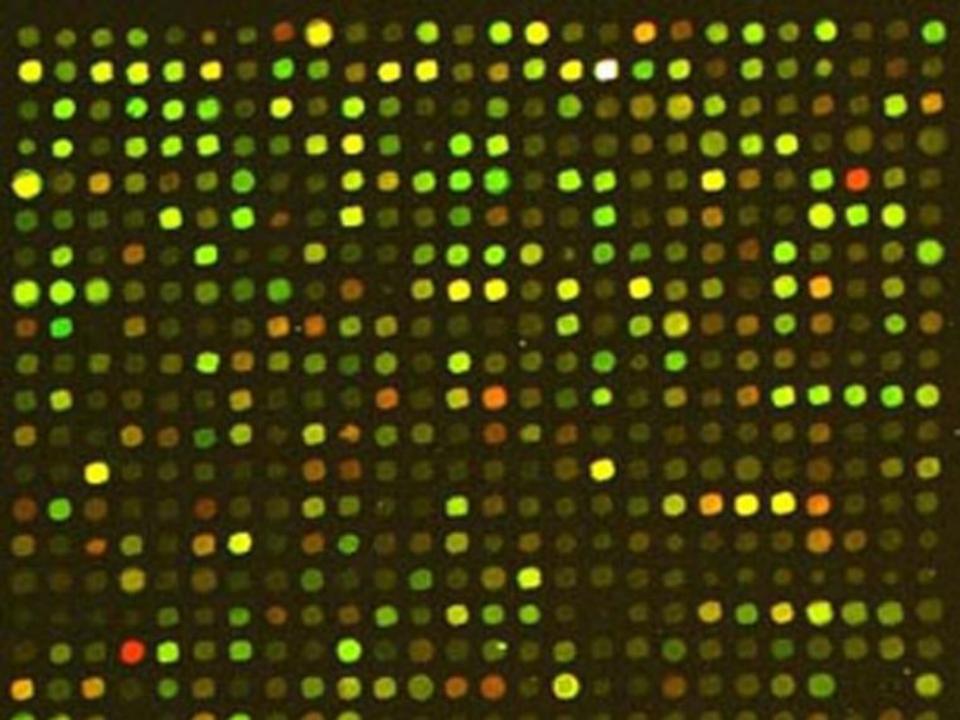


Mark Musen and the NCBO team Stanford University

Big Data Needs Big Ontology!







GO:0016265 : death [12861 gene products]

GO:0040007 : growth [13918 gene products]

GO:0032502 : developmental process [49009 gene products]

□ ■ CO:0002376 : immune system process [12542 dens products]

GO:0051234: establishment of localization [55604 gene products]

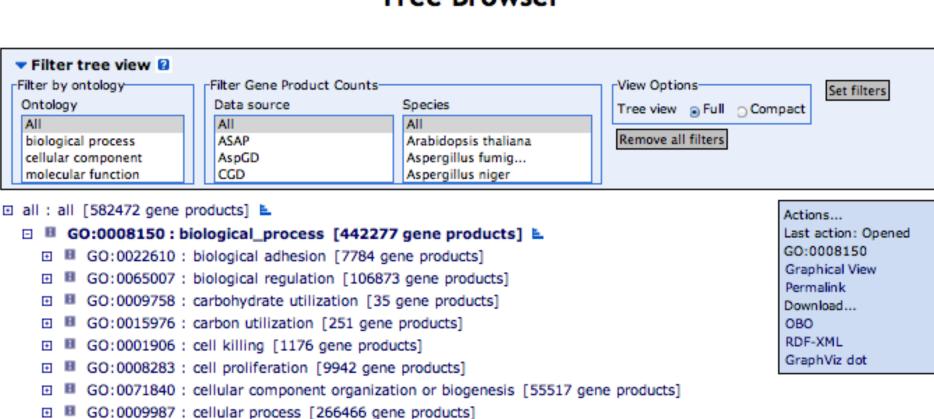
STATE OF THE PARTY.

APPEN



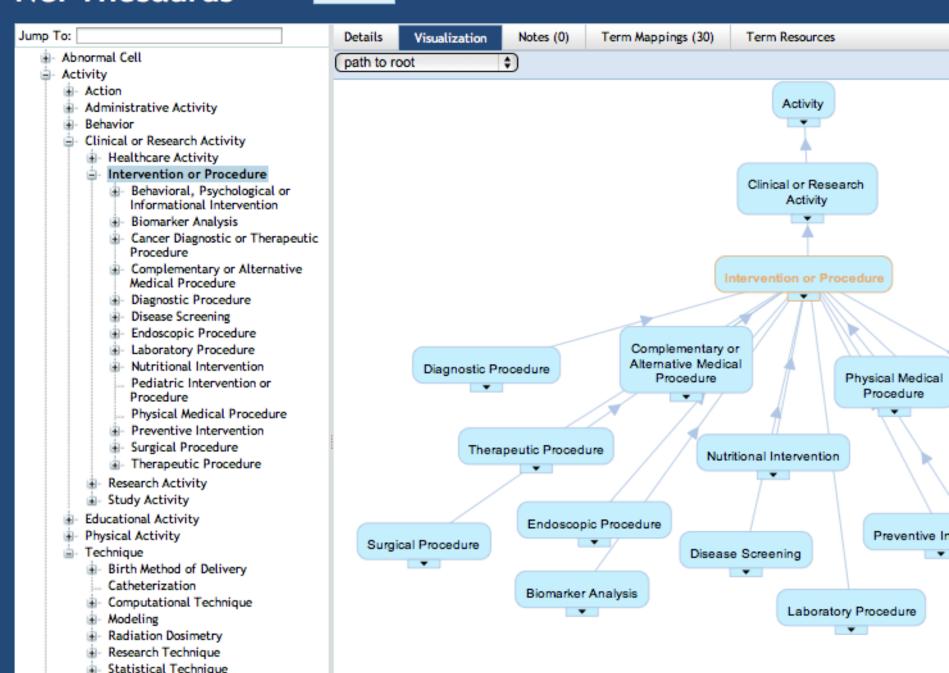
| Search Browse | BLAST Homol | og Annotations Tools & | Resources Help | |
|---------------|-------------|-------------------------|--------------------|--|
| Search GO | ● te | erms ogenes or proteins | exact match Submit | |

Tree Browser



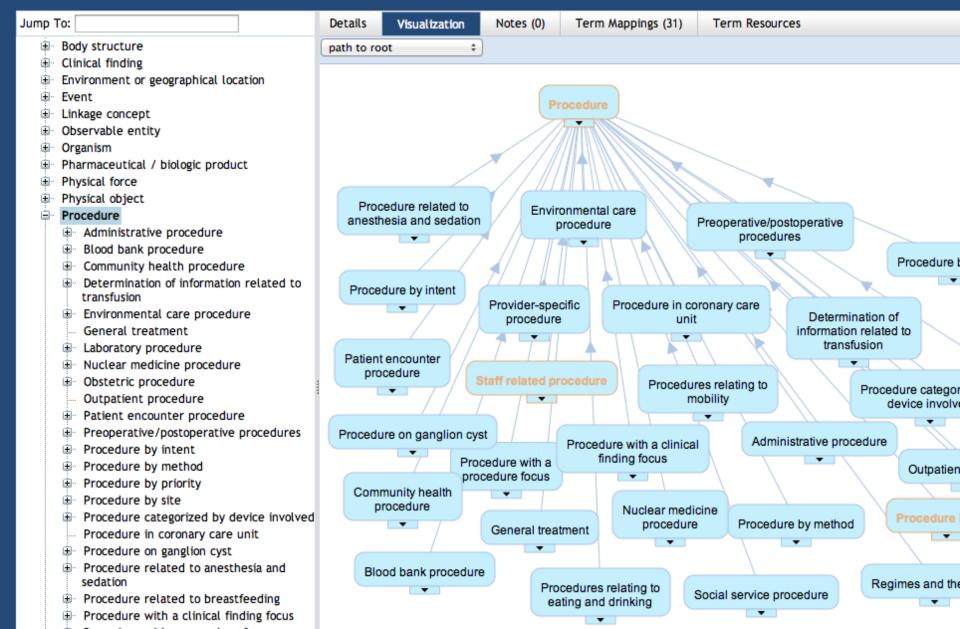
NCI Thesaurus





SNOMED Clinical Terms





The National Center for Biomedical Ontology

- We create and maintain a library of biomedical ontologies and terminologies.
- We build tools and Web services to enable the use of ontologies and terminologies.
- We collaborate with scientific communities that develop and use ontologies and terminologies in biomedicine.



Go to Resource Index



NCBO User Profile

Jin-Dong Kim
Database Center for Life
Science

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Video

Learn about Biomedical Ontologies. Watch a series of introductory videos.



Browse ontologies in BioPortal!

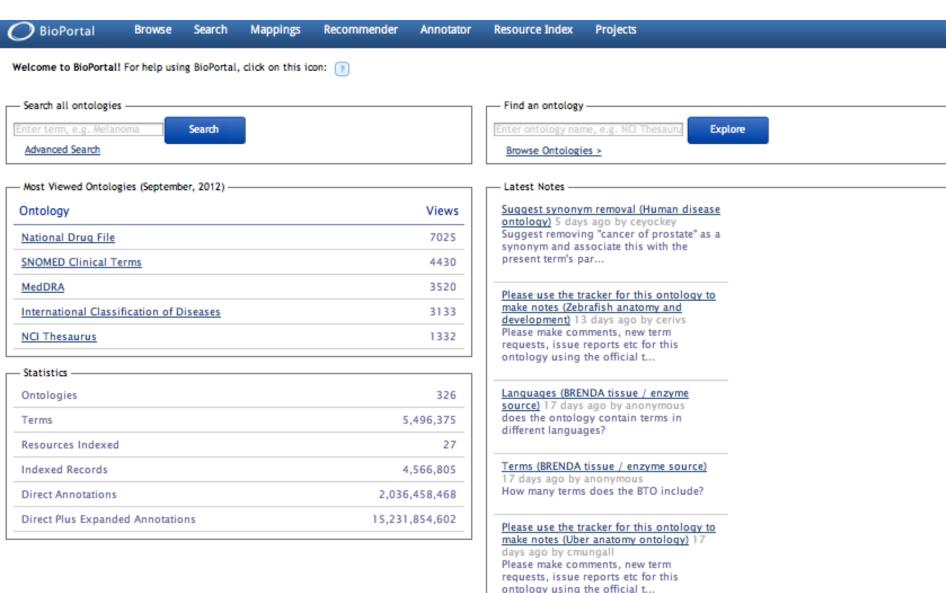
0

Go to Protégé

BioPortal allows users to browse, search and visualize ontologies.



BioPortal Ontology Repository

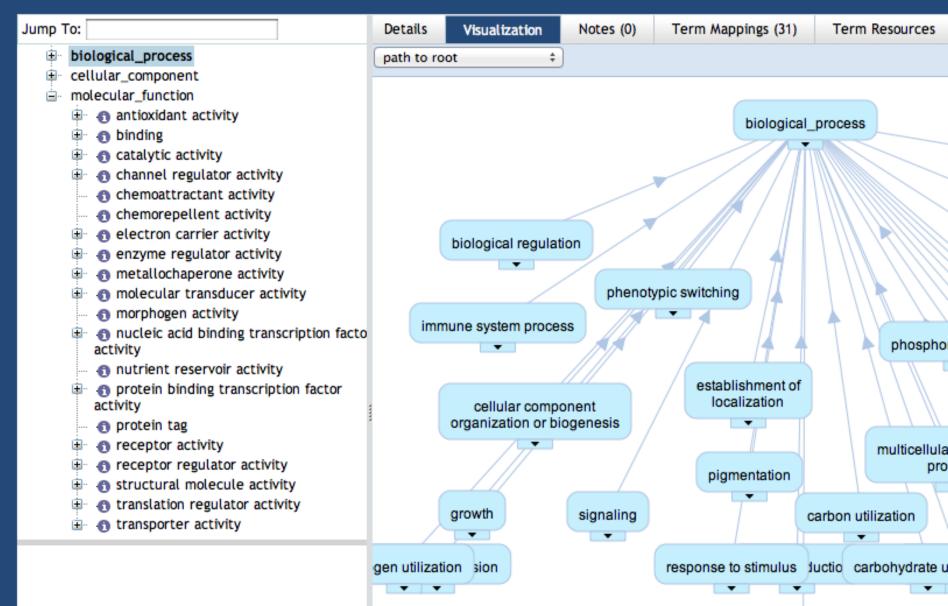


http://bioportal.bioontology.org

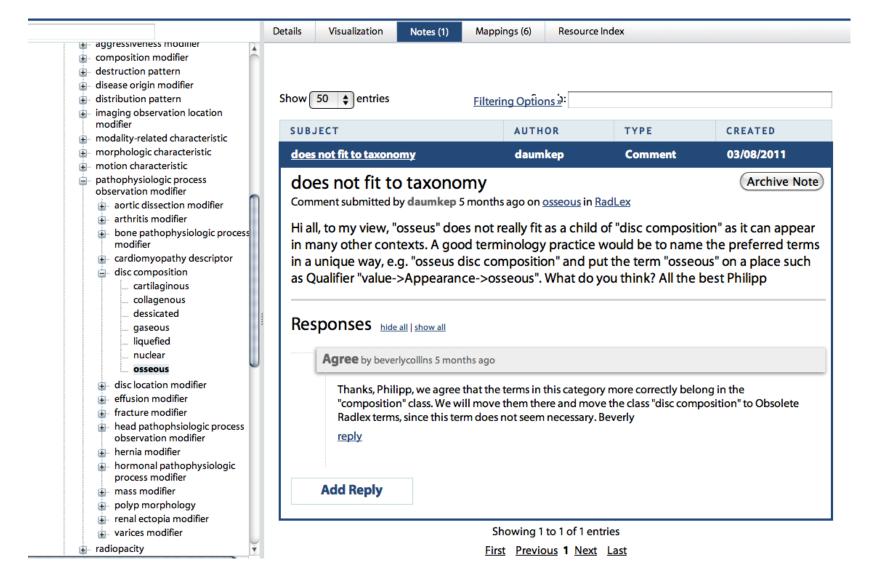
| ◯ BioPortal B | pings Recommender Annotator | | | Resource Inde | ex Projects | Sign In Help Feedt | | |
|---|-----------------------------|------------|-------|---------------|-------------|------------------------|------------|---|
| Browse rowse the library of ontologi | es 💡 | | | | | | | |
| ew: Configure which ontolog | gies you see in BioPortal | | | | | | | |
| FILTER BY CATEGORY | All Categories | 7 | | | Submit | New Ontology | | |
| FILTER BY GROUP ? | All Groups | • , | | | | | | |
| FILTER BY TEXT | | | | | | | | |
| | | | | | | | | Subscribe to all updat |
| ONTOLOGY NAME | | VISIBILITY | TERMS | NOTES | REVIEWS | PROJECTS | UPLOADED | CONTACT |
| ABA Adult Mouse Brain (ABA | A) | Public | 913 | 0 | 0 | 7 | 08/08/2009 | Allen Institute for Brain Science |
| Adverse Event Reporting or | ntology (AERO) | Public | 374 | 1 | 0 | 3 | 11/13/2012 | Melanie Courtot |
| African Traditional Medicin | e (ATMO) | Public | 223 | 2 | 2 | 4 | 06/28/2009 | Ghislain Atemezing |
| AI/RHEUM (AIR) | | Public | 681 | 0 | 0 | 2 | 02/05/2010 | May Cheh |
| Amino Acid (amino-acid) | | Public | 46 | 0 | 0 | <u>5</u> | 07/02/2010 | Nick Drummond, Georgina Moulton, Robert Stevens, Phil Lord |
| Amphibian gross anatomy (| (AAO) | Public | 1,603 | 0 | 0 | <u>5</u> | 07/22/2011 | David Blackburn |
| mphibian taxonomy (ATO) |) | Public | 6,135 | 0 | 0 | 3 | 11/02/2009 | AmphiAnat list |
| natomical Entity Ontology | (AEO) | Public | 250 | 0 | 0 | 3 | 06/01/2012 | EMAP Administrators |
| nimal natural history and | life history (ADW) | Public | 360 | 0 | 0 | 2 | 08/31/2010 | Animal Diversity Web technical staff |
| apollo-akesios (apollo) | | Public | 3 | 0 | 0 | 1 | 09/30/2010 | Jeremy Espino |
| Ascomycete phenotype onto | ology (APO) | Public | 328 | 0 | 0 | 4 | 03/01/2012 | SGD curators |
| Basic Formal Ontology (BFO) | | Public | 39 | 0 | 1 | 21 | 07/24/2009 | Holger Stenzhorn |
| Basic Vertebrate Anatomy (anatomy) | basic-vertebrate-gross- | Public | 99 | 0 | 0 | 4 | 01/16/2007 | |
| Bilateria anatomy (BILA) | | Public | 114 | 0 | 0 | 4 | 03/03/2010 | Thorsten Heinrich |
| Bio-health ontological know (ibrosis (OntoKBCF) | ledge base- cystic | Public | 405 | 0 | 0 | 3 | 09/18/2012 | Xia Jing |
| BioAssay Ontology (BAO) | | Public | 1,292 | 0 | 0 | 7 | 04/12/2012 | Stephan Schurer |
| cioinformatics operations, ormats and topics (EDAM) | | Public | 2,719 | 0 | 2 | <u>6</u> | 07/05/2012 | Jon Ison |
| Bioinformatics Web Service | Ontology (OBIWs) | Public | 195 | 0 | 0 | 1 | 07/17/2012 | jie zheng |
| Biological imaging methods | (FBbi) | Public | 624 | 0 | 0 | <u>6</u> | 06/09/2011 | Chris Woodcock |
| | (max) | | 244 | | 2 | | 00/04/0040 | Trish Whetzel, Csongor Nyulas, Natashi |

Gene Ontology

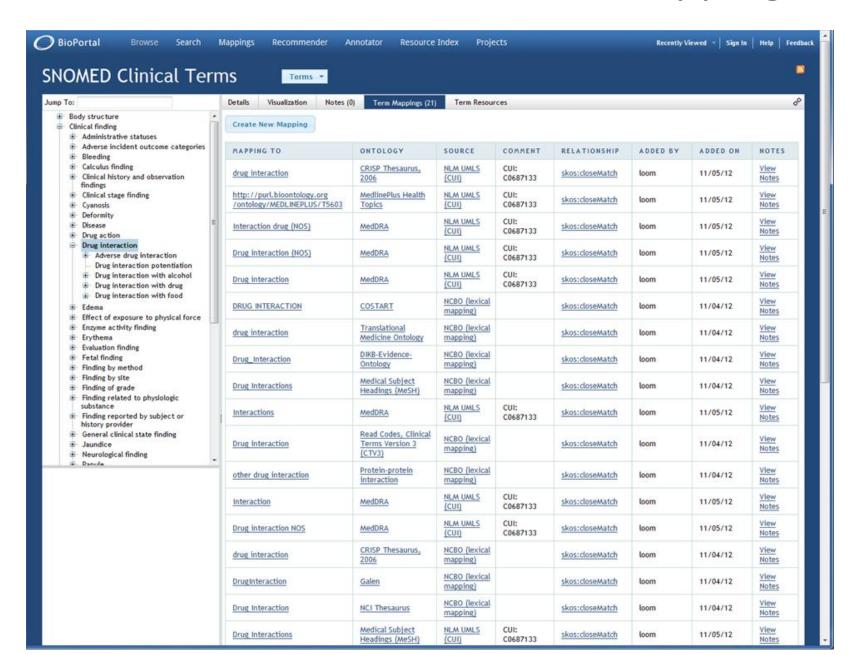




Community Comments in BioPortal

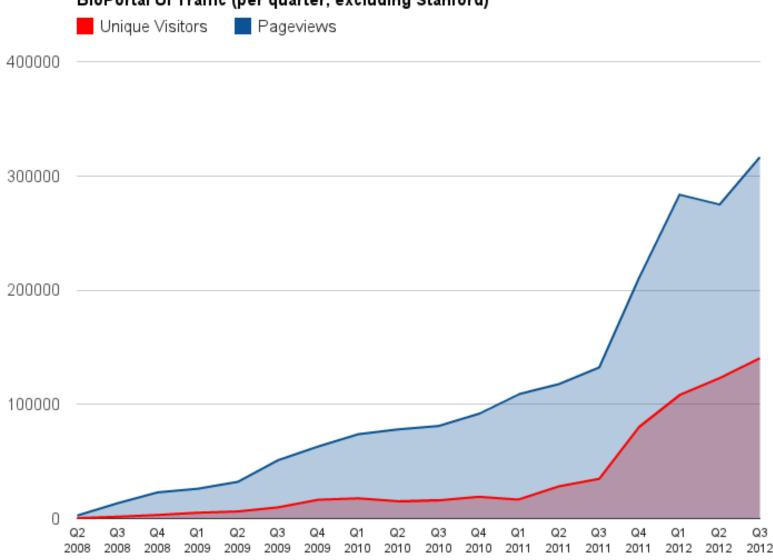


Users can view and create mappings



BioPortal Traffic – Volume

BioPortal UI Traffic (per quarter, excluding Stanford)



BioPortal UI Traffic – Top Users

Thomson Financial

Japan Network Information Center

Department of Veterans Affairs

National Health Service (UK)

University of British Columbia

University of Florida

MRC Human Genome Mapping Project Resource Centre

University of California, San Diego

Mayo Foundation for Medical Education and Research

University of California, Davis

University of Pennsylvania

BioPortal is building an online community of users who

- Develop, upload, and apply ontologies
- Map ontologies to one another
- Comment on ontologies via "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 "push" technology
- Incorporate BioPortal services into their own technologies

Browse

Search

Mappings

Recommender

Annotator

Resource Index

Annotator

Get annotations for biomedical text with terms from the ontologies ?

insert sample text

Cyclic nucleotide phosphodiesterases (PDEs) are enzymes that regulate the cellular levels of the second messengers, cAMP and cGMP, by controlling their rates of degradation. There are 11 different PDE families, with each family typically having several different isoforms and splice variants. These unique PDEs differ in their three-dimensional structure, kinetic properties, modes of regulation, cellular expression, and inhibitor sensitivities. Current data suggest that individual isozymes modulate distinct regulatory pathways in the cell.

Select Ontologies

GO ×

clear selection select from list

Select UMLS Semantic Types

Type here to select UMLS semantic types

Include Mappings:

Automatic

Manual

Include Ancestors Up To Level: All \$

Get Annotations

Browse

Search

Mappings

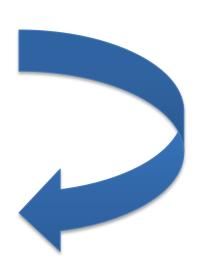
Annotator

Get annotations for biomedical text with terms from the ontologies 🕞



insert sample text

Cyclic nucleotide phosphodiesterases (PDEs) are enzymes that regulate the cellular levels of the second messengers, cAMP and cGMP, by controlling their rates of degradation. There are 11 different PDE families, with each family typically having several different isoforms and splice variants. These unique PDEs differ in their three-dimensional structure, kinetic properties, modes of regulation, cellular expression, and inhibitor sensitivities. Current data suggest that individual isozymes modulate distinct regulatory pathways in the cell.



Get Annotations

Annotations total results 5 (direct 2 / ancestor 3

| TERM filter | ONTOLOGY filter | TYPE filter | CONTEXT | MATCHED TERM filter | MATCHED ONTOLOG |
|--------------------|-----------------|-------------|--|---------------------|-----------------|
| cell | Gene Ontology | direct | regulatory pathways in the cell | cell | Gene Ontology |
| metabolic process | Gene Ontology | ancestor | controlling their rates of degradation. There are 11 different | catabolic process | Gene Ontology |
| cellular_component | Gene Ontology | ancestor | regulatory pathways in the cell | cell | Gene Ontology |
| catabolic process | Gene Ontology | direct | controlling their rates of degradation. There are 11 different | catabolic process | Gene Ontology |
| biological_process | Gene Ontology | ancestor | controlling their rates of degradation. There are 11 different | catabolic process | Gene Ontology |

NCBO Annotator

- Typically called as Web service by client programs
- Takes as input some text; generates as output links between the text and terms in any or all NCBO ontologies
- Enables high-throughput
 - Annotation of textual metadata
 - Text mining of electronic medical records
 - Concept-based indexing and retrieval of online data sets



BioPortal

Browse

Search

Mappings

Recommender

Annotator

Resource Index

Projects

Search Resource Index Search biomedical resources 🕞

Start typing to find terms to search the index with

Search Resource Index

select ontologies to search

railable December

caNanoLab

| Available Resources | | |
|---------------------|-----------------|--|
| RESOURCE | DESCRIPTION | |
| | | |

This database provides a searchable and browsable list of aging related genes and their effects ... [more]

AgingGenesDB (via NIF)

ArrayExpress is a public repository for microarray data, which is aimed at storing MIAME-compli ... [more]

ArrayExpress ARRS GoldMiner ARRS GoldMiner provides instant access to images published in selected peer-reviewed radiology ... [more]

Biositemaps

Biositemaps represent a mechanism for computational biologists and bio-informaticians to openly ... [more] caNanoLab is a data sharing portal designed to facilitate information sharing in the biomedical ... [more] The Cell Centered Database is a publicly accessible resource for high resolution 2D, 3D and 4D ... [more]

Cell Centered Database (via NIF)

Database of Genotypes and The database of Genotypes and Phenotypes (dbGaP) was developed to archive and distribute the re ... [more] Phenotypes DrugBank

Online Mendelian Inheritance in Man

ClinicalTrials.gov Conserved Domain Database (CDD)

Clinical Trials, gov provides regularly updated information about federally and privately support ... [more] The Conserved Domain Database (CDD) contains protein domain models imported from outside source ... [more

DrugBank is offered to the public as a freely available resource. Use and re-distribution of th ... [more]

A gene expression/molecular abundance repository supporting MIAME compliant data submissions, ... [more] Molecular Imaging and Contrast Agent Database

ModelDB provides an accessible location for storing and efficiently retrieving computational ne ... [more]

Pathway Commons is a convenient point of access to biological pathway information collected fro ... [more]

OMIM is a comprehensive, authoritative, and timely compendium of human genes and genetic phenot ... [more]

Gene Expression Omnibus DataSets MICAD

ModelDB (via NIF)

Pathway Commons



NCBO User Profile

Jin-Dong Kim
Database Center for Life
Science

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Video

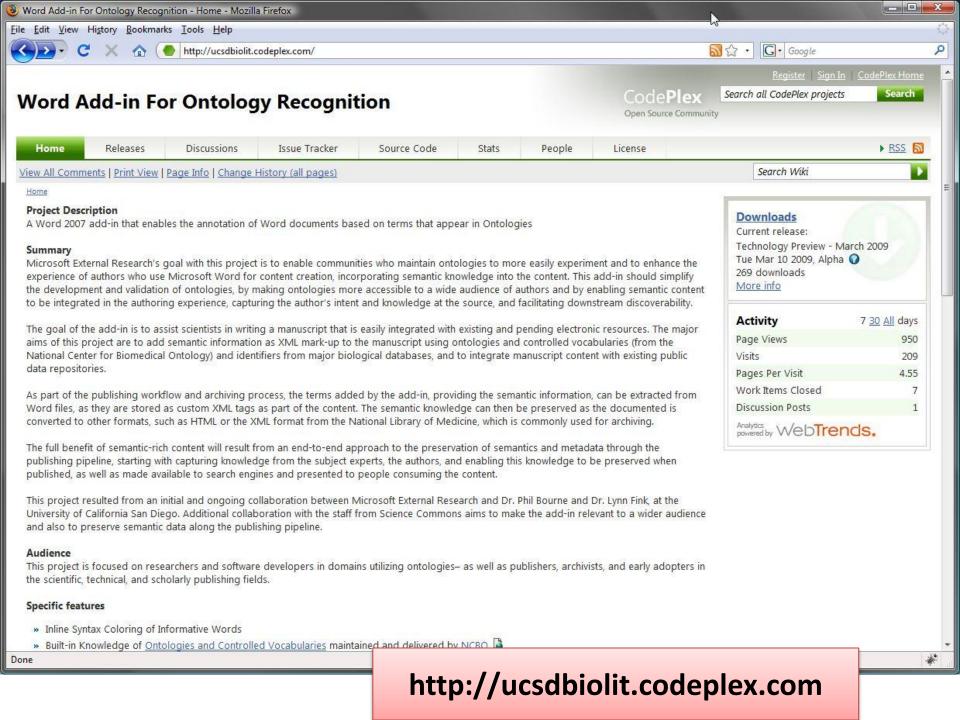
Learn about Biomedical Ontologies. Watch a series of introductory videos.



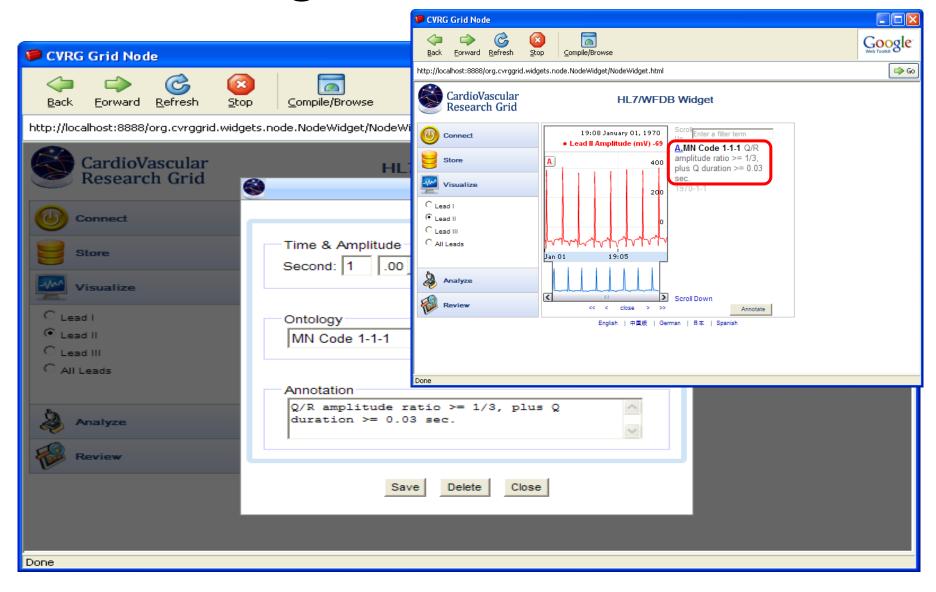
Browse ontologies in BioPortal!

BioPortal allows users to browse, search and visualize ontologies.

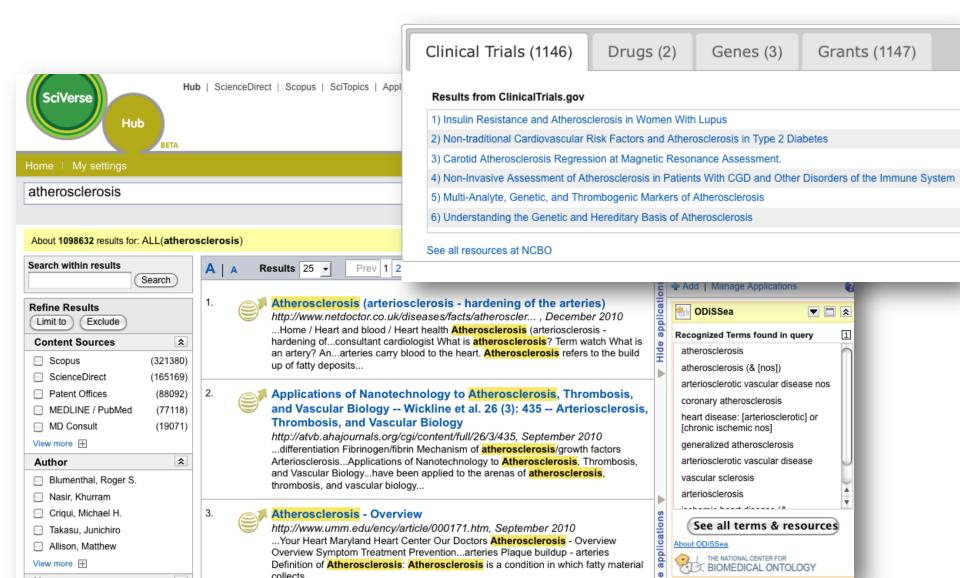




ECG Gadget uses NCBO services



Elsevier SciVerse uses NCBO technology for ontology-based IR



Simbios uses **NCBO** Ontology Web Widgets



7. Classify your project (required)

- a) Choose one or more terms from the Biological Resource Ontology that most closely describe your project. These will be used in searches so others can more easily find your work.
- Search for a term by starting to enter it in the box below all terms that contain what you entered will be shown below the box. Select one of these, or. . .
- 2) Alternatively, you can try browsing the <u>Biological Resource Ontology</u>. Only terms under BRO:Resource are allowed. You are most likely to find suitable terms under BRO:Resource → BRO:Software → BRO:Modeling_and_Simulation. Enter the term or any part of it in the box below without the BRO prefix and continue as in 1).

Restriction: only terms from the BRO:Resource branch of the Biological Resource Ontology allowed.

Samples: Neuromuscular_Model, Molecular_Dynamics

Ontology terms:

 Computational_Model
 Remove

 Protein_Model
 Remove

 Standalone_Application
 Remove

 Structure-Based_Protein_Classification
 Remove

Add (click after choosing a term)

b) Please also choose one or more keywords for your project. These will also be used in searches so others can more easily find your project.

Keywords: (required)

allosteric communication Remove allostery Remove

Add (one at a time)

To save changes, click Save Project Info here or at the bottom of the page.

8. Short Purpose/Synopsis (required)

Please provide a synopsis of your project. This will be displayed in the search results.

Restriction: 255 characters

Samples:

- 1) Provide an easy-to-use application for manipulating RNA structures
- Provides the code base for creating, simulating, and visualizing threedimensional finite-element models of skeletal muscle.
- Geometric models in VTK/XML PolyData format for download, for use in cardiovascular applications.

Collaborations with other NCBCs

- Simbios uses NCBO Web widgets for accessing ontology-derived value sets
- I2b2 uses NCBO ontology repository to build its ontology "hive"
- NCBO Annotator uses mgrep named-entity recognizer from NCIBI
- NA-MIC Slicer system uses NCBO for semantic annotation of images
- NCBO has collaborated with CCB and NCIBI to develop BioSitemaps

"Collaborating R01" Program

- Nine NIH-funded collaborations with NCBO to date
- Collaborators have benefitted from
 - Early access to NCBO technology to enable their research
 - Opportunity to shape some of our work
- NCBO has benefitted from
 - Continuous feedback on our work
 - Actual code that we have incorporated into our technology (e.g., support for ontology "views")

Driving Biological Projects (DBPs)

- Nine DBPs supported by NCBO to date
- DBPs allow NCBO to target "applications pull"
- Lots of important contributions
 - Structural visualization of clinical-trial results
 - Tool for annotating electrophysiological signals
 - Concept-based search of data regarding therapeutic nanoparticles in caNanoLab
 - Annotation of microarray data in Array Express
 - General-purpose tool for ontology-based enrichment analysis

Our initial proposal did not predict ...

- The exponential increase in the use of semantic technology in biomedicine
- The huge user community that NCBO would attract
- The time that it would take for our technology to become sufficiently robust for other investigators to stake their success on us
- All the applications that investigators would find for our technology
- That people would take us for granted

The National Center for Biomedical Ontology

- We create and maintain a library of biomedical ontologies and terminologies.
- We build tools and Web services to enable the use of ontologies and terminologies.
- We collaborate with scientific communities that develop and use ontologies and terminologies in biomedicine.



