is-a
- Laterality
- orientation
- part
- part of
- partition
- physical state
- Rank of tissue



Ontologies are not like journal articles

- It is difficult to judge methodological soundness simply by inspection
- We may wish to use an ontology even though some portions
 - Are not well designed
 - Make distinctions that are different from those that we might want

Ontologies are not like journal articles

- The utility of ontologies
 - Depends on the task
 - May be highly subjective
- The expertise and biases of reviewers may vary widely with respect to different portions of an ontology
- Users should want the opinions of more than 2–3 hand-selected reviewers
- Peer review needs to scale to the entire user community

Community-Based Annotation as Peer Review

- Makes ontology evaluation a democratic process
- Assumes users' application of ontologies will lead to insights not achievable by inspection alone
- Assumes end-users will be motivated to comment on and engage in dialog about ontologies in the repository



NCI Thesaurus

- Abnormal_Cell_Kind
- Activity_Kind
- Anatomy_Kind
- Biological_Process_Kind
- Chemicals_and_Drugs_Kind
- Chemotherapy_Regimen_Kind
- Diagnostic_and_Prognostic_Factors_Kind
- Diagnostic_Anatomy_Kind
- Diagnostic_Findings_and_Disorders_Kind
- Equipment_Kind
- Findings_and_Disorders_Kind
- Disease, Disorder or Finding**
- Gene_Kind
- Gene_Product_Kind
- Molecular_Abnormality_Kind
- NCI_Kind
- Organism_Kind
- Pathway_Kind
- Properties_or_Attributes_Kind
- Retired_Kind

Filter Notes

Filter Text:

Filter By Type:

Comment: Class appropriateness Bill Bug at 11/20/07 14:57

Having a disjunction in a class name is an odd thing to have

[Reply](#)

Explanation: Re: Class appropriateness MaryAnn Martone at 11/20/07 20:47

[New Thread](#)

An ontology of “marginal notes”



Classes Slots Forms Instances Queries

CLASS BROWSER

For Project: ● changes_marginalia

Class Hierarchy

- AnnotatableThing
 - Annotation
 - Advice
 - Comment
 - Example
 - Explanation
 - MarginalNote
 - Class_Notes
 - Class_Compositionality
 - Class_Merge
 - Class_Relationship_Type
 - Class_Sibling
 - Class_Split
 - Class_Subclass
 - Class_Superclass
 - ▶ ● Component_Notes
 - ▶ ● MarginalNote_Notes
 - ▶ ● Ontology_Notes
 - ▶ ● Property_Notes

CLASS EDITOR

For Class: ● Class_Sibling (instance of :STANDARD-CLASS)

Name: Class_Sibling

Documentation: Notes about sibling(s) of a class

Constraints

Role: Concrete ●

Template Slots

Name	Cardin...	Type	Other Facets
annotates	multiple	Instance of Ontolo...	inverse-slot=associatedAnnotations
associatedAnno...	multiple	Instance of Annota...	inverse-slot=annotates
author	single	String	
body	single	String	
context	single	String	
created	single	Instance of Timest...	
modified	single	Instance of Timest...	
related	single	String	

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SUBJECTS

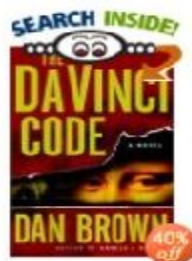
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THE NEW YORK TIMES®
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MAGAZINES

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REAL NAME

There is no question that everybody should read this book. It is very entertaining and full of very peculiar facts (assuming that they are true). The writer skillfully turns religious history (highly sensitive and mostly boring subject to read) into a page turning thriller. I highly recommend it.

I have a general advise though, make sure that you read it on a weekend, as you will not be able to put it down. I read it on a business trip with near disastrous consequences.

Was this review helpful to you? yes no [\(Report this\)](#)★☆☆☆☆ **Don't Take It as Gospel**, November 9, 2003Reviewer: [Leslie Strang Akers](#) (Riverside, CA) - [See all my reviews](#)

In the beginning I was intrigued by the premise set down in THE DA VINCI CODE, but my initial interest turned first to annoyance and then by the time I got to the info on Disney was laughing so hard at the absurdity of the whole novel. First of all, this is a work of fiction, so let's deal with that part. Far from being the taut, fast-paced thriller that the potenti reader is lead to believe it is, TDVC is turgid, jerky, and filled with clichés. The characters are characterless and stupid, merely cardboard for the author to push around like pawn a chessboard. Langford, a Harvard professor, can't distinguish between backwards English and a Semitic language. Sophie, a French police cryptologist, doesn't have the brains to figure out that an armor truck from a Swiss bank might be lo-jacked. These are only two of the many idiotic things the main characters aren't intelligent enough to figure out. The characters ponder clues ad nauseum, which turns a 300-page book into 454 pages. I don't know if the author is writing down to his audience, or if he really thinks that gifted peop are idiot savants. Whatever it is, it's exasperating.

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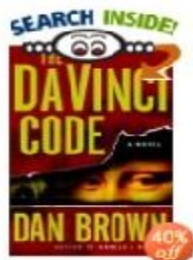
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by [Dan Brown](#) "Robert Langdon awoke slowly..." [\(more\)](#)



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Solution
Snapshot

 **cBIO: National Center for Biomedical Ontology**

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The MGED Ontology

by [EMBL](#) "The primary purpose of the MGED Ontology is to provide standard" ([more](#))

MGED



Search inside
this Ontology

List Price: ~~\$\$~~

Price: ~~\$0~~ & Eligible for **FREE Access**

You Save: ~~\$0~~

Availability: Usually available 24/7 on cBIO.org

Edition: Pragmatic

2 of 3 people found the following review helpful:

★★★★★ **A Great resource**, Aug 11, 2004

Reviewer: [Catherine](#) Ball (Stanford, CA USA) - [See all my reviews](#)

TOP 500 REVIEWER REAL NAME

MGED Ontology aims to facilitate the sharing of microarray data generated by functional genomics and proteomics experiments....

Was this review helpful to you? yes no ([Report this](#))

1 of 1 people found the following review helpful:

☆☆☆☆☆ **Needs considerable improvement**, November 9, 2003

Reviewer: [Barry](#) Smith (Buffalo, NY) - [See all my reviews](#)

MGED ontology is indeed an essential part of any solution to the problems of Microarray analysis - but only if it is understood in the right sort of way. Ontological engineering, should in every case go hand in hand with a sound ontological theory....

Open ratings for ontologies

- Any user can
 - rate an ontology
 - add a “marginal note”
- Ontology evaluation becomes a community-based initiative
- *A web of trust* can enable users to filter comments or ratings to avoid “noise”

Possible Review Criteria

- What is the level of user support?
- What documentation is available?
- What is the granularity of the ontology content in specific areas?
- How well does the ontology cover a particular domain?
- In what applications has the ontology been used successfully? Where has it failed?

Users can make proposals for changes

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Proposals for Mutation Type Core:Editorial Note

Current Value

As with the more specific types of isogenic status listed as TaxonRank instances, there is an implied subsumptive graph amongst the entries in this list that can be represented as a hierarchy when the final representation technique is resolved for this information. For example, Transgenic_insertion is a type of Insertion, Knock-out_targeted_mutatation is a type of Targeted_mutation, etc. The mutation type instances are a modified version of the collection of mutation types employed by the IMSR. The primary difference is additional, distinct types have been specified, when an IMSR definition included multiple types in its definition.

Proposal: Proposal For Change

Jon Doe at 11/08/07 14:46

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[Reply](#)

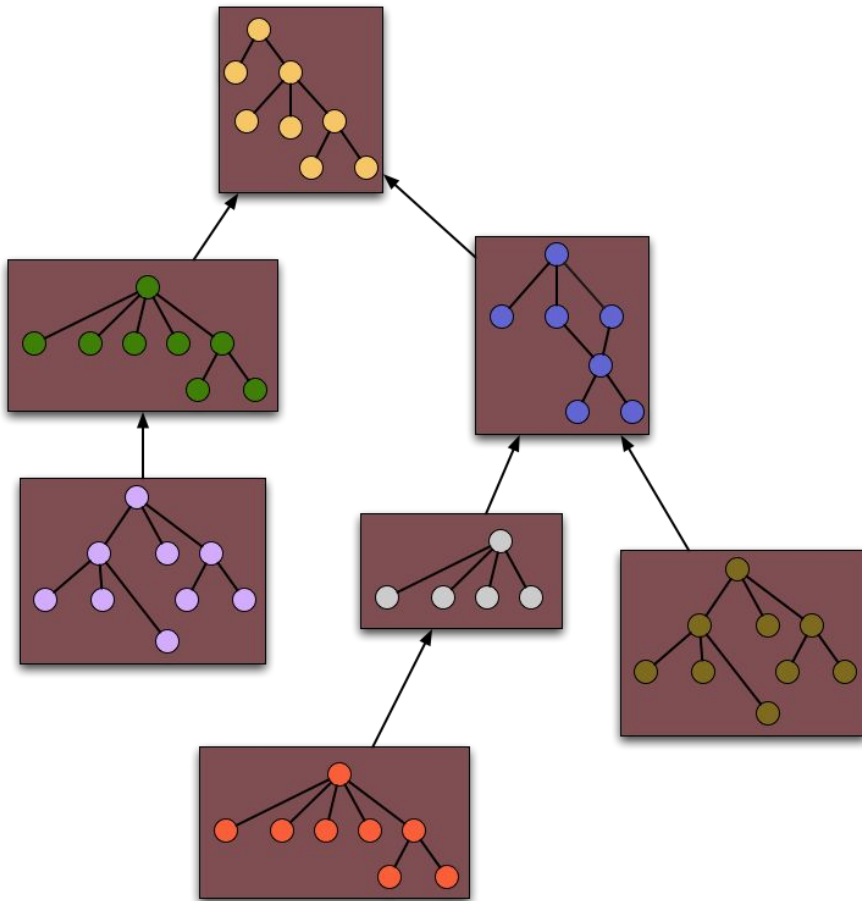
Explanation: Proposal For Change

Jon Doe at 11/08/07 14:46

New Proposal

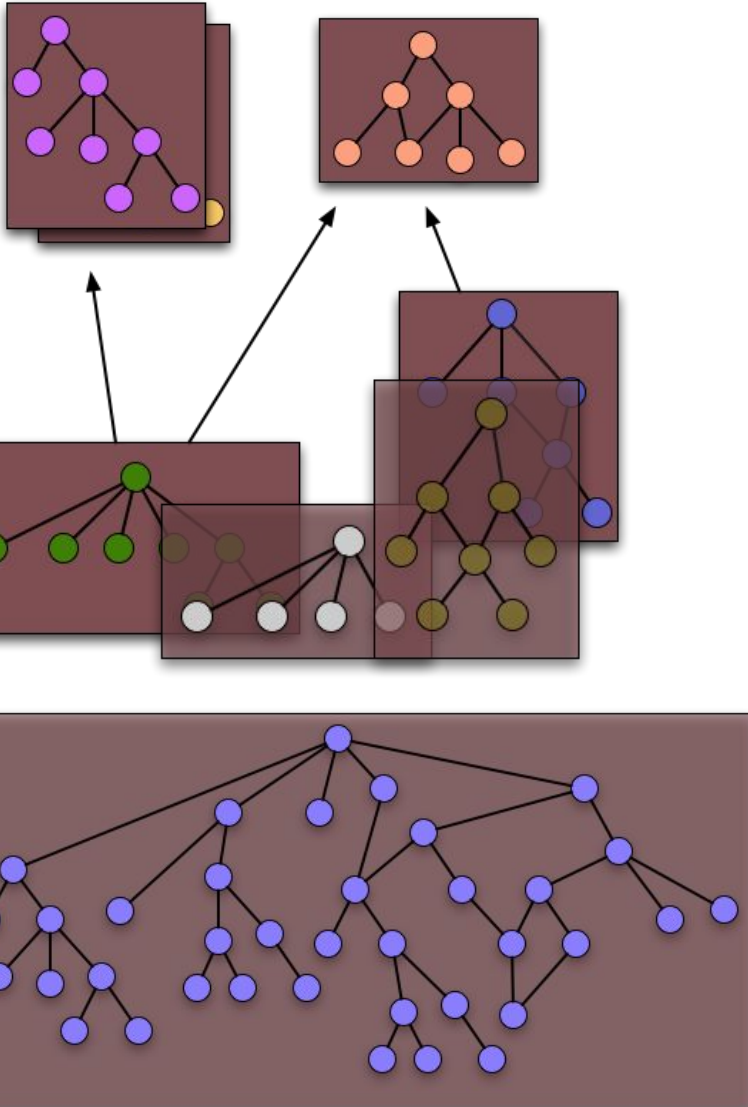
Close

The Ideal World



- " The same **language**
- " No overlap in **coverage**
- " No new **versions**
- " A single **extension tree**
- " Small reusable **modules**

The “Bad” News: The Real World



~~" The same language~~

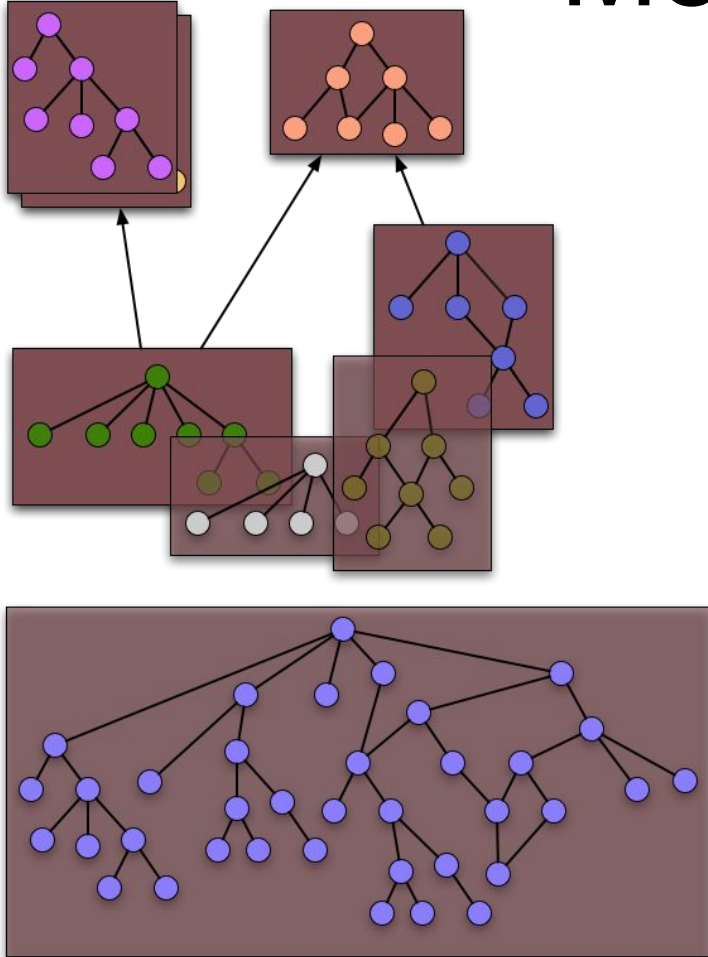
~~" No overlap in coverage~~

~~" No new versions~~

" A single extension tree

~~" Small reusable modules~~

PROMPT: Dealing with the Messy World



- Find similarities and differences between ontologies
- Compare versions of ontologies
- Extract meaningful portions of ontologies
- Integrate in an ontology-editing environment



● Classes ■ Slots ■ Forms ◆ Instances ▲ Queries Prompt

Source classes

cmu

- :THING
- ▶ ● :SYSTEM-CLASS
- ▼ ● Publication
 - Article
 - ▶ ● Book
 - TechReport
 - ▼ ● Thesis
 - MastersThesis
 - PhdThesis
- Academic_Mission
- Email
- Industrial_Mission
- Office
- ▶ ● Organisation
- ▼ ● Person
 - Employee
- ▶ ● Project
- Research_Deliverables
- Research_Group
- Sex
- ▶ ● Work_Activity
- Academic_Org

Suggestions User-defined mappings

Candidate Mappings

Map	Arg1	Arg2	Param
map	● Employee cmu	● Employee umd	
map	● Proceedings cmu	● Proceedings umd	
map	● EMail cmu	● Email umd	
map	● Research_Group cmu	● ResearchGroup umd	
map	● Publication cmu	● Location umd	
map	● Publication cmu	● Publication umd	
map	● Organisation cmu	● Organization umd	
map	● Article cmu	● Article umd	
map	● Book cmu	● Book umd	
map	● Thesis cmu	● Thesis umd	
map	● MastersThesis cmu	● MastersThesis umd	
map	● PhdThesis cmu	● Thesis umd	
map	● Person cmu	● Person umd	
map	● Meeting cmu	● Thing umd	
map	● Director cmu	● Director umd	
map	● Faculty cmu	● Faculty umd	
map	● Teaching cmu	● Thing umd	

Reason for selected suggestion

frames have identical names

Create Mapping

Target classes

umd

- Advertisement
- ▶ ● Article
- Book
- Dictionary
- Editorial
- Manual
- ▶ ● Periodical
 - Proceedings
 - Regulation
 - Specification
 - TechnicalReport
- ▼ ● Thesis
 - DoctoralThesis
 - MastersThesis
- Review
- rdfs_:comment
- PhoneCall
- Software
- Speech
- Location
- PhysicalObject
- Gender

Users can view mappings uploaded from PROMPT in BioPortal

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Home Browse Search

Zebrafish Anatomy

Visualization Class/Type Details Marginal Notes **Mappings** Resources

Subscribe

New Mapping active

Mapping To	Mapped By	Mapped On	Marginal Notes
Adult Mouse Anatomy => blood	Nick Griffith	10/3/07	View Margin Notes

- ...oboInOwl:DbXref
- ...oboInOwl:Definition
- + oboInOwl:ObsoleteClass
- ...oboInOwl:Subset
- ...oboInOwl:Synonym
- ...oboInOwl:SynonymType
- ...Kupffer's vesicle
- ...adaxial cell
- ...anterior axial hypoblast
- ...ball
- blood**
- ...cardiac ventricle
- ...epiphysis
- + floor plate
- ...retinal ganglion cell layer
- ...heart primordium
- ...hindbrain
- ...hypochochord
- ...hypothalamus
- ...intermediate cell mass of me
- ...lens

New Mapping For Abnormal Cell Kind

Point-to-Point Mapping from Abnormal Cell Kind to :

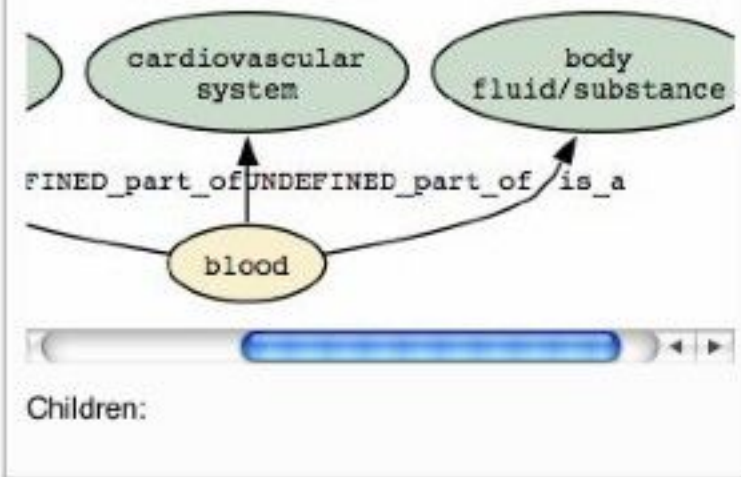
Ontology:

Search:

Close

blood

Preview of blood, MA_0000059



Create

Close

Users can push changes to RSS feeds

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 **BioPortal**


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BIRNLex

Visualization Class/Type Details Marginal Notes **Mappings** Resources **Subscribe** 

New Mapping

Mapping To	Mapped By
Adult Mouse Anatomy => blood	Nick Griffith

Feed To Subscribe To: 

Ontology Branch Concept

... birn_annot:KnownApplica
... birn_annot:ReferenceURL
... obo_annot:AbbrevSource
... obo_annot:DefinitionSour
... obo_annot:ExternalSourc
... TaxonRank
- MaturationStage
 + Adult
 + Elderly
 + Newborn
 + Infant
 + Juvenile
 - Prenatal
 ... Human Prenatal
 ... Mouse Prenatal
Mutation Type
+ core:Collection
... core:Concept
... core:ConceptScheme
... dcmitype:Image
... foaf:Document


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Subscribed Feeds

Comment: [Xenopus anatomy and development => Cell](#)

"I think the definition for this class would..."

Change: [Pathway Ontology => Diabetes Pathway:Definition](#)

"A broad condition with various manifestations that is associated..."

Proposal: [NCI Thesaurus => Cell:Definition](#)

Addition: [NCI Thesaurus => Blood](#)

Statistics

Total Number of Ontologies 72

NCBO Library 59

Remote 13

Number of Classes/Types 300109*

*ontologies which have been parsed and indexed

My Ontologies

[Xenopus anatomy and development =>](#)

7 new comments added

3 proposals added

[Sequence types and features =>](#)

7 new comments added

3 proposals added

[Pathway ontology =>](#)

7 new comments added

3 proposals added

Proposals

Your proposal for [Xenopus anatomy and development => adult](#) is still pending.

Your proposal for [Xenopus anatomy and development => embryo](#) has been approved.

A proposal has been created for [Sequence types and features => mutation](#).

BioPortal will support specialized views on the repository

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BioPortal

Home Browse Search

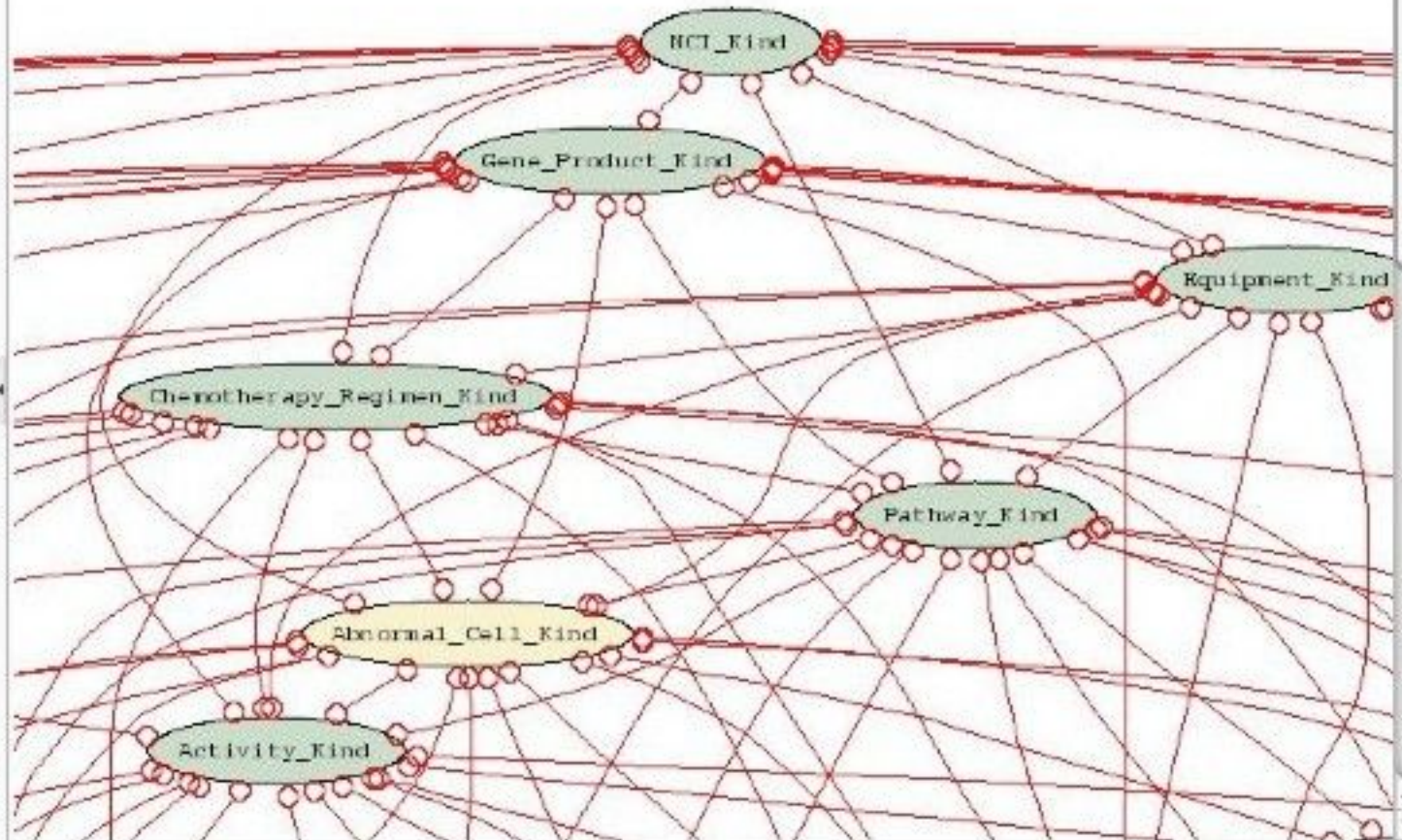
Filter By:

- All
- OBO Foundry
- HL7
- UMLS
- WHO






Ontology	Format	Version
NCI Thesaurus	OWL	
Adult Mouse Anatomy	OBO	
Zebrafish anatomy and development	OBO	
Galen	OWL	
FMA	OBO	
BIRNLex	OWL	

NCI Thesaurus





- + Abnormal_Cell_Kind
- + Activity_Kind
- Anatomy_Kind
 - Anatomic Structure, Syst
 - + Body Fluid or Substan
 - Body Part
 - + Body Region
 - + Body Cavity
 - + Embryologic Structure
 - + Microanatomic Structu
 - + Organ
 - + Organ System
 - + Other Anatomic Conce
- + Biological_Process_Kind
- + Chemicals_and_Drugs_Kind
- + Chemotherapy_Regimen_Kir
- + Diagnostic_and_Prognostic_
- + EO_Anatomy_Kind
- + EO_Findings_and_Disorders
- + Equipment_Kind
- + Findings_and_Disorders_Kin



- + Childhood Hepatic Neoplasms
 - ... Focal Nodular Hyperplasia
 - ... Hepatic Carcinoid Tumor
 - ... Hepatic Dysplastic Nodule
 - ... Hepatic Fibroma
 - ... Hepatic Hemangioma
 - ... Hepatic Inflammatory Myofibroblastic Tumor
 - ... Hepatic Leiomyoma
 - ... Hepatic Lipoma
- + Hepatic Lymphoma
 - ... Hepatic Mesenchymal Hamartoma
- + Hepatic Sarcoma
 - ... Hepatic Vascular Disorder
- + Hepatocellular Adenoma
 - ... Hepatocellular Carcinoma
 - ... Intrahepatic Bile Duct Adenoma
 - ... Intrahepatic Bile Duct Cystadenoma
 - ... Intrahepatic Bile Duct Papillary Cystadenoma
- + Intrahepatic Cholangiocarcinoma
- + Metastatic Malignant Neoplasm of Liver
- + Non-Neoplastic Hepatic Disease

	<p>PubMed is a service of the U.S. National Library of Medicine that includes over 17 million citations from MEDLINE and other life science journals for biomedical articles back to the 1950s. PubMed includes links to full text articles and other related resources.</p>	<p>Elements:29</p>
	<p>ArrayExpress is a public repository for microarray data, which is aimed at storing MIAME-compliant data in accordance with MGED recommendations. The ArrayExpress Data Warehouse stores gene-indexed expression profiles from a curated subset of experiments in the repository.</p>	<p>Elements:8</p>
	<p>ClinicalTrials.gov provides regularly updated information about federally and privately supported clinical research in human volunteers. ClinicalTrials.gov gives you information about a trial's purpose, who may participate, locations, and phone numbers for more details. The information provided on ClinicalTrials.gov should be used in conjunction with advice from health care professionals. Before searching, you may want to learn more about clinical trials.</p>	<p>Elements:206</p>
	<p>A gene expression/molecular abundance repository supporting MIAME compliant data submissions, and a curated, online resource for gene expression data browsing, query and retrieval.</p>	<p>Elements:7</p>
	<p>ARRS GoldMiner provides instant access to images published in selected peer-reviewed radiology journals. This new, web-based system allows viewers to search for images by findings, anatomy, imaging technique, and patient age and sex.</p>	<p>Elements:2</p>

- + Childhood Hepatic Neoplasms
 - ... Focal Nodular Hyperplasia
 - ... Hepatic Carcinoid Tumor
 - ... Hepatic Dysplastic Nodule
 - ... Hepatic Fibroma
 - ... Hepatic Hemangioma
 - ... Hepatic Inflammatory Myofibroblastic Tumor
 - ... Hepatic Leiomyoma
 - ... Hepatic Lipoma
- + Hepatic Lymphoma
 - ... Hepatic Mesenchymal Hamartoma
- + Hepatic Sarcoma
 - ... Hepatic Vascular Disorder
- + Hepatic Vascular Disorder
 - ... Hepatocellular Adenoma
 - ... Hepatocellular Carcinoma
 - ... Intrahepatic Bile Duct Adenoma
 - ... Intrahepatic Bile Duct Cystadenoma
 - ... Intrahepatic Bile Duct Papillary Cystadenoma
- + Intrahepatic Cholangiocarcinoma
 - ... Intrahepatic Cholangiocarcinoma
- + Metastatic Malignant Neoplasms
 - ... Metastatic Malignant Neoplasms
- + Non-Neoplastic Hepatic Disorders
 - ... Non-Neoplastic Hepatic Disorders

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<table border="1"> <thead> <tr> <th>Element ID</th> <th>Annotation Context</th> <th>Element Link</th> </tr> </thead> <tbody> <tr> <td>E-GEOD-4731</td> <td>description</td> <td>View Element</td> </tr> <tr> <td>E-GEOD-5230</td> <td>title</td> <td>View Element</td> </tr> <tr> <td>E-GEOD-5230</td> <td>description</td> <td>View Element</td> </tr> <tr> <td>E-MEXP-199</td> <td>title</td> <td>View Element</td> </tr> <tr> <td>E-MEXP-199</td> <td>description</td> <td>View Element</td> </tr> <tr> <td>E-MEXP-84</td> <td>description</td> <td>View Element</td> </tr> <tr> <td>E-SMDB-2975</td> <td>description</td> <td>View Element</td> </tr> <tr> <td>E-TABM-36</td> <td>description</td> <td>View Element</td> </tr> </tbody> </table>			Element ID	Annotation Context	Element Link	E-GEOD-4731	description	View Element	E-GEOD-5230	title	View Element	E-GEOD-5230	description	View Element	E-MEXP-199	title	View Element	E-MEXP-199	description	View Element	E-MEXP-84	description	View Element	E-SMDB-2975	description	View Element	E-TABM-36	description	View Element
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E-GEOD-5230	title	View Element																											
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E-SMDB-2975	description	View Element																											
E-TABM-36	description	View Element																											
 ClinicalTrials.gov	ClinicalTrials.gov provides regularly updated information about federally and privately supported clinical research in human volunteers. ClinicalTrials.gov gives you information about a trial's purpose, who may participate, locations, and phone numbers for more details. The information provided on ClinicalTrials.gov should be used in conjunction with advice from health care professionals. Before searching, you may want to learn more about clinical trials.	Elements:206																											
 Gene Expression Omnibus	A gene expression/molecular abundance repository supporting MIAME compliant data submissions, and a curated, online resource for gene expression data browsing, query and retrieval.	Elements:7																											

1 / 4	Experiment : E-GEOD-4612	Submitter(s) : Katzenellenbogen	Lab : Hadassah University Hospital
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Experiment Design Type : individual genetic characteristics , co expression , time series

(Generated description): Experiment with 12 hybridizations, using 12 samples of species [Mus musculus], using 12 arrays of array design [Affymetrix GeneChip® Mouse Genome 430A 2.0 [Mouse430A_2]], producing 12 raw data files and 12 transformed and/or normalized data files.

(Submitter's description 1): We studied the molecular mechanisms of hepatocellular carcinoma (HCC) initiation and promotion using the Mdr2-knockout (Mdr2-KO) mice at pre-cancerous stages of liver disease. These mice lack the liver-specific P-glycoprotein responsible for phosphatidylcholine transport across the canalicular membrane. Portal inflammation ensues at an early age followed by the development of HCC between the ages of 12 and 15 months. Liver tissue samples of Mdr2-KO and control Mdr2-heterozygotes mice aged 3 and 12 months, were subjected to histological, biochemical and gene expression profiling analysis using Affymetrix Mouse Genome Array. The RNA samples from Mdr2-KO and control heterozygous mice aged 3 and 12M (3 males in each experimental group) were subjected to genome scale gene expression profiling with Affymetrix Mouse Array. The gene expression values were extracted with the help of MAS 5.0 software, and analyzed by cluster analysis, and by fold change filtering

[Retrieve data >>](#)

[Experimental protocols >>](#)

[Providers >>](#)






[Array design used >>](#)

- [Experiment's directory in the FTP >>](#)
- [MAGE-ML : \(.gz \(661 KB\) \)](#)
- [Sample annotation : \(.txt .xls \)](#)
- [Experiment design : \(.png .svg \)](#)
- [Detailed sample annotation : \(.txt .xls \)](#)

[Bibliographic references >>](#)

[Samples >>](#)

- + Childhood Hepatic Neoplasms
 - Focal Nodular Hyperplasia
 - Hepatic Carcinoid Tumor
 - Hepatic Dysplastic Nodule
 - Hepatic Fibroma
 - Hepatic Hemangioma
 - Hepatic Inflammatory Myofibroblastic Tumor
 - Hepatic Leiomyoma
 - Hepatic Lipoma
- + Hepatic Lymphoma
- Hepatic Mesenchymal Hamartoma
- + Hepatic Sarcoma
- + Hepatic Vascular Disorder
 - Hepatocellular Adenoma
 - Hepatocellular Carcinoma**
 - Intrahepatic Bile Duct Adenoma
 - Intrahepatic Bile Duct Cystadenoma
 - Intrahepatic Bile Duct Papillary Cystadenoma
- + Intrahepatic Cholangiocarcinoma
- + Metastatic Malignant Neoplasm of Liver
- + Non-Neoplastic Hepatic Disorder

	<p>PubMed is a service of the U.S. National Library of Medicine that includes over 17 million citations from MEDLINE and other life science journals for biomedical articles back to the 1950s. PubMed includes links to full text articles and other related resources.</p>	<p>Elements:29</p>
	<p>ArrayExpress is a public repository for microarray data, which is aimed at storing MIAME-compliant data in accordance with MGED recommendations. The ArrayExpress Data Warehouse stores gene-indexed expression profiles from a curated subset of experiments in the repository.</p>	<p>Elements:8</p>
	<p>ClinicalTrials.gov provides regularly updated information about federally and privately supported clinical research in human volunteers. ClinicalTrials.gov gives you information about a trial's purpose, who may participate, locations, and phone numbers for more details. The information provided on ClinicalTrials.gov should be used in conjunction with advice from health care professionals. Before searching, you may want to learn more about clinical trials.</p>	<p>Elements:206</p>
	<p>A gene expression/molecular abundance repository supporting MIAME compliant data submissions, and a curated, online resource for gene expression data browsing, query and retrieval.</p>	<p>Elements:7</p>
	<p>ARRS GoldMiner provides instant access to images published in selected peer-reviewed radiology journals. This new, web-based system allows viewers to search for images by findings, anatomy, imaging technique, and patient age and sex.</p>	<p>Elements:2</p>

Gene Expression Omnibus



e.g., element GDS1989

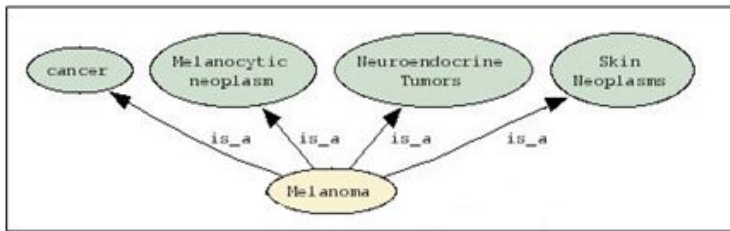
Entis Web Service API

Resource element fetching

```
<title>
  Melanoma progression
</title>

<description>
  Analysis of tissue specimens representing benign nevus, atypical nevus, melanoma in situ, vertical growth phase (VGP) melanoma, and metastatic growth phase (MGP) melanoma. Results identify expression signatures that distinguish benign and atypical nevi and melanomas in situ from VGPs and MGPs.
</description>
```

Annotation



172 closure annotations and 171 useful.

Examples:

Cancer, concept (DOID:162) in ontology *Human disease*

Skin Neoplasms, concept (DOID:3165) in ontology *Human disease*

Transitive closure

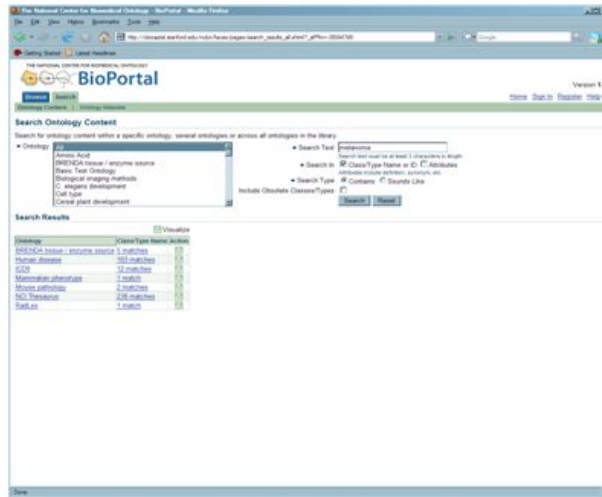
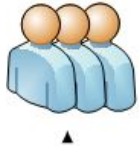
```
<title>
  Melanoma progression
</title>

<description>
  Analysis of tissue specimens representing benign nevus, atypical nevus, melanoma in situ, vertical growth phase (VGP) melanoma, and metastatic growth phase (MGP) melanoma. Results identify expression signatures that distinguish benign and atypical nevi and melanomas in situ from VGPs and MGPs.
</description>
```

23 direct annotations (4 title, 19 description)

Example:

Melanoma, concept (DOID:1909) in ontology *Human disease*.

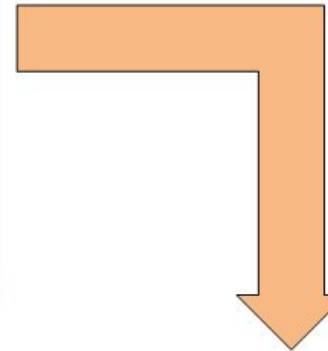


Bioportal search for “melanoma”

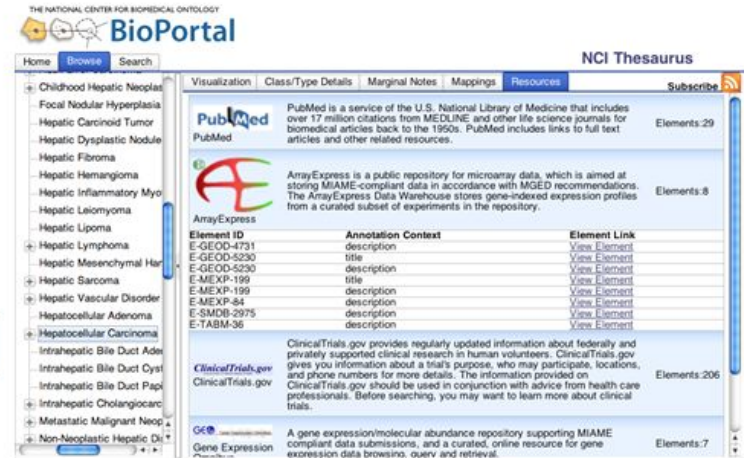
362 matches in all Bioportal ontologies.

Example:

Melanoma, concept (DOID:1909) in ontology *Human disease*.



OBS index request

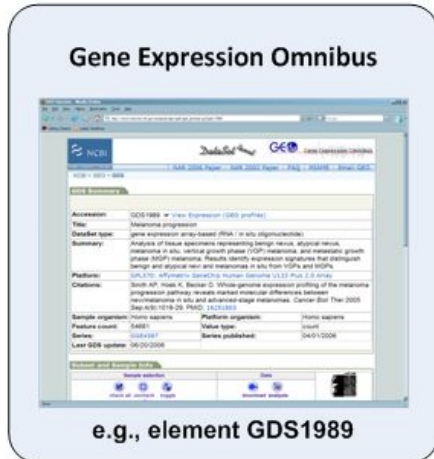


Onrez index results display

- 227 PubMed elements,
- 3 ArrayExpress elements,
- 969 ClinicalTrials.gov elements,
- 10 ARRS GoldMiner elements,
- 3 Gene Expression Omnibus elements.

Example:

Melanoma progression element (GDS1989)








Gene Expression Omnibus

e.g., element GDS1989



Link to the original resource element

- + Childhood Hepatic Neoplasms
 - ... Focal Nodular Hyperplasia
 - ... Hepatic Carcinoid Tumor
 - ... Hepatic Dysplastic Nodule
 - ... Hepatic Fibroma
 - ... Hepatic Hemangioma
 - ... Hepatic Inflammatory Myofibroblastic Tumor
 - ... Hepatic Leiomyoma
 - ... Hepatic Lipoma
- + Hepatic Lymphoma
 - ... Hepatic Mesenchymal Hamartoma
- + Hepatic Sarcoma
 - ... Hepatic Vascular Disorder
- + Hepatic Vascular Disorder
 - ... Hepatocellular Adenoma
 - ... **Hepatocellular Carcinoma**
 - ... Intrahepatic Bile Duct Adenoma
 - ... Intrahepatic Bile Duct Cystadenoma
 - ... Intrahepatic Bile Duct Papillary Cystadenoma
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If we build it, will they come?

A problem in both technology and sociology

- How can we identify communities of likely early adopters?
- How will we know when we will have sufficient functionality to entice early adopters to adopt?
- How can we measure the affects of our technology on the way that science gets done?
- How can we engage in participatory design of technology that potential users cannot even imagine?

BioPortal User Group

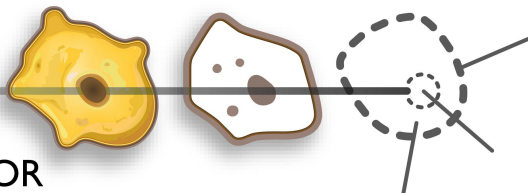
- CTSAAs
- Immunology
- Imaging
- RadLex
- W3C HCLSIG
- BioPAX
- CVRGrid
- caBIG
- HL7
- MODs
- GO Consortium
- BIRN

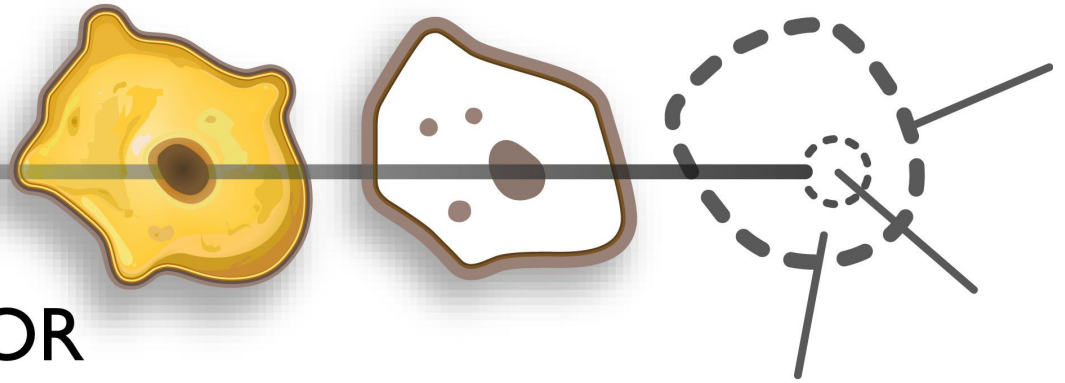
BioPortal can build an online community of users who

- Develop, upload, and apply ontologies
- Map ontologies to one another
- Comment on ontologies via “marginal notes” to give feedback
 - To the ontology developers
 - To one another
- Make proposals for specific changes to ontologies
- Stay informed about ontology changes and proposed changes via active feeds

Goals for the NCBO

- Providing technology for ontology archiving, access, browsing, visualization, peer review, mapping, versioning
- Making most biomedical ontologies accessible via a common portal
- Educating the community about principles of ontology development and use
- Serving as a generalizable model for the formalization of knowledge in e-science





NATIONAL CENTER FOR

BIOMEDICAL ONTOLOGY

<http://bioontology.org>