

# Aplicación de la Web Semántica en Biología Molecular

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<http://tinyurl.com/2u6vhqe>



Información en Biología Molecular

Life Sciences Semantic Web (LSSW)

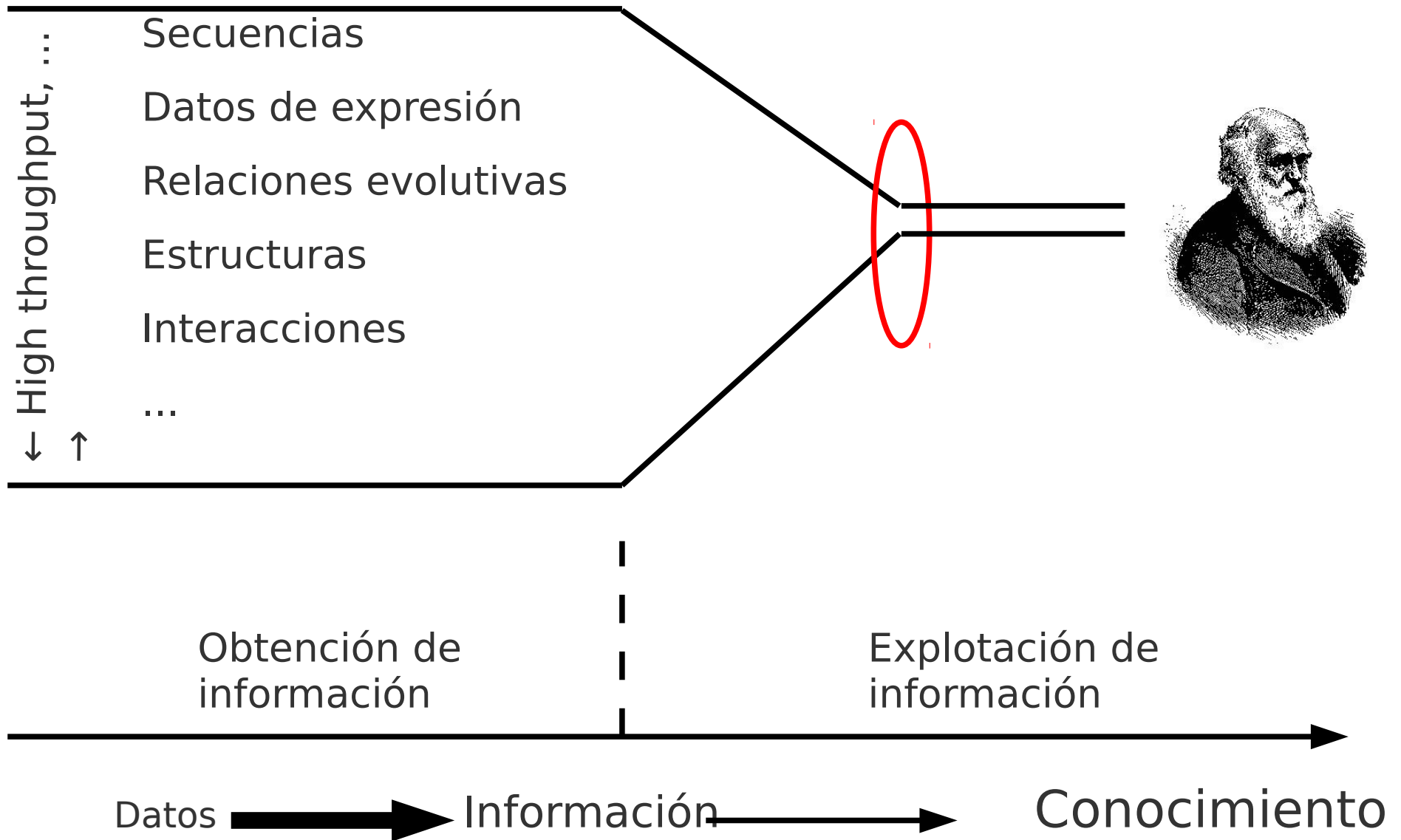
Open Biological and Biomedical Ontologies -OBO- (!  
LSSW)

¿Por qué no se aplica la LSSW?

Conclusiones

# **Información en Biología Molecular**

# Información en Biología Molecular



## Información en Biología Molecular

... por la naturaleza misma de la información biológica

Compleja (¡Y no se puede abstraer!)

Cambiante

Producida por muchos agentes diferentes

Grandes volúmenes

Crecimiento acelerado (ej. high throughput)

... por cómo hemos representado la información biológica

Diferentes recursos, diferentes esquemas

Crisis de identidad

Para humanos, no para máquinas (Anotaciones, literatura, ...  
)

...

## Información en Biología Molecular

“It is quite depressive to think that we are spending millions in grants for people to perform experiments, produce new knowledge, hide this knowledge in a often badly written text and then spend some more millions trying to second guess what the authors really did and found”

Teresa K. Attwood, Douglas B. Kell, Philip McDermott, James Marsh, Steve R. Pettifer, and David Thorne. Calling international rescue: knowledge lost in literature and data landslide! *The Biochemical journal*, 424(3):317-333, December 2009.

## Información en Biología Molecular

Q708Y0 *Saccharomyces cerevisiae*

Ortólogos de Q708Y0 en *Schizosaccharomyces pombe*

Si algun ortólogo esta en el núcleo (¡O partes del núcleo!), las proteínas que interaccionan con él por fosforilación

Obtener los procesos de regulación en los que participan esas proteínas

¿Afecta alguno de esos procesos al ciclo celular?

¿Cuál?



# Información en Biología Molecular



.....  
.....  
..... PR 2 is located in A ...  
.....

..... A is part of B .....  
.....  
.....X has part Y .....  
M has parts only M and N ...  
.....  
.....F appears only in H .....  
.....

.....  
.....  
PR1 is ortholog of PR2 ...  
.....

.....  
.....  
..... B is part of C ...  
.....

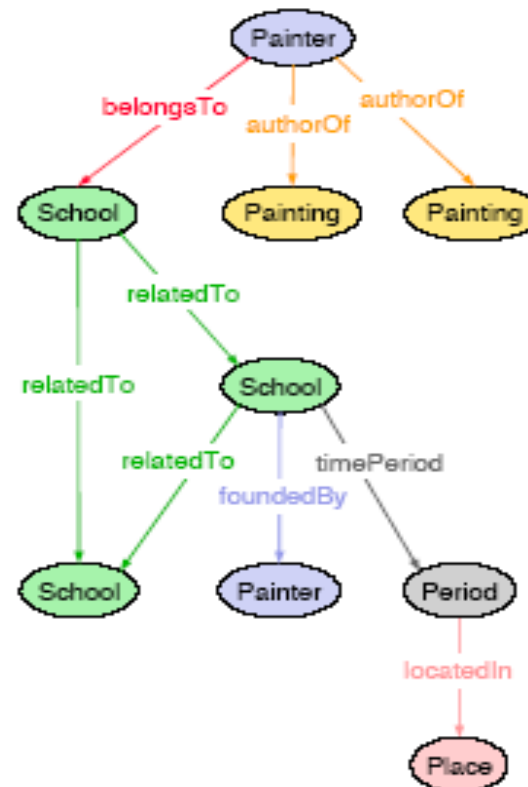
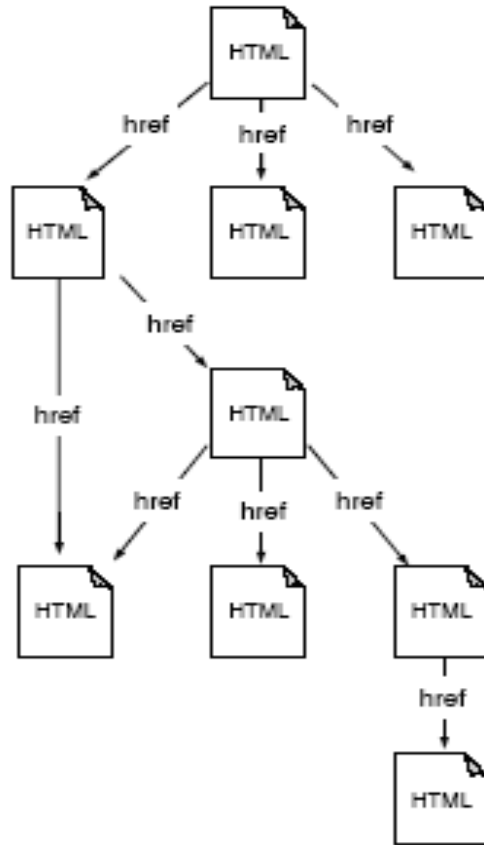
Grandes volúmenes de  
información desperdigados  
en la web



# **Life Sciences Semantic Web**

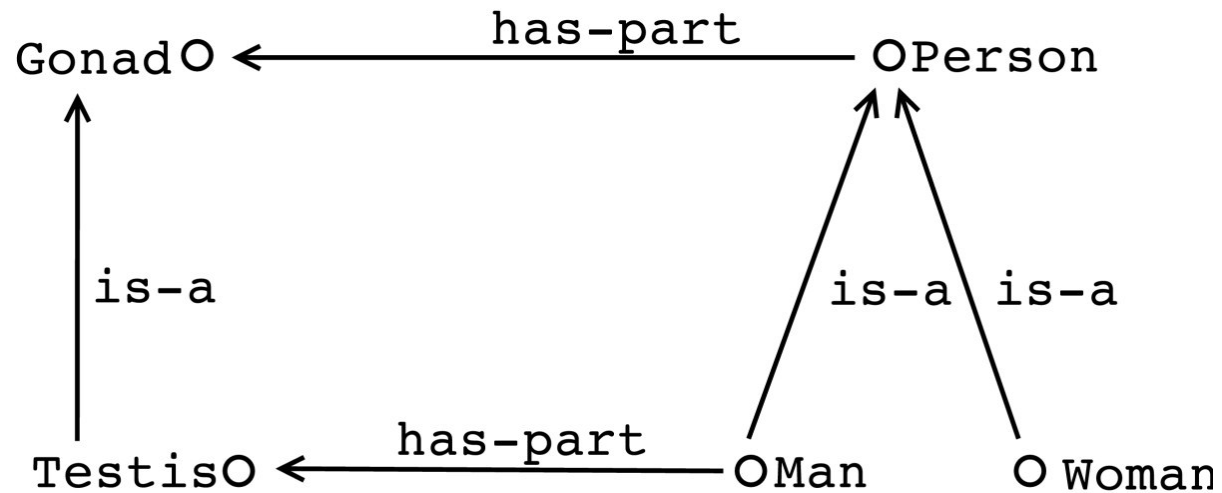
# Life Sciences Semantic Web

Web Semántica



## Life Sciences Semantic Web

### Ontología



## Life Sciences Semantic Web

Biología Molecular (Bioinformática)

Necesidades muy concretas de gestión de  
información

Muchos usuarios dispuestos a crear contenido  
semántico



“Test case” perfecto para la Web Semántica

W3C Health Care and Life Sciences Interest Group

<http://www.w3.org/blog/hcls>

## **Life Sciences Semantic Web**

Globally unique and resolvable names for biological entities

Consistent standards for data representation

Consistent standards for knowledge representation

Standard interface definitions for data retrieval and processing

Benjamin M. Good and Mark D. Wilkinson. The life sciences semantic web is full of creeps! *Brief Bioinform*, 7(3):275-286, September 2006.

## Life Sciences Semantic Web

Globally unique and resolvable names for biological entities:  
**LSID(?)**

Consistent standards for data representation: **RDF**

Consistent standards for knowledge representation: **OWL**

Standard interface definitions for data retrieval and processing: Semantic Web Services

Benjamin M. Good and Mark D. Wilkinson. The life sciences semantic web is full of creeps! *Brief Bioinform*, 7(3):275-286, September 2006.

## Life Sciences Semantic Web

### LSID (Life Sciences Identifiers): URNs

`urn:lsid:ipni.org:names:30000959-2`

### URIs

#### Bio2RDF (BANFF manifesto)

`http://tinyurl.com/39m9qru`

#### HCLS IG

`http://www.w3.org/2001/sw/hcls/notes/uris/`

#### Neurocommons

`http://neurocommons.org/page/URIs`

### Shared Names

`http://neurocommons.org/page/Shared_names`

...

## Life Sciences Semantic Web

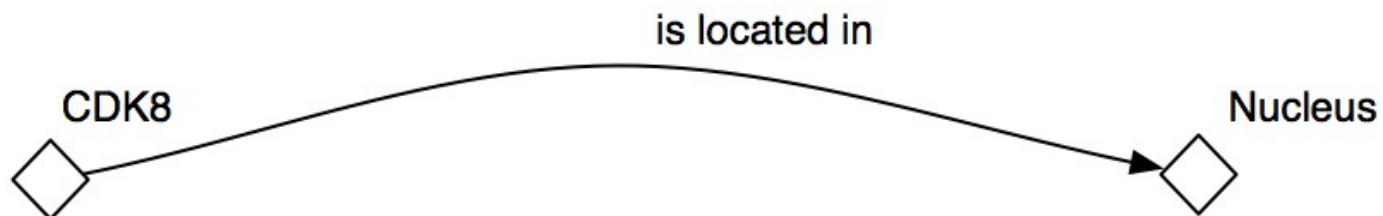
### RDF (Resource Description Framework)

Standard para representar datos en la WS

<http://www.w3.org/RDF/>

SPARQL para consultas

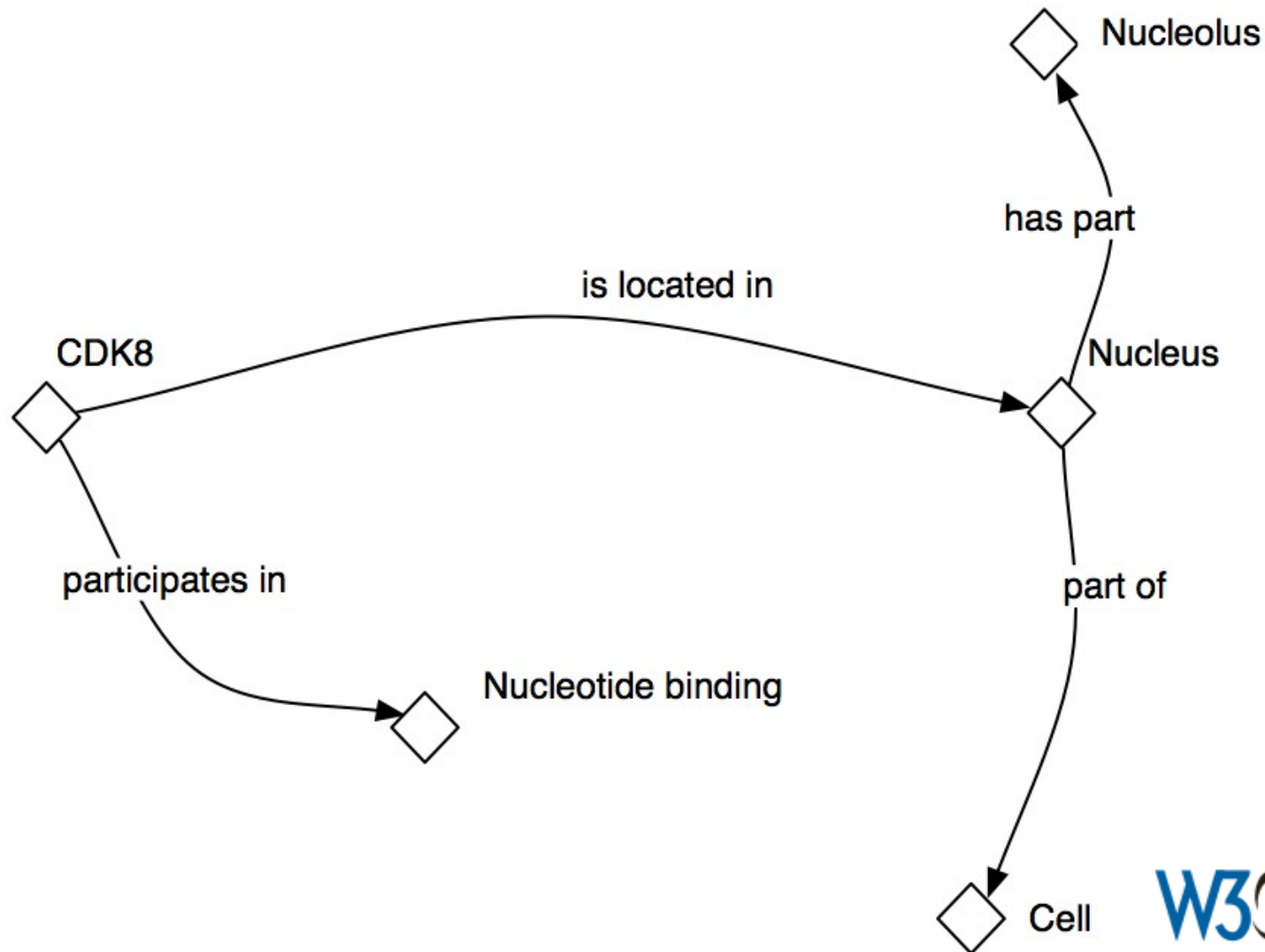
<http://www.w3.org/TR/rdf-sparql-query/>





# Life Sciences Semantic Web

RDF (Resource Description Framework)



# Life Sciences Semantic Web

Bio2RDF <http://bio2rdf.org/>

physical association, direct interaction between two molecules of uniprotkb:P05067-4|refseq:NP\_958817|entrezgene/locuslink:351 identified by anti tag coip, anti bait coip [irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE] at Bio2RDF

Search

[Find intranamespace links](#) [Find global links](#) Links Namespace

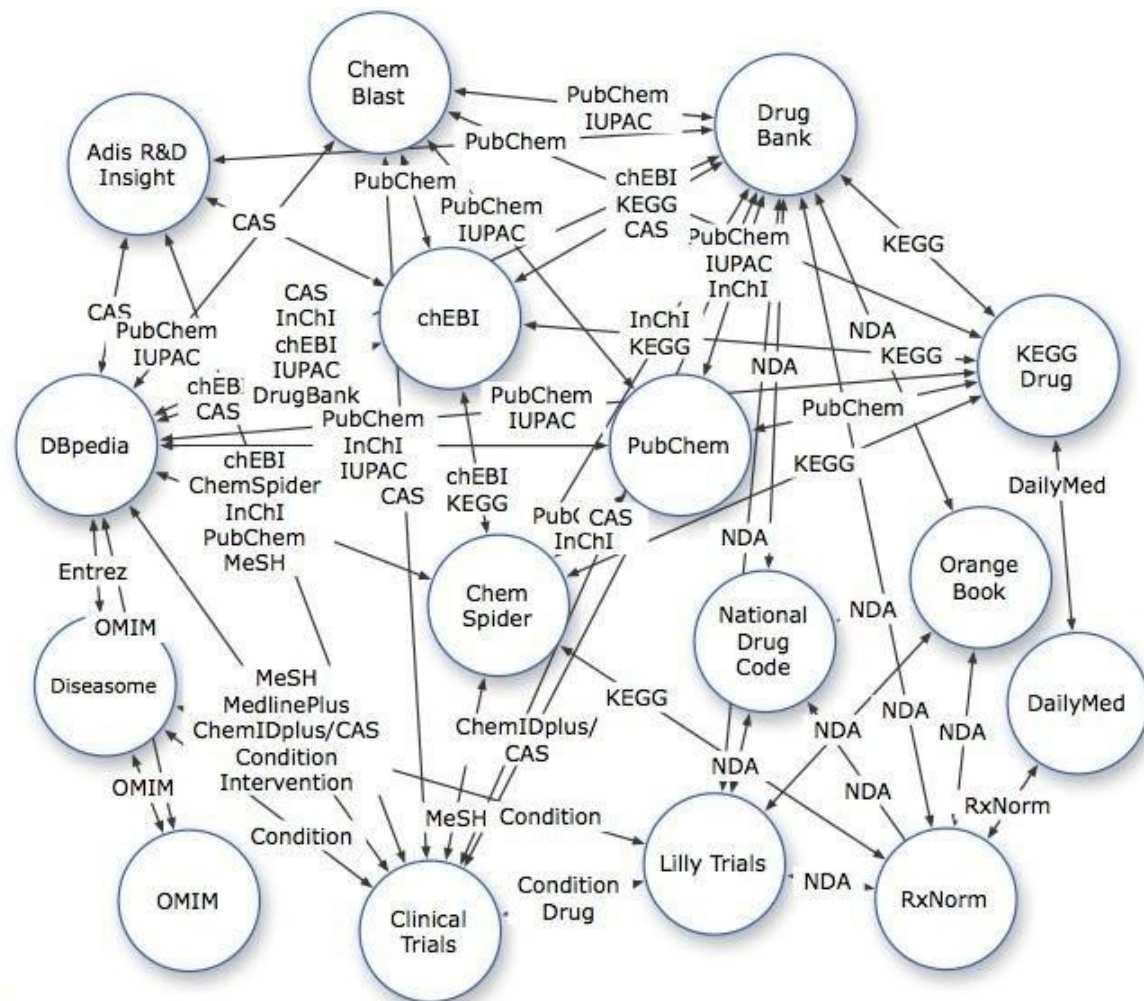
<http://bio2rdf.org/irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE>

Subject	Predicate	Object
<a href="http://bio2rdf.org/irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE">http://bio2rdf.org/irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE</a>	<a href="http://bio2rdf.org/irefindex:hpr">http://bio2rdf.org/irefindex:hpr</a>	10
	<a href="http://bio2rdf.org/irefindex:lpr">http://bio2rdf.org/irefindex:lpr</a>	10
	<a href="http://bio2rdf.org/irefindex:np">http://bio2rdf.org/irefindex:np</a>	1
	<a href="http://bio2rdf.org/irefindex:number_of_interactors">http://bio2rdf.org/irefindex:number_of_interactors</a>	2
	<a href="http://bio2rdf.org/irefindex:source">http://bio2rdf.org/irefindex:source</a>	<a href="http://bio2rdf.org/irefindex:2f6643738f03d32dc361ec3b513fd88c">http://bio2rdf.org/irefindex:2f6643738f03d32dc361ec3b513fd88c</a>
		<a href="http://bio2rdf.org/mi:0469">http://bio2rdf.org/mi:0469</a>
		<a href="http://bio2rdf.org/mi:0923">http://bio2rdf.org/mi:0923</a>
	<a href="http://bio2rdf.org/ns/bio2rdf:article">http://bio2rdf.org/ns/bio2rdf:article</a>	<a href="http://bio2rdf.org/pubmed:16193067">http://bio2rdf.org/pubmed:16193067</a>
	<a href="http://bio2rdf.org/ns/bio2rdf:method">http://bio2rdf.org/ns/bio2rdf:method</a>	<a href="http://bio2rdf.org/mi:0006">http://bio2rdf.org/mi:0006</a>
		<a href="http://bio2rdf.org/mi:0007">http://bio2rdf.org/mi:0007</a>
	<a href="http://creativecommons.org/ns#license">http://creativecommons.org/ns#license</a>	<a href="http://bio2rdf.org/license/irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE">http://bio2rdf.org/license/irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE</a>
	<a href="http://purl.org/dc/elements/1.1/identifier">http://purl.org/dc/elements/1.1/identifier</a>	irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE
	<a href="http://semanticscience.org/ontology/hasParticipant">http://semanticscience.org/ontology/hasParticipant</a>	<a href="http://bio2rdf.org/irefindex:q3lluSzu8RGEONqITBEVFyMpfaq9606">http://bio2rdf.org/irefindex:q3lluSzu8RGEONqITBEVFyMpfaq9606</a>
	<a href="http://www.w3.org/1999/02/22-rdf-syntax-ns#type">http://www.w3.org/1999/02/22-rdf-syntax-ns#type</a>	<a href="http://bio2rdf.org/mi:0407">http://bio2rdf.org/mi:0407</a>
		<a href="http://bio2rdf.org/mi:0915">http://bio2rdf.org/mi:0915</a>
		<a href="http://semanticscience.org/ontology/HomolInteraction">http://semanticscience.org/ontology/HomolInteraction</a> (External link)
	<a href="http://www.w3.org/2000/01/rdf-schema#label">http://www.w3.org/2000/01/rdf-schema#label</a>	physical association, direct interaction between two molecules of uniprotkb:P05067-4 refseq:NP_958817 entrezgene/locuslink:351 identified by anti tag coip, anti bait coip [irefindex:/sMGTLOhS3hkGyRkP0kAgNqG9qE]
	<a href="http://www.w3.org/2002/07/owl#sameAs">http://www.w3.org/2002/07/owl#sameAs</a>	<a href="http://bio2rdf.org/intact:EBI-1781246">http://bio2rdf.org/intact:EBI-1781246</a>
		<a href="http://bio2rdf.org/intact:EBI-1782329">http://bio2rdf.org/intact:EBI-1782329</a>
		<a href="http://bio2rdf.org/intact:EBI-2126823">http://bio2rdf.org/intact:EBI-2126823</a>

# Life Sciences Semantic Web

Linking Open Drug Data (LODD) <http://esw.w3.org/HCLSIG/LODD>

Ganador del triplify challenge <http://triplify.org/Challenge/2009>



# Life Sciences Semantic Web

BioGateway <http://www.semantic-systems-biology.org/biogateway>

The screenshot displays the BioGateway web interface. At the top, there is a search bar with the query "Get psoriasis proteins" and a LIMIT of 40. Below the search bar, the query is shown in SPARQL format:

```
SELECT distinct ?protein_name ?disease_description
  ?interacts_with ?encoded_by
WHERE {
  GRAPH <uniprot_sprot> {
    ?protein_id ssb:disease ?disease_description.
    ?protein_id ssb:mnemonic ?protein_name.
    OPTIONAL {
      ?protein_id ssb:interacts_with ?interactor.
      ?interactor ssb:mnemonic ?interacts_with.
      ?interactor ssb:encoded_by ?encoded_by.
    }
  }
  FILTER regex(?disease_description, 'psoriasis').
}
```

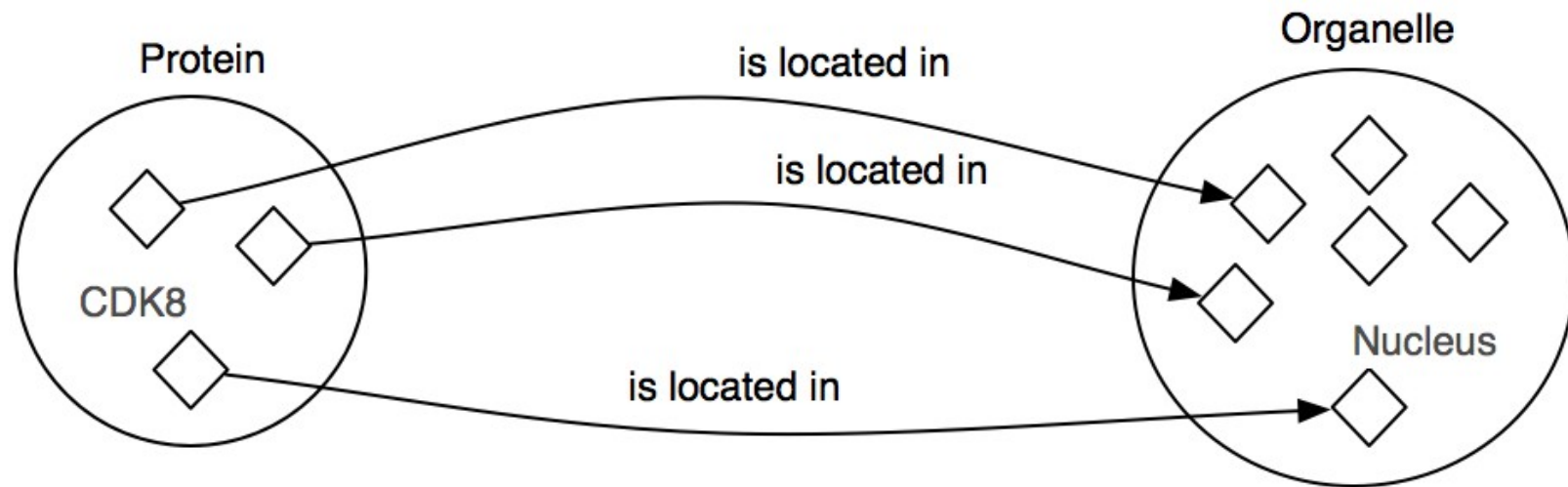
The results are displayed in a table with the following columns: protein\_name, disease\_description, interacts\_with, and encoded\_by. The first row shows the protein 1C06\_HUMAN, which is associated with the disease description "Genetic variation in HLA-C is associated with susceptibility to psoriasis 1 (PSORS1) [MIM%3A177900]. Psoriasis is a chronic inflammatory".

Below the table, a graph visualization shows the relationships between the proteins and diseases. The graph is centered on "Results" and shows connections to various entities like NALP1\_HUMAN, CASP5\_HUMAN, 1C06\_HUMAN, TNFB\_HUMAN, K1C17\_HUMAN, and CCDC85B. The graph is interactive, with sliders for "Degrees of Separation", "Scaling", and "Link Length", and a checkbox for "AutoFit".

Erick Antezana, Ward Blondé, Mikel Egaña, Alistair Rutherford, Robert Stevens, Bernard De Baets, Vladimir Mironov, Martin Kuiper. BioGateway a semantic systems biology tool for the life sciences. BMC bioinformatics 2009, 10(Suppl 10):S11

## Life Sciences Semantic Web

OWL (Web Ontology Language)



## Life Sciences Semantic Web

OWL (Web Ontology Language)

Autodescriptivo (self-descriptive)

datos + esquema en el mismo idioma

OWA (Open World Assumption)

el conocimiento en biología molecular es  
necesariamente incompleto

Semantica monotónica

~~UNA (Unique Name Assumption)~~

diferentes entradas en diferentes recursos se  
refieren a la misma entidad

## Life Sciences Semantic Web

OWL (Web Ontology Language)

Inferencia (“Reasoning”) completa y “eficiente”

Consultas

Clase-subclase

Individuo → clase

Consistencia

URIs para entidades

Editores (Protégé, TopBraid composer, ...)

APIs (OWL API, ...)

Razonadores (Pellet, FaCT++, Racer, ... )

## Life Sciences Semantic Web

### Uso de OWL

Vocabulario común

almacenar/integrar/reusar conocimiento

Inferir conocimiento no evidente

Clasificación de información

Consultas expresivas

Generación de hipótesis

Consistencia de la información

Al representar un dominio el razonador nos dice las  
contradicciones que cometemos: nos obliga a definirnos

Mantenimiento de conocimiento

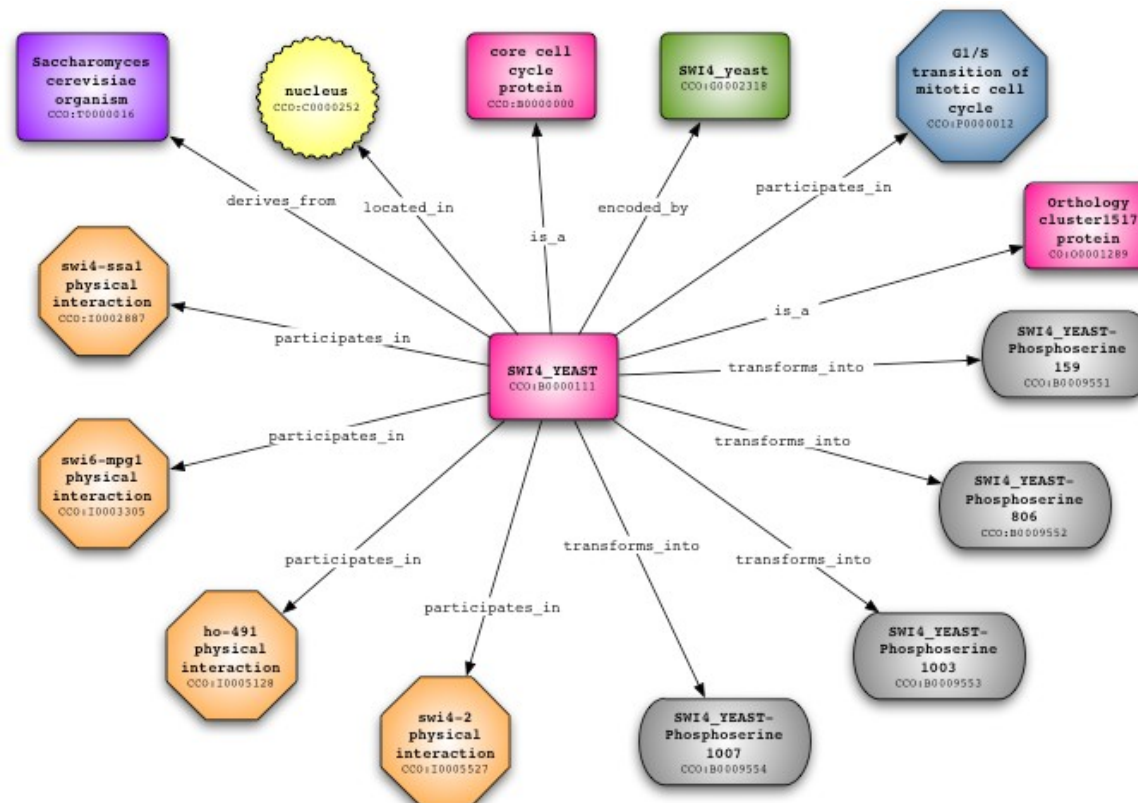




# Life Sciences Semantic Web

## Cell Cycle Ontology <http://www.semantic-systems-biology.org/cco>

Erick Antezana, Mikel Egaña, Ward Blondé, Aitzol Illarramendi, Iñaki Bilbao, Bernard De Baets, Robert Stevens, Vladimir Mironov, and Martin Kuiper. The cell cycle ontology: an application ontology for the representation and integrated analysis of the cell cycle process. *Genome Biology*, 10(5):R58+, 2009



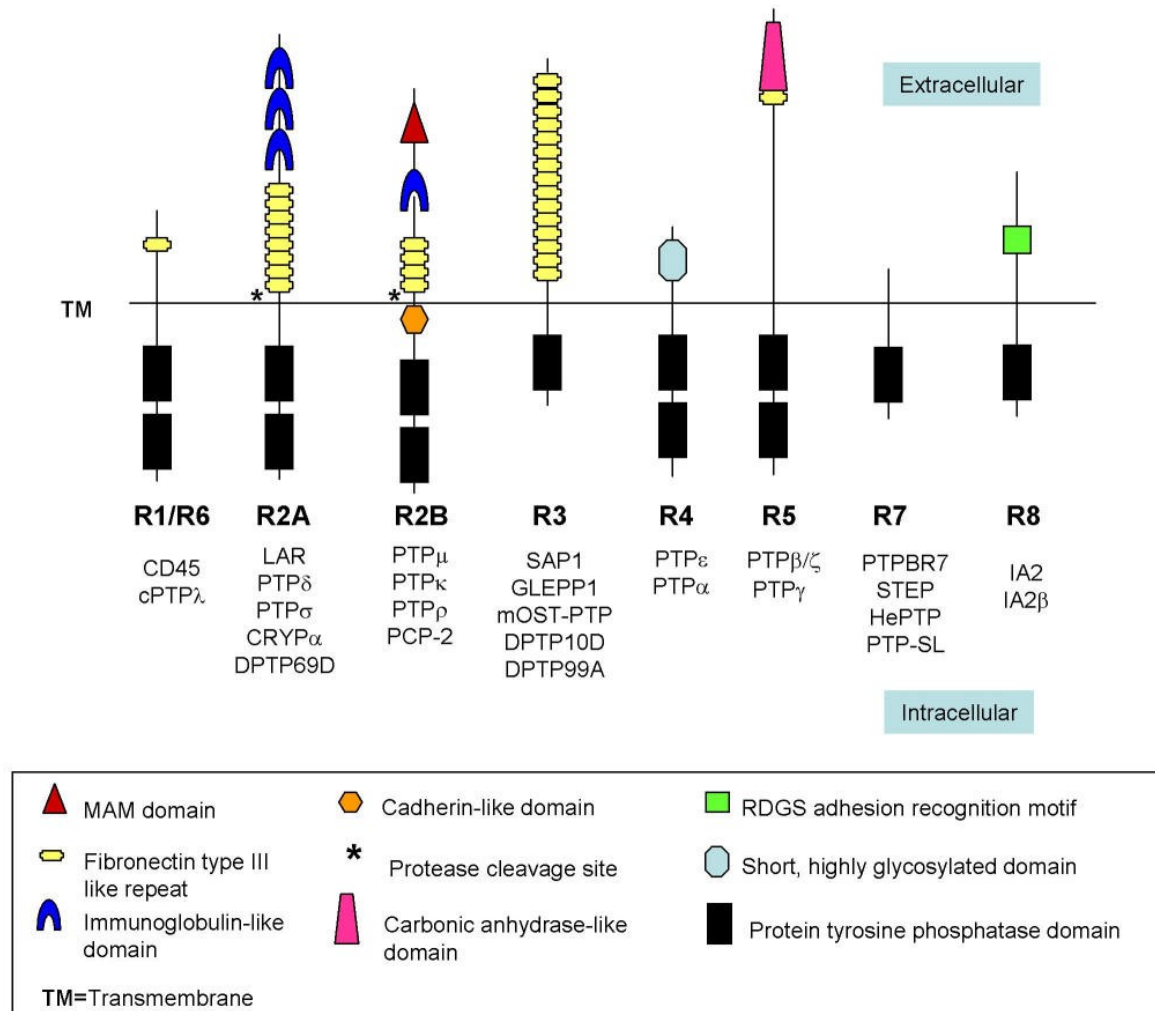
# Life Sciences Semantic Web

OBI <http://obi-ontology.org>

The screenshot displays the OBI ontology browser interface. On the left, the 'Asserted class hierarchy' for 'survival assessment' is shown, starting from 'Thing' and descending through 'entity', 'occurent', 'processual\_entity', 'planned\_process', 'investigation', and 'assay'. The 'survival assessment' class is highlighted. On the right, the 'Annotations' and 'Description' panels are visible. The 'Annotations' panel shows properties such as 'has\_curation\_status' (with value 'metadata incomplete'), 'editor\_preferred\_term' (with value 'survival assessment'), 'definition' (with value 'Survival assessment is an assay that measures the occurrence of death events in one or more organisms that are monitored over time'), 'editor\_note' (with value 'Need to point out more specifically that survival / death is measured.'), and 'label' (with value 'survival assessment'). The 'Description' panel shows 'Equivalent classes' (none listed), 'Superclasses' (including 'assay'), and 'Inferred anonymous superclasses' (none listed). The 'assay' superclass is expanded to show properties: 'has\_specified\_input some organism', 'has\_specified\_output some ('survival rate' and 'is about' some organism)', and 'achieves\_planned\_objective some 'assay objective''.

# Life Sciences Semantic Web

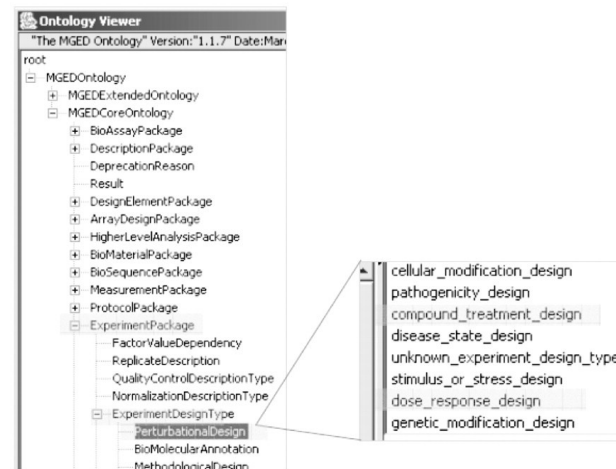
PhosphaBase <http://www.bioinf.manchester.ac.uk/phosphabase/>



## Life Sciences Semantic Web

MGED <http://mged.sourceforge.net/>

(a)



(b)

MiMiR

Introduction	Title	ExpType	Factors	Description	Users
Experiment	Category				Value
Compound	MO: PerturbationalDesign			dose_response_design	
Array	MO: PerturbationalDesign			compound_treatment_design	
Chip Type					

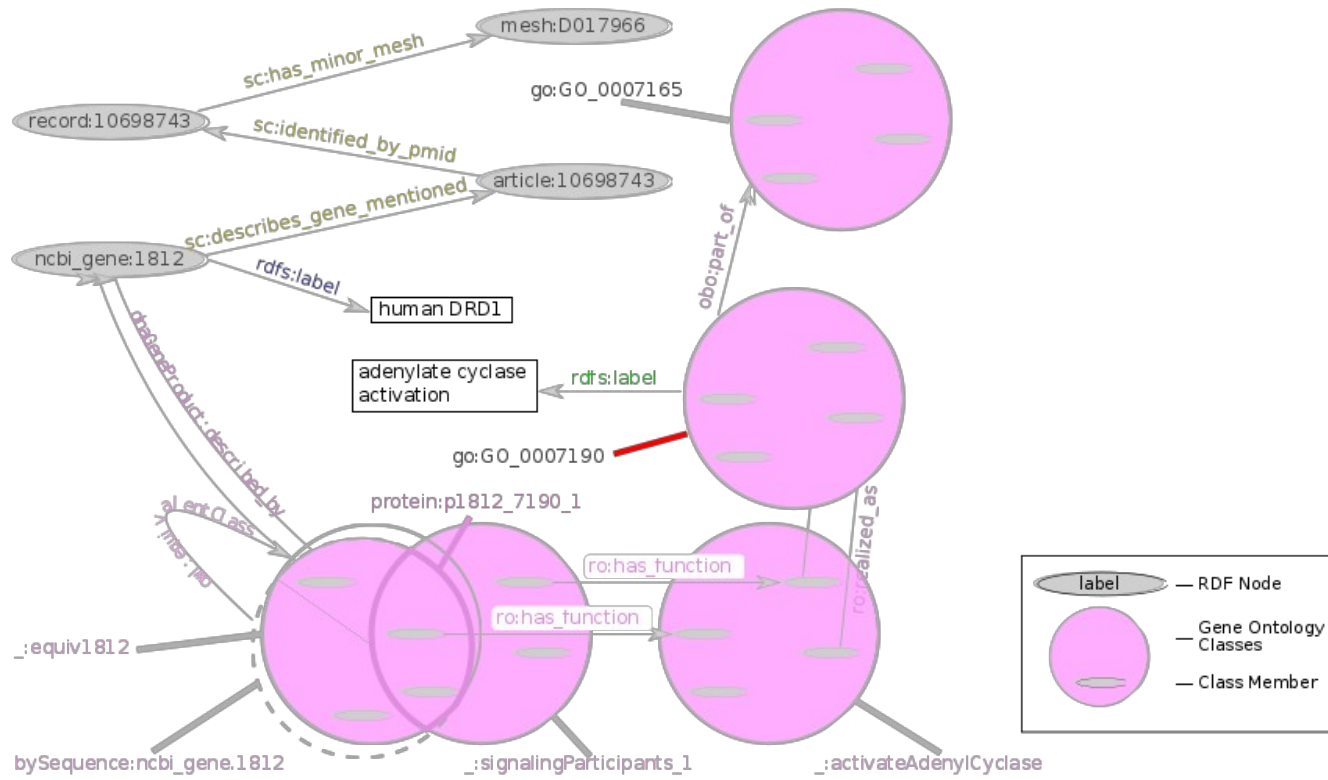
(c)

Introduction	BioSample	Ontology	Properties	Treatment	Aliquots
Experiment	Category				Value
Compound					
Array	MO: CellLine			NCI: Fibroblast	
Chip Type	MO: CellType			NCI: 3T3-L1 Cells	
BioSource					
BioSample					
Hybridisation					

# Life Sciences Semantic Web

## A prototype KB for the Life Sciences

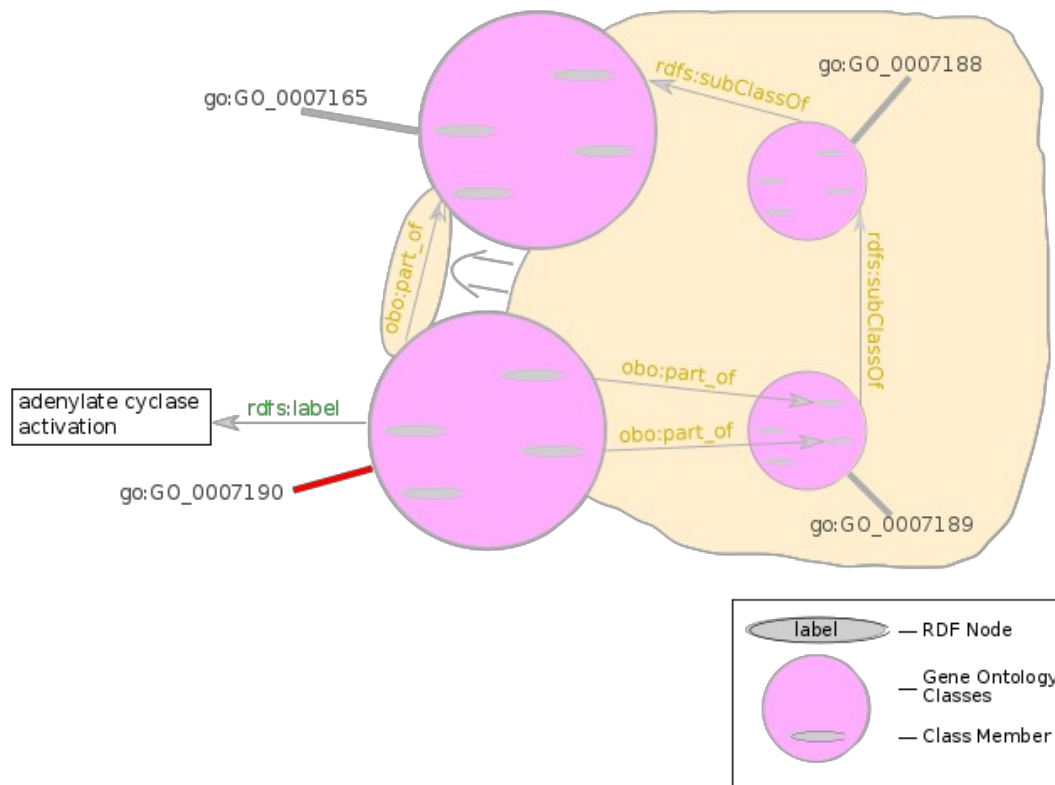
<http://www.w3.org/TR/hcls-kb/>



# Life Sciences Semantic Web

## A prototype KB for the Life Sciences

<http://www.w3.org/TR/hcls-kb/>



# **Open Biological and Biomedical Ontologies (OBO)**



# Open Biological and Biomedical Ontologies

OBO Foundry <http://www.obofoundry.org/>

Open

Common shared syntax

Unique identifier space

Versions

Delineated content

Definitions

OBO Relation Ontology








Well documented

Users

Collaboratively

# Open Biological and Biomedical Ontologies

## OBO Foundry

OBO Foundry ontologies				
Title	Domain	Prefix	File	Last changed
<a href="#">Biological process</a>	biological process	GO	<a href="#">gene_ontology_edit.obo</a> 	2010/08/29
<a href="#">Cellular component</a>	anatomy	GO	<a href="#">gene_ontology_edit.obo</a> 	2010/08/29
<a href="#">Chemical entities of biological interest</a>	biochemistry	CHEBI	<a href="#">chebi.obo</a> 	2010/08/03
<a href="#">Molecular function</a>	biological function	GO	<a href="#">gene_ontology_edit.obo</a> 	2010/08/29
<a href="#">Phenotypic quality</a>	phenotype	PATO	<a href="#">quality.obo</a> 	2010/08/19
<a href="#">PRotein Ontology (PRO)</a>	proteins	PRO	<a href="#">pro.obo</a> 	2010/08/20
<a href="#">Xenopus anatomy and development</a>	anatomy	XAO	<a href="#">xenopus_anatomy.obo</a>	2009/12/02
<a href="#">Zebrafish anatomy and development</a>	anatomy	ZFA	<a href="#">zebrafish_anatomy.obo</a> 	2010/08/06

OBO Foundry candidate ontologies and other ontologies of interest				
Title	Domain	Prefix	File	Last changed
<a href="#">Amphibian gross anatomy</a>	anatomy	AAO	<a href="#">amphibian_anatomy.obo</a>	2008/06/19
<a href="#">Amphibian taxonomy</a>	anatomy	ATO	<a href="#">amphibian_taxonomy.obo</a>	
<a href="#">Ascomycete phenotype ontology</a>	phenotype	APO	<a href="#">ascomycete_phenotype.obo</a>	2010/05/12

# Open Biological and Biomedical Ontologies

## Gene Ontology <http://geneontology.org/>

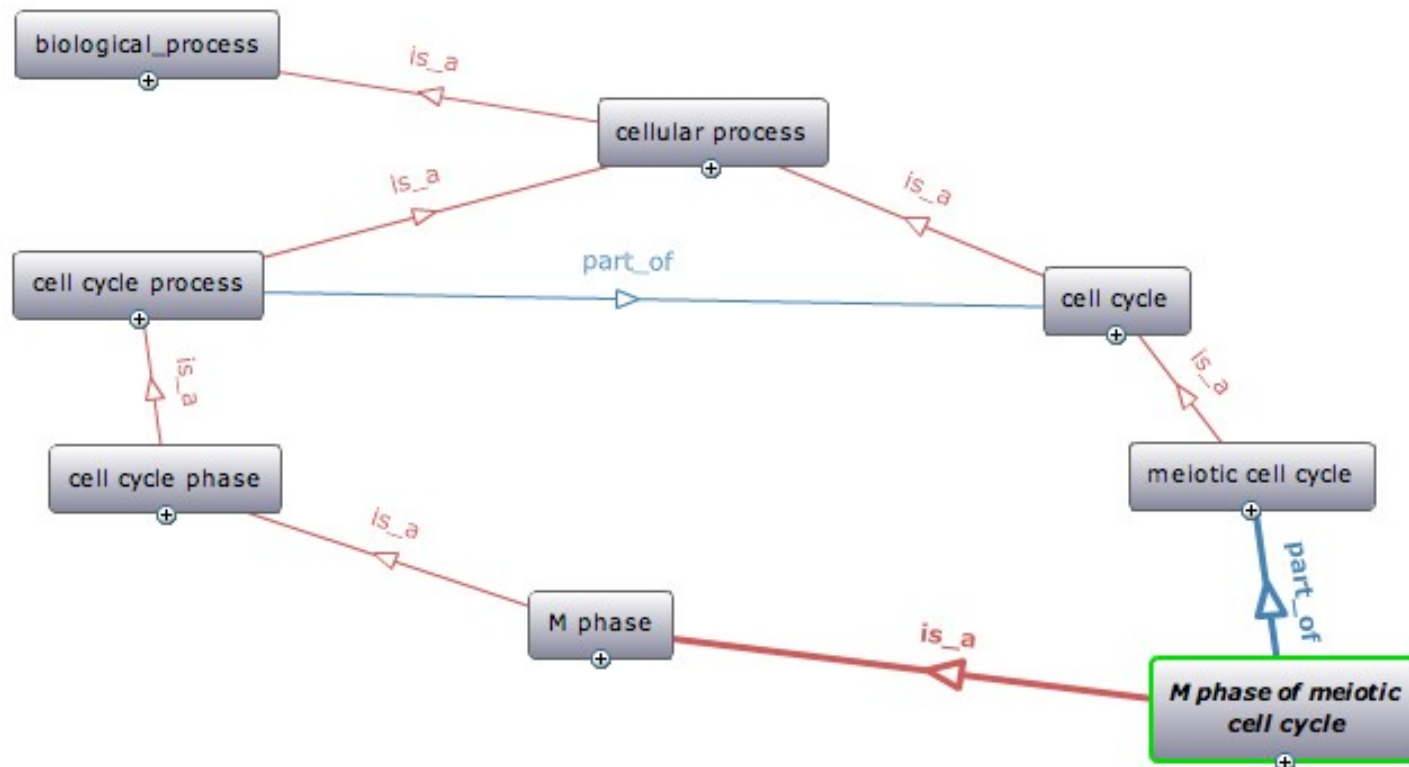
The Gene Ontology Consortium. Gene Ontology: tool for the unification of biology. *Nature Genet.* (2000) 25: 25-29

Michael Bada, Robert Stevens, Carole Goble, Yolanda Gil, Michael Ashburner, Judith A. Blake, J. Michael Cherry, Midori Harris, and Suzanna Lewis. A Short Study on the Success of the Gene Ontology . *Web Semantics: Science, Services and Agents on the World Wide Web*, 1(2):235-240, 2004.

Vocabulario controlado para describir la función molecular, el componente celular y el proceso biológico de genes (“Gene Products”)

## Open Biological and Biomedical Ontologies

Gene Ontology: ~ 32.500 términos en una estructura *is\_a*,  
*part\_of*, *regulates (+,-)*

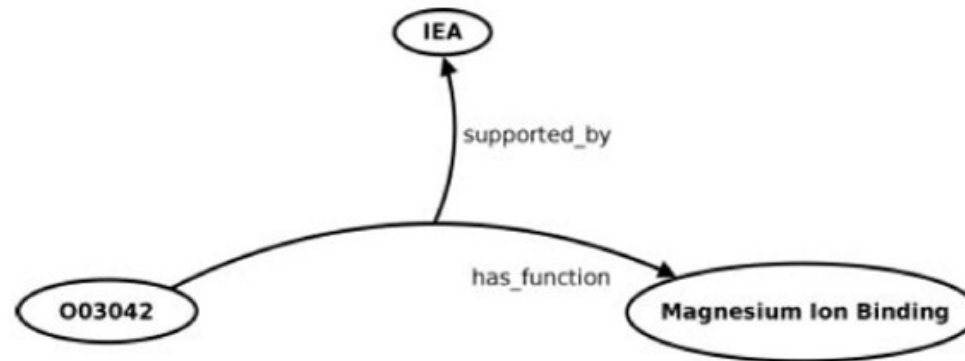


# Open Biological and Biomedical Ontologies

## Gene Ontology

Integración de recursos (anotaciones)

Gene Ontology Annotation (GOA)



## Explotación de la estructura

Ontological analysis of gene expression data: current tools, limitations, and open problems. *Bioinformatics*. 2005 Sep 15;21(18):3587-95. Epub 2005 Jun 30.

## **Open Biological and Biomedical Ontologies**

Otras ontologías importantes: ChEBI, Cell Type, Sequence Ontology, Phenotype Ontology, UberOntology, ...

“Meta Ontologías”

Basic Formal Ontology (BFO)

OBO Relation Ontology (RO)

OBO Foundry tiene mucho contenido de relativamente alta calidad, pero ...

... la mayoría de las ontologías OBO son muy pobres axiomáticamente

... OBO format

# Open Biological and Biomedical Ontologies

## OBO Format

No tiene una definición semántica clara: ¿Qué quieren decir las expresiones en OBO?

Mikel Egaña Aranguren, Sean Bechoffer, Phillip Lord, Ulrike Sattler and Robert Stevens. Understanding and using the meaning of statements in a bio-ontology: recasting the Gene Ontology in OWL. BMC Bioinformatics 2007, 8:57

Para usar inferencias, traducir a OWL ...

Christine Golbreich, Matthew Horridge, Ian Horrocks, Boris Motik, and Rob Shearer. OBO and OWL: Leveraging semantic web technologies for the life sciences. ISWC 2007, 4825:169-182, 2007

... o usar razonadores “ad hoc”:

OBO Edit reasoner

OBD-SQL reasoner

OBO Language (OBOL)

# Open Biological and Biomedical Ontologies

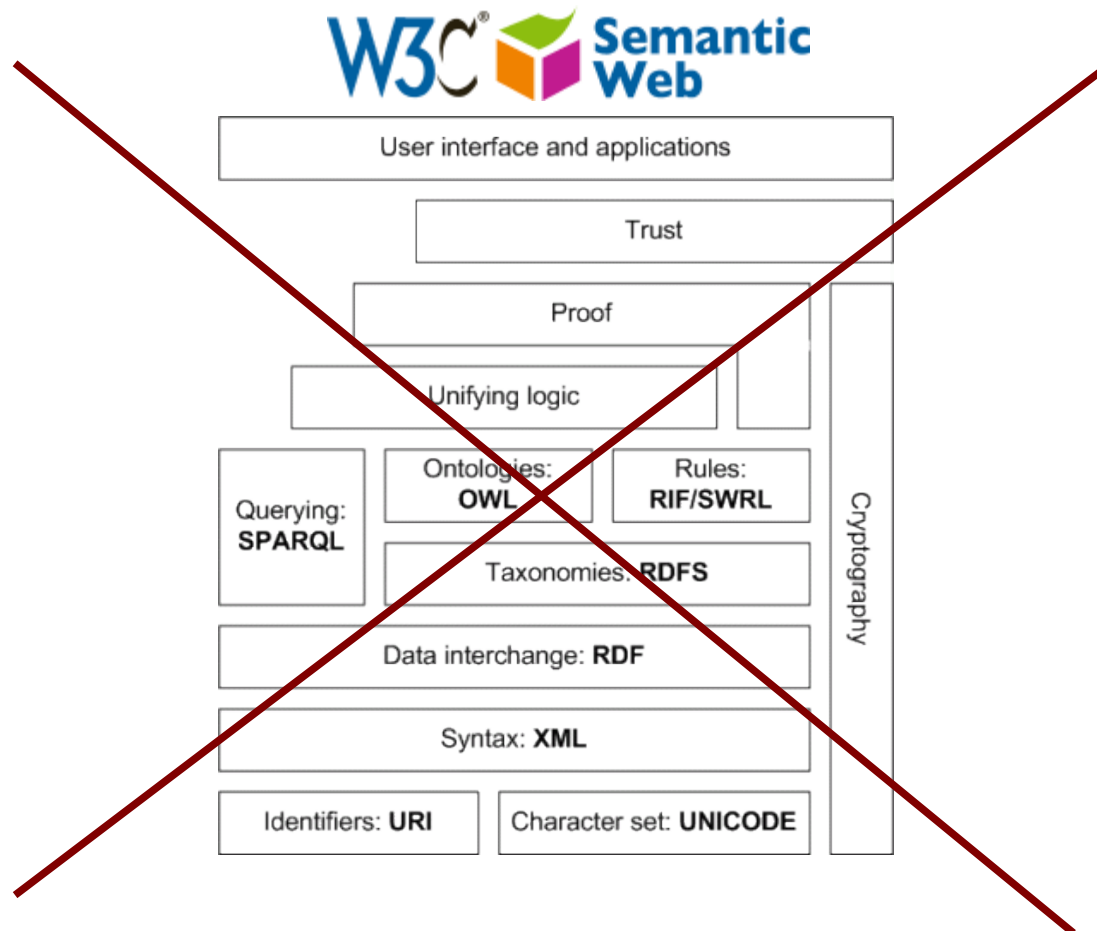
## OBO Format

```
[Term]
id: CCO:B0002060
name: NEB2_HUMAN
def: "Neurabin-2" [UniProt:Q96SB3]
synonym: "Neurabin-II" EXACT [UniProt:Q96SB3]
xref: UniProt:Q8TCR9
is_a: CCO:B0000000 ! core cell cycle protein
relationship: belongs_to CCO:T0000004 ! Homo sapiens organism
relationship: encoded_by CCO:G0005171 ! PPP1R9B_human
relationship: participates_in CCO:I0006401 ! aah62584-q96sb3 physical interaction
relationship: transforms_into CCO:B0013139 ! NEB2_HUMAN-Phosphoserine15
```



# Open Biological and Biomedical Ontologies

OBO format



**¿Por qué no se  
aplica LSSW?**

## ¿Por qué no se aplica LSSW?

Problema general en bioinformática: los biólogos crean sus propias soluciones artesanalmente y con afán de publicar

Carole Goble, The Seven Deadly Sins of  
Bioinformatics:

<http://tinyurl.com/6nvoe4>

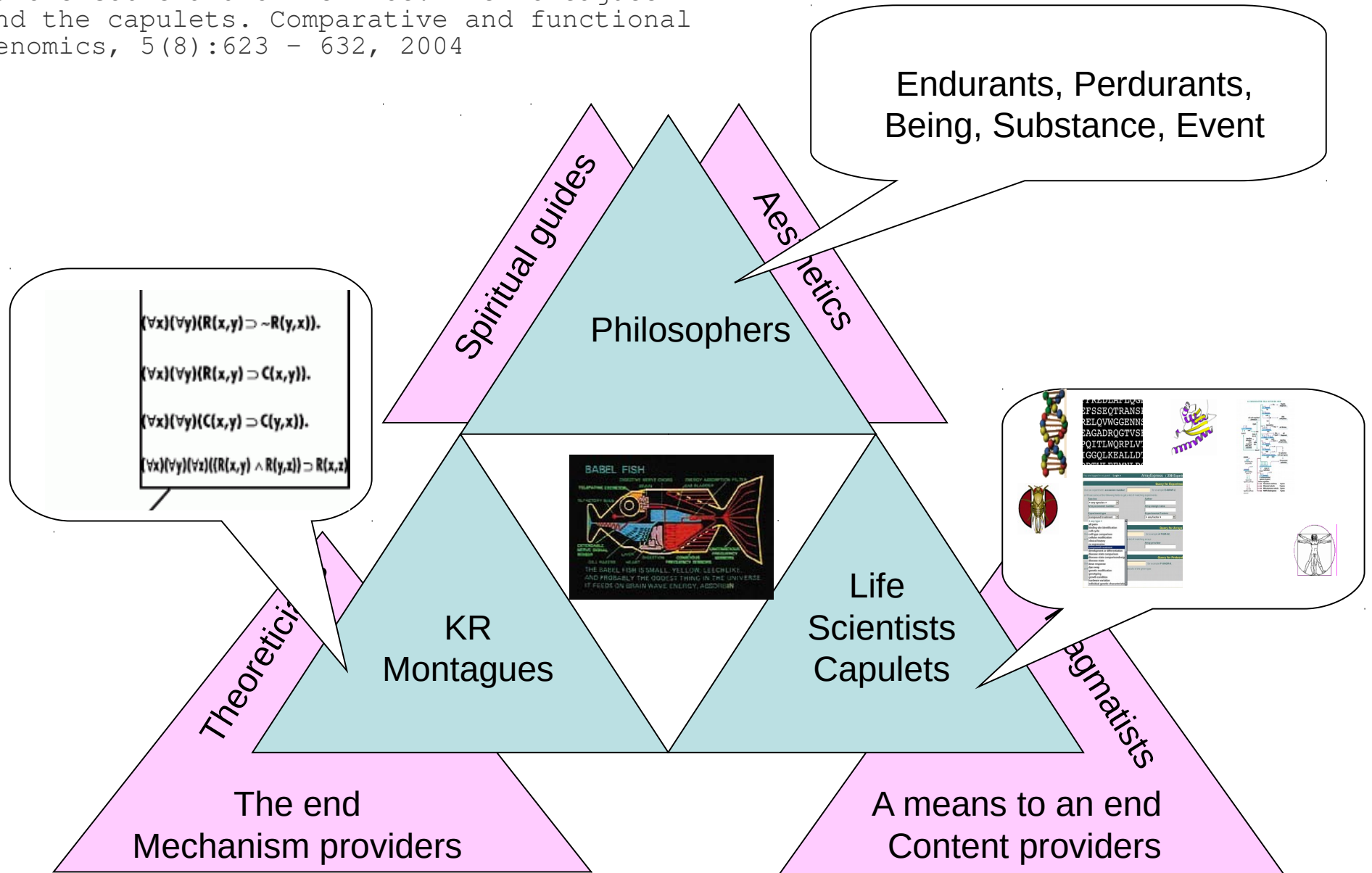
OBO es un ejemplo más de “artesanía”

Cuando surgió GO, OWL no existía (y mucho menos  
Protégé)

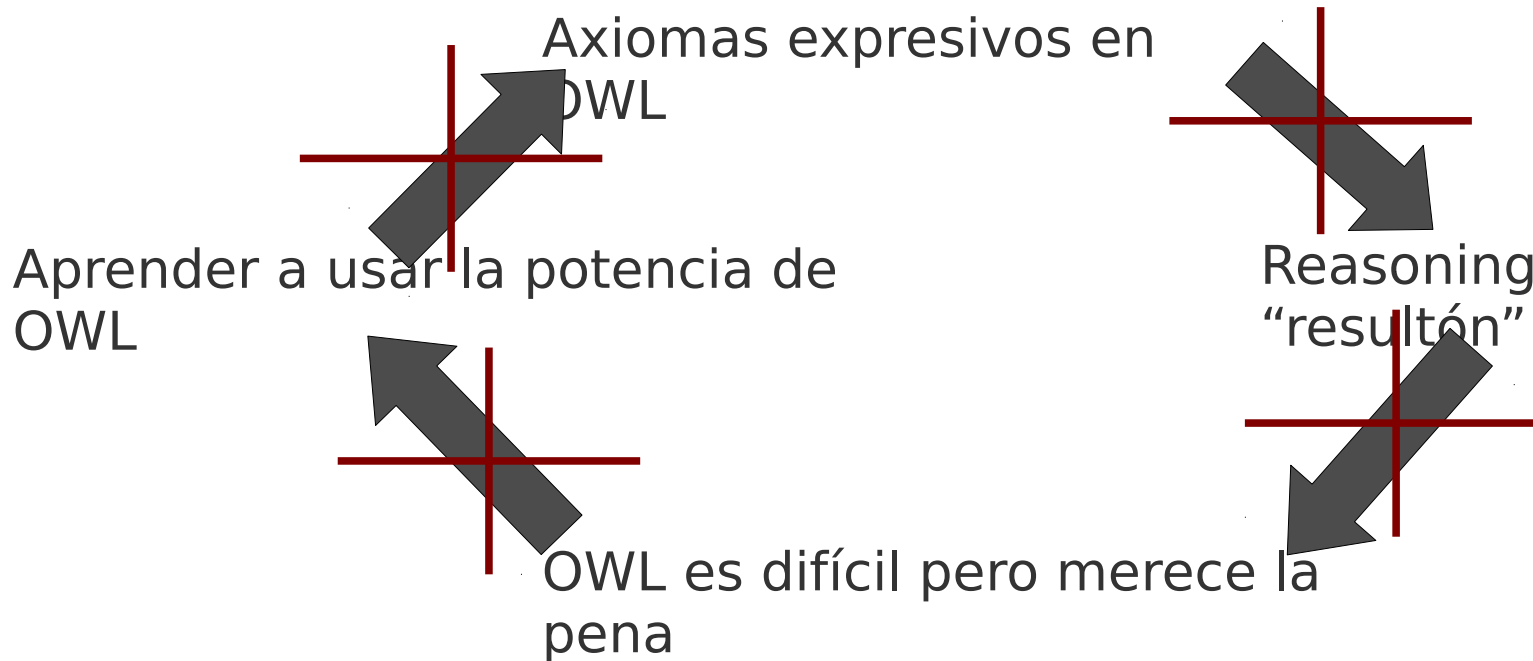
OWL es anti-intuitivo (ej. OWA, !UNA, ... )

# ¿Por qué no se aplica LSSW?

Carole Goble and Chris Wroe. The montagues and the capulets. Comparative and functional genomics, 5(8):623 - 632, 2004



## ¿Por qué no se aplica LSSW?



Las ontologías se diseñan para consumo humano (ej. Muchos axiomas "enterrados" en anotaciones en GO)

La prioridad es la integración de recursos, no la representación de conocimiento

## ¿Por qué no se aplica LSSW?

Ingeniería ontológica vs. ingeniería software

DEXA 2010 (Deusto). Philosophy Goes Information Technology - A Critical Reflection on Ontologies. Nick Falkner, University of Adelaide, Australia.

# Conclusiones

## Conclusiones

Para que LSSW se implante

IDs (Shared Names?)

Bio-ontologías axiomáticamente ricas (más funcionalidades): best practices (Ontology Design Patterns, ...)

Mikel Egaña, Alan Rector, Robert Stevens, Erick Antezana.  
Applying Ontology Design Patterns in bio-ontologies. EKAW  
2008, LNCS 5268, pp. 7-16

Bio-ontologías *consensuadas*:

Upper Level Ontology

RO

Menos “Realism”: bio-ontologías como medio, no como fin.

Phillip Lord and Robert Stevens. Adding a little reality to building ontologies for biology. PLoS ONE, 5(9):e12258, 2010.



## Conclusiones

Para que LSSW se implante:

Herramientas

¡OBO se adoptó por OBO-Edit!

Reasoning eficiente, predecible, y comprensible

Demostración de beneficios inmediatos de reasoning

Consultas

Generación de hipótesis

Mantenimiento

## Reasoning comprehensible

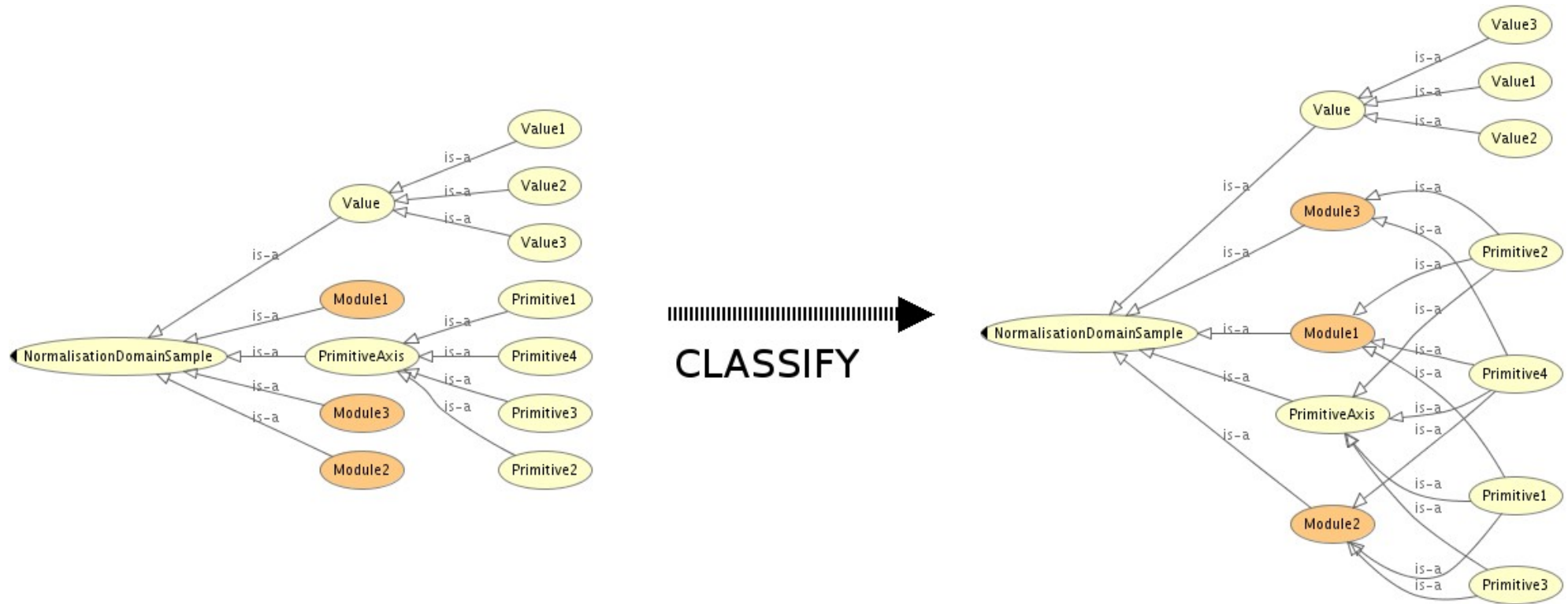
<http://owl.cs.manchester.ac.uk/explanation/>

	Employee <b>subClassOf</b> Person
1)	Employee <b>subClassOf</b> worksFor <b>some</b> Thing
2)	worksFor <b>subPropertyOf</b> memberOf
3)	memberOf <b>subPropertyOf</b> <b>inv</b> ( member )
4)	member <b>range</b> Person

# Conclusiones

## Reasoning para mantenimiento

<http://www.gong.manchester.ac.uk/odp/html/Normalisation.html>

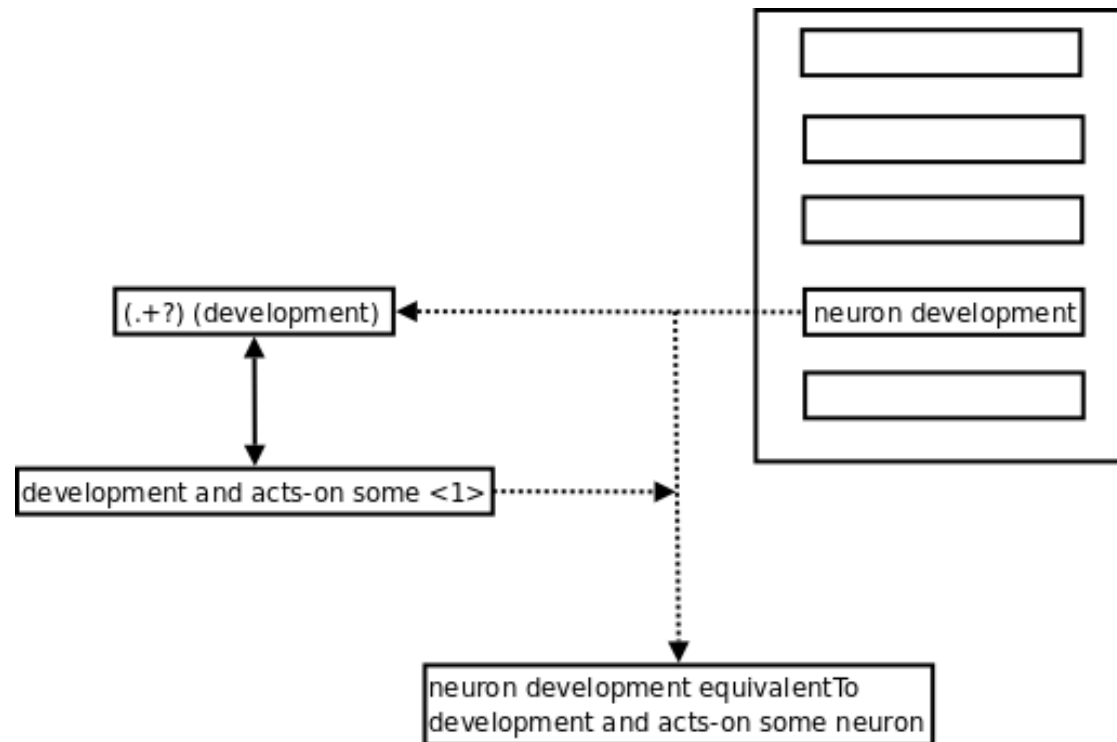


# Open Biological and Biomedical Ontologies

Reasoning para mantenimiento y consultas

Gene Ontology Next Generation (GONG): más axiomas con mínimo esfuerzo

Mikel Egaña Aranguren, Chris Wroe, Carole Goble, Robert Stevens. In situ migration of handcrafted ontologies to Reason-able Forms. Data & Knowledge Engineering 2008, 66, 147-162



## Conclusiones

Life Sciences Semantic Web:  
¿“Killer app” de Semantic Web?

**Para saber más ...**

**Para saber más ...**

Bio-ontologies SIG at ISMB (Intelligent Systems for  
Molecular Biology) <http://www.bio-ontologies.org.uk/>

SWAT4LS (Semantic Web Applications and Tools for Life  
Sciences) <http://www.swat4ls.org/>

ICBO (International Conference on Biomedical Ontology)  
<http://icbo.buffalo.edu/>

**Para saber más ...**

**EBI (European Bioinformatics Institute)** <http://www.ebi.ac.uk/>

**NCBO (National Center for Biomedical Ontology)**

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

**Journal of Biomedical Semantics** <http://www.jbiomedsem.com/>

**Semantic Systems Biology** <http://www.semantic-systems-biology.org/>

**Repositorios:**

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

<http://obo.sf.net/>

<http://www.ebi.ac.uk/ontology-lookup/>



**Mi trabajo en todo  
esto (Hasta ahora)**

## **Mi trabajo en todo esto (hasta ahora)**

Hacer SW más fácil a los biólogos

Ontology Design Patterns

OPPL 2

GONG

Tutoriales OWL

Construir recursos que usan SW

Cell Cycle Ontology

BioGateway

OGO

<http://mikeleganaaranguren.wordpress.com/publications/>

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