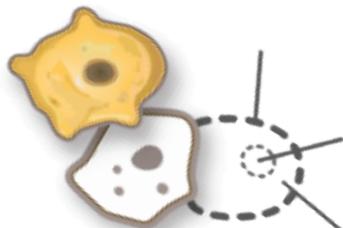


NCBO SPARQL Endpoint

**Trish Whetzel
Outreach Coordinator**



THE NATIONAL CENTER FOR
BIOMEDICAL ONTOLOGY

```

1 PREFIX omv: <http://omv.ontoware.org/2005/05/ontology
2
3 SELECT ?ont ?name ?acr
4 WHERE {
5     ?ont a omv:Ontology .
6     ?ont omv:acronym ?acr .
7     ?ont omv:name ?name .
8 }
9

```

Results: [Browse](#) [run query](#) [reset](#)

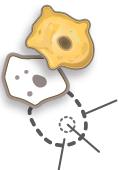
BioPortal SPARQL is a service to query BioMedical ontologies using the SPARQL standard. Ontologies have been transformed into RDF triples from their original formats (OWL, OBO and UMLS/RRF, ...) and asserted into a triple store. This service provides programmatic access to that triple store.

[Documentation](#)

[SPARQL Examples](#)

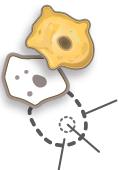
ont	name	acr
<http://bioportal.bioontology.org/ontologies/44840>	"Human disease"	"DOID"
<http://bioportal.bioontology.org/ontologies/46187>	"HOM-DATASOURCE_OSHPD"	"HOM-DATASOURCE_OSHPD"
<http://bioportal.bioontology.org/ontologies/46255>	"Common Anatomy Reference Ontology"	"CARO"
<http://bioportal.bioontology.org/ontologies/40477>	"NanoParticle Ontology"	"NPO"
<http://bioportal.bioontology.org/ontologies/42598>	"Gene Ontology"	"GO"
<http://bioportal.bioontology.org/ontologies/45037>	"Yeast phenotypes"	"YPO"
<http://bioportal.bioontology.org/ontologies/40654>	"NanoParticle Ontology"	"NPO"
<http://bioportal.bioontology.org/ontologies/46097>	"Mosquito insecticide resistance"	"MIRO"
<http://bioportal.bioontology.org/ontologies/42069>	"Cereal plant trait"	"TO"
<http://bioportal.bioontology.org/ontologies/47271>	"HOM-PCSTEST"	"HOM-PCSTEST"
<http://bioportal.bioontology.org/ontologies/45990>	"HOM-ICD9CM-ECODES"	"HOM-ICD9CM-ECODES"

<http://sparql.bioontology.org/>



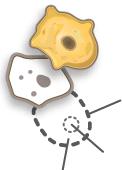
BioPortal SPARQL Content

- All ontologies from BioPortal
 - Original ontology format (OBO, OWL, UMLS/RRF) transformed into RDF
 - Updated daily
 - *Latest version* only
- Statistics
 - 393 ontologies
 - 4.2M terms
 - 2419 different predicates
 - 80M triples



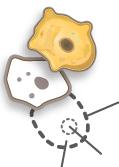
Resource Description Framework

- A collection of RDF statements represents a labeled, directed multi-graph
- RDF is modeled as a set of triples
- Triples are in the form (subject, predicate, object) where:
 - Subjects and predicates are URIs
 - Objects are either URIs or Literals
 - Literals can be typed and have a language tag
- Blank nodes (bnodes) or anonymous resources are nodes that do not have a URI or literal
- RDF is not XML, there are different serializations, e.g. N3, turtle, rdf/xml



RDF – Turtle format

```
@prefix rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#> .  
@prefix rdfs: <http://www.w3.org/2000/01/rdf-schema#> .  
@prefix owl: <http://www.w3.org/2002/07/owl#> .  
@prefix xsd: <http://www.w3.org/2001/XMLSchema#> .  
  
<http://who.int/bodysystem.owl#NervousSystem>  
    rdf:type owl:Class ;  
    rdfs:label "Nervous System"^^xsd:string ;  
    rdfs:subClassOf <http://who.int/bodysystem.owl#BodySystem> .  
  
<http://who.int/bodysystem.owl#BodySystem>  
    rdf:type owl:Class ;  
    rdfs:label "Body System"^^xsd:string .
```



RDF – N3 format

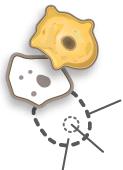
```
#(each triple block has to be in one line, splitted to fit in screen)
<http://who.int/bodysystem.owl#BodySystem>
<http://www.w3.org/1999/02/22-rdf-syntax-ns#type>
<http://www.w3.org/2002/07/owl#Class> .

<http://who.int/bodysystem.owl#BodySystem>
<http://www.w3.org/2000/01/rdf-schema#label>
"Body System"^^<http://www.w3.org/2001/XMLSchema#string> .

<http://who.int/bodysystem.owl#NervousSystem>
<http://www.w3.org/1999/02/22-rdf-syntax-ns#type>
<http://www.w3.org/2002/07/owl#Class> .

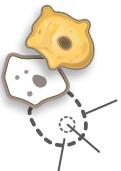
<http://who.int/bodysystem.owl#NervousSystem>
<http://www.w3.org/2000/01/rdf-schema#label>
"Nervous System"^^<http://www.w3.org/2001/XMLSchema#string> .

<http://who.int/bodysystem.owl#NervousSystem>
<http://www.w3.org/2000/01/rdf-schema#subClassOf>
<http://who.int/bodysystem.owl#BodySystem> .
```



RDF – RDF/XML format

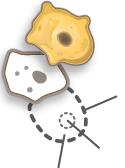
```
<?xml version="1.0" encoding="utf-8"?>
<rdf:RDF xmlns:rdf="http://www.w3.org/1999/02/22-rdf-syntax-ns#">
  <ns1:Class xmlns:ns1="http://www.w3.org/2002/07/owl#"
    rdf:about="http://who.int/bodysystem.owl#BodySystem">
    <ns2:label xmlns:ns2="http://www.w3.org/2000/01/rdf-schema#"
      rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Body System</ns2:label>
  </ns1:Class>
  <ns3:Class xmlns:ns3="http://www.w3.org/2002/07/owl#"
    rdf:about="http://who.int/bodysystem.owl#NervousSystem">
    <ns4:label xmlns:ns4="http://www.w3.org/2000/01/rdf-schema#"
      rdf:datatype="http://www.w3.org/2001/XMLSchema#string">Nervous System</ns4:label>
    <ns5:subClassOf xmlns:ns5="http://www.w3.org/2000/01/rdf-schema#"
      rdf:resource="http://who.int/bodysystem.owl#BodySystem"/>
  </ns3:Class>
</rdf:RDF>
```



SPARQL

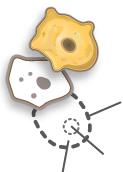
- W3C standard query language for RDF
- Structure

```
# prefixes, optional
PREFIX owl: <http://www.w3.org/2002/07/owl#>
...
# result clause {SELECT, DESCRIBE, CONSTRUCT, ASK}
SELECT ...
# dataset definition, optional
FROM ...
# query pattern(s)
WHERE {
    ...
}
# query modifiers, optional
ORDER BY ...
```



BioPortal Metadata

- Virtual ontology identifier
 - Stable identifier across all versions of the ontology
 - All versions of an ontology are linked via this ID
- Ontology version identifier
 - Unique for each ontology version
 - Most metadata linked directly to the ontology version



Virtual Ontology

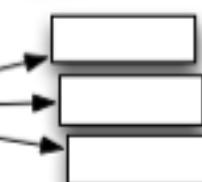
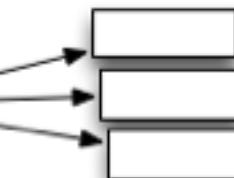
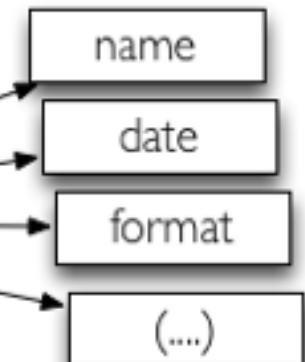


meta:hasVersion

Versions



ontology
/46896



```
PREFIX meta: <http://bioportal.bioontology.org/metadata/def/>
SELECT DISTINCT ?graph
WHERE {
    [ ] meta:hasDataGraph ?graph
}
ORDER BY ?graph
```

graph

<http://bioportal.bioontology.org/ontologies/3.0_OWL>

<http://bioportal.bioontology.org/ontologies/AAO>

<http://bioportal.bioontology.org/ontologies/ABA> ↗

<<http://bioportal.bioontology.org/ontologies/ACGT>> 

<http://bioportal.bioontology.org/ontologies/ADW>

<http://bioportal.bioontology.org/ontologies/AEO_x1> ↗

<http://bioportal.bioontology.org/ontologies/AERO_x1>

<<http://bioportal.bioontology.org/ontologies/AIR>> ↗

<<http://bioportal.bioontology.org/ontologies/AMINO-ACID>> 

<http://bioportal.bioontology.org/ontologies/AMINO_ACID-ZH_CN> 

<http://bioportal.bioontology.org/ontologies/

Select ontology abbreviations

@prefix rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns# .	ONTOLOGY ID:	1353
@prefix xsd: <http://www.w3.org/2001/XMLSchema# .	BIOPORTAL PURL:	http://purl.bioontology.org/ontology/SNOMEDCT
@prefix foaf: <http://xmlns.com/foaf/0.1# .	STATUS:	
@prefix rdfs: <http://www.w3.org/2000/01/rdf-schema# .	FORMAT:	UMLS
@prefix dc: <http://purl.org/dc/terms# .	CATEGORIES:	Health
@prefix dct: <http://purl.org/dc/terms# .	GROUPS:	Unified Medical Language System WHO Family of International Classifications Cancer Biomedical Informatics Grid
@prefix owl: <http://www.w3.org/2002/07/owl# .	CONTACT:	Vivian A. Auld, auld@nlm.nih.gov
@prefix bio: <http://purl.org/bio/ontology# .	HOME PAGE:	http://ihtsdo.org
@prefix meta: <http://ihtsdo.org/ontologies/meta# .	PUBLICATIONS PAGE:	http://ihtsdo.org
@prefix graphs: <http://ihtsdo.org/ontologies/graphs# .	DOCUMENTATION PAGE:	http://ihtsdo.org
@prefix omv: <http://ihtsdo.org/ontologies/omv# .	DESCRIPTION:	SNOMED Clinical Terms

```

<http://bioportal.bioontology.org/ontologies/1353> .
| meta:administeredBy "National Library of Medicine" ;
| meta:codingScheme "SNOMED Clinical Terms" ;
| meta:fileNames "#1353" ;
| meta:filePath "/1353" ;
| meta:hasContactEmail "Vivian.A.Auld@nlm.nih.gov" ;
| meta:hasContactName "Vivian A. Auld" ;
| meta:hasDataGraph "http://ihtsdo.org/ontologies/1353" ;
| meta:id 46896 ;
| meta:internalVersion "1.0" ;
| meta:isFlat false ;
| meta:isFoundry false ;
| meta:isMetadataOnly false ;
| meta:isRemote false ;
| meta:isVersionOfVirtualOntology <http://bioportal.bioontology.org/ontologies/1353> ;
| meta:statusID 3 ;
| meta:targetTerminologies "SNOMEDCT"^^xsd:string ;
| meta:timestampCreation "2012-02-13T13:47:00"^^xsd:dateTime ;
| meta:urlHomepage "http://ihtsdo.org"^^xsd:string ;
| meta:urlPublications "http://ihtsdo.org"^^xsd:string ;
| omv:acronym "SNOMEDCT"^^xsd:string ;
| omv:creationDate "2011-07-31T00:00:00"^^xsd:datetime ;
| omv:description "SNOMED Clinical Terms"^^xsd:string ;
| omv:documentation "http://ihtsdo.org"^^xsd:string ;
| omv:lastUpdate "2012-02-13T13:47:00"^^xsd:dateTime ;
| omv:version "1.0"^^xsd:string .

```

```

PREFIX xsd: <http://www.w3.org/2001/XMLSchema#>
PREFIX meta: <http://bioportal.bioontology.org/metadata/def/>
PREFIX omv: <http://omv.ontoware.org/2005/05/ontology#>

SELECT ?version ?name ?creation WHERE {
  ?version a omv:Ontology .
  ?version omv:name ?name .
  ?version meta:timestampCreation ?creation .
  FILTER (xsd:dateTime(?creation) > "2012-01-01T00:00:00"^^xsd:dateTime)
} ORDER BY ?name

```

version	name	creation
< http://bioportal.bioontology.org/ontologies/47573 >	"Adverse Event Reporting ontology"	"2012-05-31T18:06:33"
< http://bioportal.bioontology.org/ontologies/47302 >	"Adverse Event Reporting ontology"	"2012-04-25T18:06:40"
< http://bioportal.bioontology.org/ontologies/47035 >	"Adverse Event Reporting ontology"	"2012-03-09T18:08:24"
< http://bioportal.bioontology.org/ontologies/47022 >	"Adverse Event Reporting ontology"	"2012-03-07T18:07:48"
< http://bioportal.bioontology.org/ontologies/47058 >	"Adverse Event Reporting ontology"	"2012-03-13T18:10:57"
< http://bioportal.bioontology.org/ontologies/47432 >	"Adverse Event Reporting ontology"	"2012-05-11T18:06:38"
< http://bioportal.bioontology.org/ontologies/47523 >	"Adverse Event Reporting ontology"	"2012-05-23T18:06:28"
< http://bioportal.bioontology.org/ontologies/47581 >	"Anatomical Entity Ontology"	"2012-06-01T18:06:44"
< http://bioportal.bioontology.org/ontologies/46991 >	"Ascomycete phenotype ontology"	"2012-03-01T18:05:58"
< http://bioportal.bioontology.org/ontologies/47544 >	"BOOK"	"2012-05-27T19:19:21"
< http://bioportal.bioontology.org/ontologies/47507 >	"BRENDA tissue / enzyme source"	"2012-05-22T18:03:00"

All ontologies updated since *DATE*

```
PREFIX owl: <http://www.w3.org/2002/07/owl#>
PREFIX rdfs: <http://www.w3.org/2000/01/rdf-schema#>

SELECT DISTINCT ?graph WHERE {
GRAPH ?graph {
    <http://mouse.brain-map.org/atlas/index.html#RHP> a owl:Class .
}
}
```

graph

- <http://bioportal.bioontology.org/ontologies/H1_NMOABA_x2> ↗
- <<http://bioportal.bioontology.org/ontologies/NMOABA>> ↗
- <http://bioportal.bioontology.org/ontologies/H1_NMOABA> ↗
- <http://bioportal.bioontology.org/ontologies/H1_NMOABA_x3> ↗
- <http://bioportal.bioontology.org/ontologies/H1_NMOABA_x1> ↗
- <<http://bioportal.bioontology.org/ontologies/ABA>> ↗

Find term in all ontologies

```

PREFIX owl: <http://www.w3.org/2002/07/owl#>
PREFIX rdfs: <http://www.w3.org/2000/01/rdf-schema#>
PREFIX rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>

SELECT *
FROM <http://bioportal.bioontology.org/ontologies/ABA>
WHERE {
    ?s rdf:type owl:Class .
    ?s rdfs:label ?label .
}

```

s	label
<http://mouse.brain-map.org/atlas/index.html#Alp6b>	"Agranular insular area, posterior
<http://mouse.brain-map.org/atlas/index.html#BA>	"Bed nucleus of the accessory olfactory tract"
<http://mouse.brain-map.org/atlas/index.html#NTSm>	"Nucleus of the solitary tract, medial
<http://mouse.brain-map.org/atlas/index.html#CA2>	"Field CA2"
<http://mouse.brain-map.org/atlas/index.html#LD>	"Lateral dorsal nucleus of thalamus"
<http://mouse.brain-map.org/atlas/index.html#SBPV>	"Subparaventricular zone"
<http://mouse.brain-map.org/atlas/index.html#SSp-II5>	"Primary somatosensory area, low
<http://mouse.brain-map.org/atlas/index.html#Alv6a>	"Agranular insular area, ventral par-
<http://mouse.brain-map.org/atlas/index.html#Alv6b>	"Agranular insular area, ventral par-
<http://mouse.brain-map.org/atlas/index.html#BA1>	"Bed nucleus of the accessory olfactory tract"
<http://mouse.brain-map.org/atlas/index.html#CA1>	"Field CA1"

Select all terms from the
ABA Adult Mouse Anatomy

```

PREFIX owl: <http://www.w3.org/2002/07/owl#>
PREFIX skos: <http://www.w3.org/2004/02/skos/core#>
SELECT DISTINCT ?termURI ?prefLabel
  FROM <http://bioportal.bioontology.org/ontologies/EHDA>
  FROM <http://bioportal.bioontology.org/ontologies/globals>
WHERE {
    ?termURI a owl:Class;
    skos:prefLabel ?prefLabel .
}

```

termURI	prefLabel
<http://purl.obolibrary.org/obo/EHDA_6843> ↗	"endodermal epithelium"
<http://purl.obolibrary.org/obo/EHDA_6634> ↗	"lateral wall"
<http://purl.obolibrary.org/obo/EHDA_634> ↗	"cavity"
<http://purl.obolibrary.org/obo/EHDA_8119> ↗	"genital tubercle"
<http://purl.obolibrary.org/obo/EHDA_8816> ↗	"medulla oblongata"
<http://purl.obolibrary.org/obo/EHDA_6481> ↗	"right part"
<http://	

Select URI and preferred label from all terms

```

PREFIX owl: <http://www.w3.org/2002/07/owl#>
PREFIX rdfs: <http://www.w3.org/2000/01/rdf-schema#>

SELECT DISTINCT ?parent ?label WHERE {
GRAPH <http://bioportal.bioontology.org/ontologies/ABA> {
    <http://mouse.brain-map.org/atlas/index.html#RHP> rdfs:subClassOf ?parent .
    ?parent rdfs:label ?label .
}
}

```

parent

label

http://mouse.brain-map.org/atlas/index.html#HPF	HPF
"Hippocampal formation"	

Preferred Name	Retrohippocampal region
ID	RHP
Full Id	http://mouse.brain-map.org/atlas/index.html#RHP
Is Defined By	http://mouse.brain-map.org/atlas/coronal/RHP.html
Label	Retrohippocampal region
Disjoint With	Hippocampal region
Sub Class Of	Hippocampal formation

Get parent of given term

```

PREFIX owl: <http://www.w3.org/2002/07/owl#>
PREFIX rdfs: <http://www.w3.org/2000/01/rdf-schema#>
PREFIX rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>

SELECT *
FROM <http://bioportal.bioontology.org/ontologies/BODYSYSTEM>
WHERE {
    ?s rdf:type owl:Class .
    ?s rdfs:label ?label .
    OPTIONAL { ?s rdfs:subClassOf ?parent . }
}

```

		System"	
<http://who.int/bodysystem.owl#DigestiveSystem> ↗	"Digestive System"	<http://who.int/bodysys	
<http://who.int/bodysystem.owl#AutonomicNervousSystem> ↗	"Autonomic Nervous System"	<http://who.int/bodysys	
<http://who.int/bodysystem.owl#PeripheralNervousSystem> ↗	"Peripheral Nervous System"	<http://who.int/bodysys	
<http://who.int/bodysystem.owl#BodySystem> ↗	"Body System" ...		
<http://who.int/bodysystem.owl#VisualSystem> ↗	"Visual System" <http://who.int/bodysys		
<http://who.int/bodysystem.owl#UrinarySystem> ↗	"Urinary System" <http://who.int/bodysys		
<http://who.int/bodysystem.owl#FemaleGenitalSystem> ↗	"Female Genital System" <http://who.int/bodysys		
<http://who.int/bodysystem.owl#MetabolicSystem> ↗	"Metabolic" <http://who.int/bodysys		

Select all terms and their parent

```
SELECT DISTINCT ?p
FROM <http://bioportal.bioontology.org/ontologies/SNOMEDCT>
WHERE {
    [] ?p [] .
}
```

P

- <<http://bioportal.bioontology.org/ontologies/umls/hasSTY>> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/focus_of> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/procedure_context_of> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/has_specimen_source_morphology> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/has_scale_type> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/indirect_device_of> ↗
- <<http://purl.bioontology.org/ontology/SNOMEDCT/SUBSETMEMBER>> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/uses_access_device> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/uses_energy> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/has_associated_morphology> ↗
- <http://purl.bioontology.org/ontology/SNOMEDCT/has_intent> ↗

Select distinct properties

```
SELECT DISTINCT ?p WHERE {  
  GRAPH <http://bioportal.bioontology.org/ontologies/ABA> {  
    <http://mouse.brain-map.org/atlas/index.html#RHP> ?p ?o  
  }  
}
```

P

<<http://www.w3.org/2000/01/rdf-schema#isDefinedBy>> ↗

<<http://www.w3.org/2000/01/rdf-schema#subClassOf>> ↗

<<http://www.w3.org/1999/02/22-rdf-syntax-ns#type>> ↗

<<http://bioportal.bioontology.org/metadata/def/prefLabel>> ↗

<<http://www.w3.org/2002/07/owl#disjointWith>> ↗

<<http://www.w3.org/2000/01/rdf-schema#label>> ↗

Select properties for term

```
PREFIX owl: <http://www.w3.org/2002/07/owl#>
SELECT (COUNT(?term) as ?c)
FROM <http://bioportal.bioontology.org/ontologies/SNOMEDCT>
WHERE { ?term a owl:Class. }
```

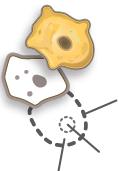
c

"395036"

Metrics ?

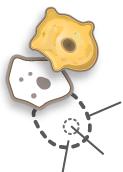
NUMBER OF CLASSES:	395036
NUMBER OF INDIVIDUALS:	0
NUMBER OF PROPERTIES:	41
MAXIMUM DEPTH:	32
MAXIMUM NUMBER OF SIBLINGS:	10010
AVERAGE NUMBER OF SIBLINGS:	1
CLASSES WITH A SINGLE SUBCLASS:	29002
CLASSES WITH MORE THAN 25 SUBCLASSES:	2053
CLASSES WITH NO DEFINITION:	394352

Count terms in SNOMED



Performance Tips and Tricks

- Completely unbound patterns (`?g ?s ?p ?o`) are not allowed
- To optimize queries, use UNIONs instead of FILTERs
- If using FILTER on literals it is better if the filter is not applied to millions of rows
- To prevent combinatorial explosions of results, consider use CONSTRUCT or DESCRIBE (any M-N relationship can provoke this)

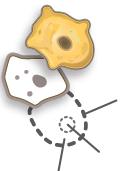


SPARQL Code Repository

- <https://github.com/ncbo/sparql-code-examples>

[sparql-code-examples](#) /

name	age	message
java	4 months ago	Update java/src/org/ncbo/stanford/sparql/examples/SimpleTest.java [msalvadores]
javascript	2 months ago	Simplified in-browser SPARQL call [Josh Mandel]
perl	4 months ago	sparql.bioontology.org/sparql is now the production SPARQL endpoint [msalvadores]
python	2 months ago	Eliminate references to alphasparql [Josh Mandel]
ruby	4 months ago	Ruby examples for querying the BioPortal SPARQL endpoint [palexander]
.gitignore	9 months ago	Javascript, Java, Perl, Python examples [msalvadores]
COPYING	9 months ago	README and LICENSE [msalvadores]
README	2 months ago	Eliminate references to alphasparql [Josh Mandel]



BioPortal SPARQL Endpoint

- Documentation:

http://www.bioontology.org/wiki/index.php/SPARQL_BioPortal

- Query interface: <http://sparql.bioontology.org/>

- Example queries:

<http://sparql.bioontology.org/examples>

- Sample code:

<https://github.com/ncbo/sparql-code-examples>