



Introducing federated queries with Wikidata

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The Gene Wiki project, circa 2008

Summarized knowledge via crowdsourcing

ITK (gene)

Function

This gene encodes an intracellular tyrosine kinase expressed in T-cells. The protein is thought to play a role in T-cell proliferation and differentiation.^{[2][3]}

Structure

The protein contains the following domains, which are often found in intracellular kinases:^[4]

- N-terminus – PH (pleckstrin homology domain)
- BTK – Bruton's tyrosine kinase Cys-rich motif
- SH3 – (Src homology 3)
- SH2 – (Src homology 2)
- C-terminus – tyrosine kinase, catalytic domain

Interactions

ITK (gene) has been shown to interact with FYN,^{[5][6]} Wiskott-Aldrich syndrome protein,^{[7][8]} KDR/KBSB,^{[9][10]} PLCG1,^{[10][11]} Lymphocyte cytosolic protein 2,^{[11][12]} Linker of activated T cells,^{[12][13]} Karyopherin alpha 2,^[14] Grb2^[15] and Peptidylprolyl isomerase A.^[15]

References

- ^ Gibson S, Leung B, Squire JA, Hill M, Arima N, Goss P, Hogd D, Mills GB (September 1993). "Identification, cloning, and characterization of a novel human T-cell-specific tyrosine kinase located at the hematopoietin complex on chromosome 5q". *Blood* 82 (5): 1561–72. PMID 8354206.
- ^ Kosaka Y, Felices M, Berg LJ (October 2006). "Itk and Th2 responses: action but no reaction". *Trends Immunol* 27 (10): 453–60. doi:10.1016/j.it.2006.08.006. PMID 16931159.
- ^ "Entrez Gene: ITK: IL2-inducible T-cell kinase".
- ^ Hawkins J, Marcy A (July 2001). "Characterization of the Itk tyrosine kinase: comparison of its catalytic domains to enzymatic activity". *Protein Expr Purif* 23 (2): 211–9. doi:10.1006/pepro.2001.1447. PMID 11437598.
- ^ a b c Bunnell, S. C., Dlehn, M., Yaffe, M. B., Findell, P., Cantley, L. C., Berg, L. J. (Jan. 2000). "Biochemical interactions integrating Itk with the T cell receptor-initiated signaling cascade". *J. Biol. Chem.* (UNITED STATES) 275 (3): 2219–30. ISSN 0021-9258. PMID 10536929.
- ^ a b c d e f g h i j k l m n o p q r s t u v w x y z Bunnell, S. C., Dlehn, M., Yaffe, M. B., Findell, P., Cantley, L. C., Berg, L. J. (Jan. 2000). "Biochemical interactions integrating Itk with the T cell receptor-initiated signaling cascade". *J. Biol. Chem.* (UNITED STATES) 275 (3): 2219–30. ISSN 0021-9258. PMID 10536929.
- ^ Perez-Villar, J., Kanner, S. B. (Dec. 1999). "Regulated association between the tyrosine kinase Emt1/Tsk and phospholipase-C gamma 1 in human T lymphocytes". *J. Immunol.* (UNITED STATES) 163 (12): 6435–41. ISSN 0021-1767. PMID 10580303.
- ^ Shim, Eun Kyung, Moon Chang Suk, Lee Gi Yeon, Ha Yun Jung, Chae Suhn-Kee, Lee Jong Ran (Sep 2004). "Association of the Src homology 2 domain containing leukocyte phosphatase with the p70 S6 kinase 1 (p70S6K1) with the p85 subunit of phosphatidylinositol 3-kinase". *FEBS Letters* (Netherlands) 575 (1-3): 35–40. doi:10.1016/j.febslet.2004.07.090. PMID 15388330. ISSN 0014-5793.
- ^ Shan, X., Wang, R. L. (Oct 1999). "Itk/Emt1/Tsk activation in response to CD3 cross-stimulation in Jurkat T cells requires ZAP-70 and Lat and is independent of membrane proximal". *J. Biol. Chem.* (UNITED STATES) 274 (41): 29323–30. ISSN 0021-9258. PMID 10506192.
- ^ Perez-Villar, J., Juan, J., White, J., Lopez-Soler, J., Diaz, J., Kanner, S. B. (Oct 2000). "The Itk kinase is required for the T cell receptor-induced phosphorylation of the p85 subunit of PI(3,4,5)P3-kinase". *Mol Cell Biol* 20 (20): 8231–40. ISSN 0270-7306. PMID 11006030.
- ^ a b c d e f g h i j k l m n o p q r s t u v w x y z Perez-Villar, J., Juan, J., White, J., Lopez-Soler, J., Diaz, J., Kanner, S. B. (Oct 2000). "The Itk kinase is required for the T cell receptor-induced phosphorylation of the p85 subunit of PI(3,4,5)P3-kinase". *Mol Cell Biol* 20 (20): 8231–40. ISSN 0270-7306. PMID 11006030.

Data imported
from structured
databases

Reelin

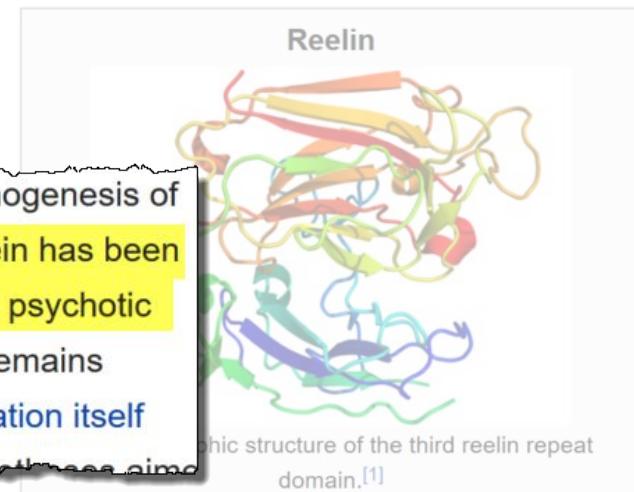
From Wikipedia, the free encyclopedia

Reelin is a large secreted extracellular matrix glycoprotein that helps regulate processes of neuronal migration and positioning in the developing brain by controlling cell–cell interactions. Besides this important role in early development, reelin continues to work in the adult brain. It modulates synaptic plasticity by [2][3] It also stimulates dendrite[4] migration of neuroblasts general zones. It is found not only in the tissues.

Reelin has been suggested to be expression of the protein has been bipolar disorder, but the cause of this observation remains uncertain as studies show that psychotropic medication itself affects reelin expression. Moreover, epigenetic hypotheses aimed at explaining the changed levels of reelin expression[6] are controversial.[7][8] Total lack of reelin causes a form of lissencephaly. Reelin may also play a role in Alzheimer's disease, temporal lobe epilepsy and autism.

Reelin's name comes from the abnormal reeling gait of *reeler* mice,[9] which were later found to have a deficiency of this brain protein and were homozygous for mutation of the RELN gene. The

Reelin has been suggested to be implicated in pathogenesis of several brain diseases. The expression of the protein has been found to be significantly lower in schizophrenia and psychotic bipolar disorder, but the cause of this observation remains uncertain as studies show that psychotropic medication itself



3D ribbon diagram of the third reelin repeat domain.[1]

Available structures

PDB Ortholog search: PDBe , RCSB

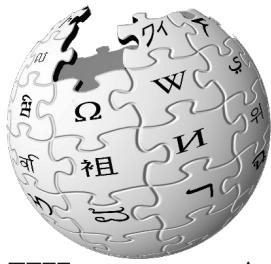
List of PDB id codes

[show]

Identifiers

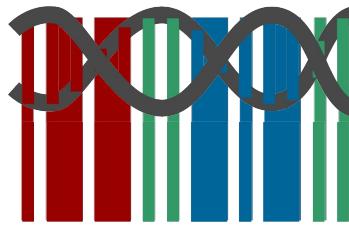
Symbols RELN ; LIS2; PRO1598; RL

External OMIM: 600514 MGI: 103022



WIKIPEDIA
The Free Encyclopedia

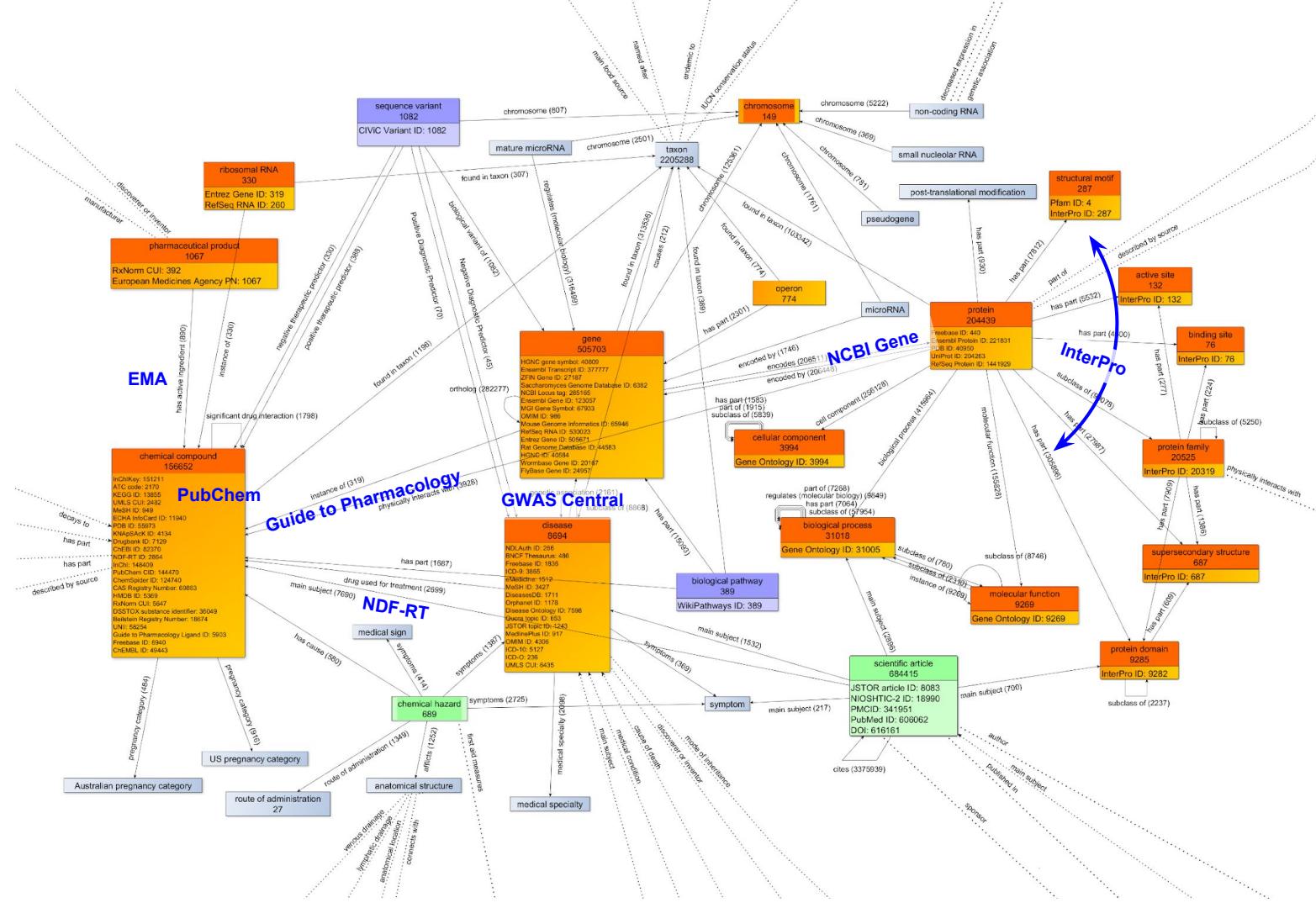
is to text



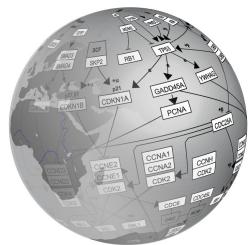
is to data

biomed
cal

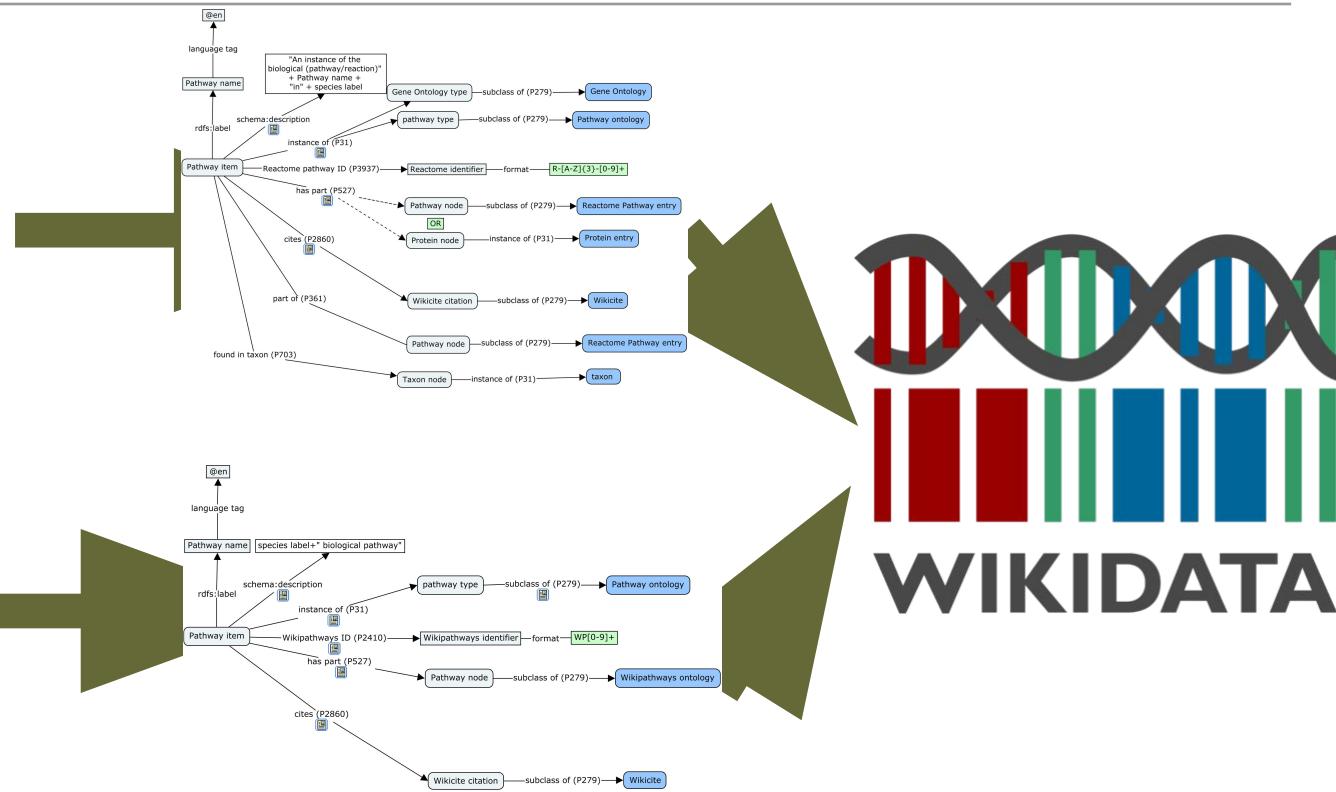
“Provide a database of the world’s knowledge that anyone can edit
- Denny Vrandečić



Pathways in Wikidata



WIKIPATHWAYS
Pathways for the People



Reactome Properties in Wikidata Properties [edit]

| Wikidata properties related to Reactome (Q2134522) | |
|---|--|
| Required properties | instance of (P31) • has part (P527) • |
| | part of (P361) • Reactome pathway ID (P3937) • |
| | found in taxon (P703) • exact match (P2888) • |
| Optional properties | cites (P2860) • |
| Reference properties | stated in (P248) • retrieved (P813) • Reactome pathway ID (P3937) • |
| Primary sources | Reactome (Q2134522) • |
| External Ontologies | Pathway Ontology (Q28864280) • Gene Ontology (Q135085) • |
| Species covered | Homo sapiens (Q15978631) • |

Wikipathways Properties in Wikidata Properties [edit]

| Wikidata properties related to biological pathway (Q4915012) | |
|---|---|
| Required properties | instance of (P31) • has part (P527) • |
| | WikiPathways ID (P2410) • found in taxon (P703) • |
| | exact match (P2888) • |
| Optional properties | connects with (P2789) • cites (P2860) • |
| Reference properties | stated in (P248) • retrieved (P813) • WikiPathways ID (P2410) • |
| Primary sources | WikiPathways (Q7999828) • |
| External Ontologies | Pathway Ontology (Q28864280) • Cell line ontology (Q21039006) • Disease Ontology (Q5282129) • |
| Species covered | Homo sapiens (Q15978631) • |

Simple data retrieval

“Retrieve genes with GWAS association with asthma”



39 genes

| gene | geneLabel | gene | geneLabel | gene | geneLabel | gene | geneLabel |
|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| Q5013317 | COL22A1 | Q18027370 | IGSF3 | Q18053559 | CDHR3 | Q14903974 | SMAD3 |
| Q14912759 | SLC22A5 | Q18045382 | HPSE2 | Q18045669 | ATG3 | Q18033889 | IL1RL1 |
| Q14914243 | PSAP | Q18048437 | IL33 | Q18035037 | RAD50 | Q17917202 | ERBB4 |
| Q14907990 | SLC30A8 | Q18051900 | PYHIN1 | Q18036984 | FBXL7 | Q18027836 | IL6R |
| Q18025002 | GAB1 | Q17709208 | ACO1 | Q18033919 | XPR1 | Q18030185 | NOTCH4 |
| Q18035589 | C6orf10 | Q18027822 | IL2RB | Q15326496 | RORA | Q18030409 | PDE4D |
| Q18054256 | GSDMA | Q18030364 | PBX2 | Q18042132 | GSDMB | Q18045645 | IKZF4 |
| Q18058487 | C5orf56 | Q18037773 | ABI3BP | Q18029145 | MKLN1 | Q18039979 | KLHL5 |
| Q18030785 | PRKG1 | Q18039623 | CTNNA3 | Q18036729 | RAP1GAP2 | Q18026947 | HLA-DQA1 |
| Q18033424 | IL18R1 | Q18046350 | ZNF665 | Q14878303 | IL13 | | |

```

1 SELECT DISTINCT ?gene ?geneLabel where {
2   ?gene wdt:P2293 wd:Q35869 . # gene has genetic association to "asthma"
3   ?gene wdt:P31 wd:Q7187 .      # gene is subclass of "gene"
4   SERVICE wikibase:label { bd:serviceParam wikibase:language "en". }
5 }
```

Data integration

“Retrieve genes with GWAS association with asthma and gene product is localized to membrane”



22 genes

| gene | geneLabel | gene | geneLabel | gene | geneLabel | gene | geneLabel |
|---------------|-----------|---------------|-----------|---------------|-----------|---------------|-----------|
| Q1491275 9 | SLC22A5 | Q1802737 0 | IGSF3 | Q1803503 7 | RAD50 | Q1802783 6 | IL6R |
| Q1491424 3 | PSAP | Q1803342 4 | IL18R1 | Q1803391 9 | XPR1 | Q1803040 9 | PDE4D |
| Q1490799 0 | SLC30A8 | Q1804538 2 | HPSE2 | Q1804213 2 | GSDMB | Q1803018 5 | NOTCH4 |
| Q1803558 9 | C6orf10 | Q1802782 2 | IL2RB | Q1803672 9 | RAP1GAP2 | Q1802694 7 | HLA-DQA1 |

```

1 SELECT DISTINCT ?gene ?geneLabel where {
2   ?gene wdt:P2293 wd:Q35869 . # gene has genetic association to "asthma"
3
4   ?gene wdt:P31 wd:Q7187 .      # gene is subclass of "gene"
5
6   ?gene wdt:P688 ?protein .      # gene encodes a protein
7   ?protein wdt:P681 ?cc .        # protein has a cellular component
8   ?cc wdt:P279*|wdt:P361* wd:Q14349455 . # cell component is 'part of' or 'subclass of' membrane
9
10 SERVICE wikibase:label { bd:serviceParam wikibase:language "en". }
11 }
```

Computing on provenance

“Retrieve genes with GWAS association with asthma and gene product is localized to membrane (non-IEA)”



15 genes

| gene | geneLabel | gene | geneLabel | gene | geneLabel |
|-----------|-----------|-----------|-----------|-----------|-----------|
| Q14912759 | SLC22A5 | Q18045382 | HPSE2 | Q17917202 | ERBB4 |
| Q14914243 | PSAP | Q18027822 | IL2RB | Q18027836 | IL6R |
| Q14907990 | SLC30A8 | Q14903974 | SMAD3 | Q18030409 | PDE4D |
| Q18027370 | IGSF3 | Q18035037 | RAD50 | Q18030185 | NOTCH4 |
| Q18033424 | IL18R1 | Q18036729 | RAP1GAP2 | Q18026947 | HLA-DQA1 |

```

6 ?gene wdt:P31 wd:Q7187 ;      # gene is subclass of "gene"
7     wdt:P688 ?protein ;      # gene encodes a protein
8     rdfs:label ?geneLabel .
9 FILTER (lang(?geneLabel) = "en")
10 ?protein p:P681 ?s .           # protein's cell component statement
11     ?s ps:P681 ?cp .          # get statement value
12     FILTER NOT EXISTS {?s pq:P459 wd:Q23190881 .} # determination method is not IEA
13     ?cp wdt:P279*|wdt:P361* wd:Q14349455 .          # statement value is 'part of' or 'subclass of' membrane
14

```

Leveraging the Disease Ontology structure

“Retrieve genes with GWAS association with any respiratory disease and gene product is localized to membrane (non-IEA)”



31 genes / 8 diseases

| diseaseGALabel | gene_counts | geneList |
|---------------------------------------|-------------|---|
| asthma | 15 | SMAD3, RAP1GAP2, IL18R1, HPSE2, SLC30A8, SLC22A5, PSAP, ERBB4, HLA-DQA1, IGSF3, IL2RB, IL6R, NOTCH4, PDE4D, RAD50 |
| chronic obstructive pulmonary disease | 5 | HLA-C, SFTPB, ANXA5, ANXA11, ATP2C2 |
| lung cancer | 3 | TGM5, VTI1A, PHACTR2 |
| interstitial lung disease | 2 | DSP, ATP11A |
| non-small-cell lung carcinoma | 2 | NALCN, DLST |
| nasopharynx carcinoma | 2 | ITGA9, TNFRSF19 |
| adenocarcinoma of the lung | 1 | BTNL2 |
| pulmonary emphysema | 1 | BICD1 |

```

1 SELECT ?diseaseGALabel (count (DISTINCT ?geneLabel) AS ?geneCounts)
2 (group_concat(DISTINCT ?geneLabel; separator = ",") AS ?geneList)
3 ?gene wdt:P2293 ?diseaseGA .                                # genes
4 ?diseaseGA wdt:P279* wd:Q3286546 .                         # to a
5
6 ?gene wdt:P31 wd:Q7187 ; wdt:P688 ?protein ;               # gene is subclass of "gene" and encodes protein
7     rdfs:label ?geneLabel .
8 FILTER (lang(?geneLabel) = "en")
9 ?protein p:P681 ?s .                                         # protein's cell component statement
10    ?s ps:P681 ?cp .                                         # get statement value
11 FILTER NOT EXISTS (?s p:P681 ?o)

```

Opportunistic integration

“Retrieve genes with GWAS association with any respiratory disease and gene product is localized to membrane (non-IEA) **and show causative chemical hazards**”



4 diseases / 6 chemical hazards

| diseaseGALabel | exposureLabel |
|---------------------|----------------------------|
| lung cancer | arsenic pentoxide exposure |
| lung cancer | HN1 exposure |
| lung cancer | mechlorethamine exposure |
| lung cancer | HN3 exposure |
| asthma | Phenacyl chloride exposure |
| pulmonary emphysema | phosgene exposure |

```

11 .cp wdt:P279 wd:Q2116751 . # statement values are part of our
12
13 ?exposure wdt:P1542 ?diseaseGA . # something causes disease
14 ?exposure wdt:P279 wd:Q2116751 . # and that something is a chemical hazard
15
16 SERVICE wikibase:label { bd:serviceParam wikibase:language "en". }
17 }
```

... and show associated pathways

“Retrieve genes with GWAS association with any respiratory disease and gene product is localized to membrane (non-IEA), show causative chemical hazards and **show pathways where they have a role.**”



16 genes / 59 pathways

| gene | pathway |
|-------|--|
| SMAD3 | Androgen receptor signaling pathway |
| SMAD3 | TGF-beta Receptor Signaling |
| SMAD3 | mechlorethamine exposure |
| HLA-C | Allograft Rejection |
| SFTPD | Regulation of toll-like receptor signaling pathway |
| | |

```

11 .cp wdt:P279 wd:Q501 . # statement values are part of rows
12
13 ?pathway wdt:P31 wd:Q4915012 ; # instance of a biological pathway
14   wdt:P527 ?gene .
15
16 SERVICE wikibase:label { bd:serviceParam wikibase:language "en". }
17 }
```

... and show remote pathway annotations

16 genes / 59 pathways

“Retrieve genes with GWAS association with any respiratory disease and gene product is localized to membrane (non-IEA), show causative chemical hazards, show pathways where they have a role and show pathway annotations in WikiPathways”

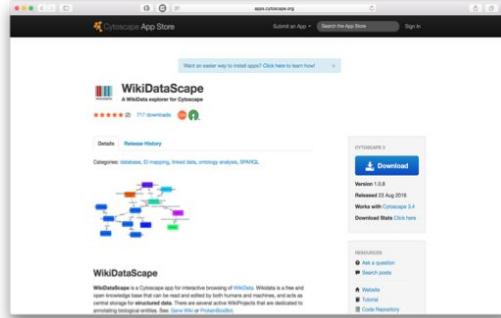


| Remote pathway annotations | |
|----------------------------|-------|
| Pathway ontology | 55 |
| Cell line ontology | 12 |
| Disease ontology | 11 |
| | |

```
11 .cp wdt:r2t wd:TP501 wd:Q1449455 .      # statements values are part of rows
12
13     wdt:P2888 ?source_pathway .
14
15     SERVICE <http://sparql.wikipathways.org/> {
16         ?wp_pathway dc:identifier ?source_pathway ;
17             wp:ontologyTag ?pwTag .
}
```

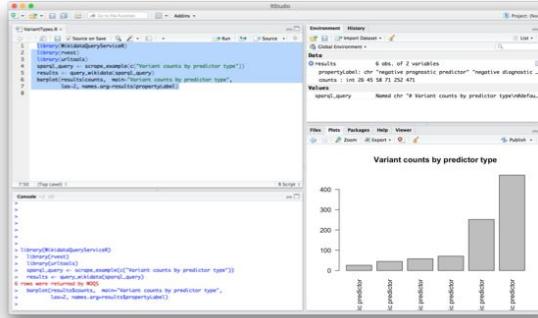
Tools using Wikidata

<http://www.wikigenomes.org>



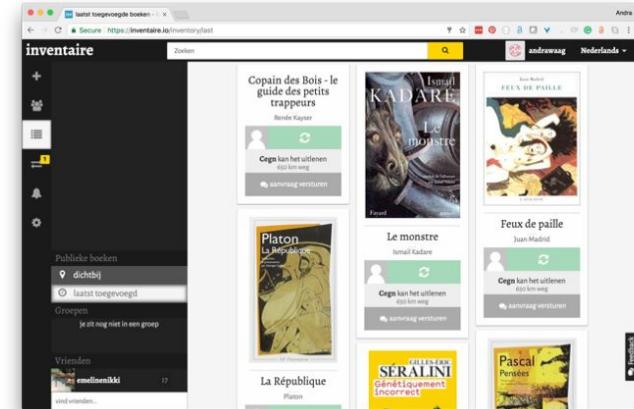
Cytoscape

Wikipedia and Wikidata
google plugin



R plugins

<http://inventaire.io>



Availability

www.wikidata.org/wiki/User:Pathwaybot

- www.wikidata.org/wiki/User:Pathwaybot/data_model - Semantic models
- www.wikidata.org/wiki/User:Pathwaybot/query_examples – Pathway examples
- www.wikidata.org/wiki/User:ProteinBoxBot/SPARQL_Examples– Genewiki example

github.com/SuLab/GeneWikiCentral

- github.com/SuLab/wikidataintegrator – python module for Wikidata
- github.com/SuLab/scheduled-bots – bot automation framework
- github.com/SuLab/Genewiki-ShEx – data models



Structure of the federated query

```
SELECT * WHERE {  
    ?localitem ?localproperty ?shareditem .  
    SERVICE <http://example.remote.sparql> {  
        ?shareditem ?remoteproperty ?remoteitem .  
    }  
}
```

Local pattern

Remote pattern

A federated query contains query patterns for both the local endpoint (green box) and a remote endpoint (blue box). The address of the remote SPARQL endpoint is expressed with the SERVICE keyword

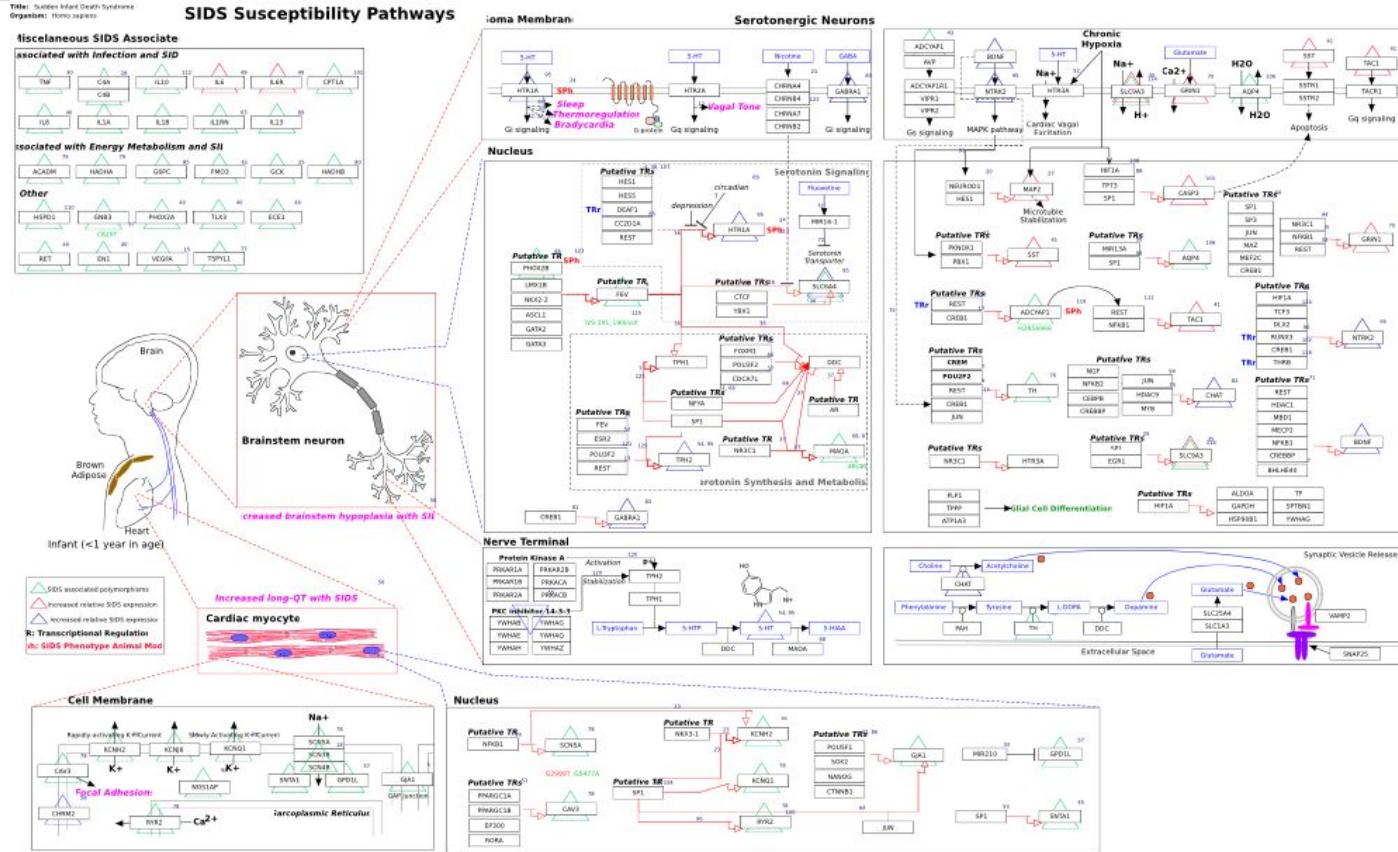
Why federate

- Incorporate more fine-grained data not captured in Wikidata
- Link to other (non-public licenses)
- Link to volatile data

Wikidata Query Service support federated queries

- From Wikidata to an external SPARQL endpoint (Wikipathways)
- From a remote SPARQL endpoint to Wikidata
- From a local SPARQL endpoint to Wikidata
- From a local Wikibase (with WDQS) to Wikidata

From Wikidata to an external SPARQL endpoint (WikiPathways)



PREFIX wp: <<http://vocabularies.wikipathways.org/wp#>>
PREFIX dcterms: <<http://purl.org/dc/terms/>>
PREFIX dc: <<http://purl.org/dc/elements/1.1/>>
SELECT DISTINCT ?metabolite1Label ?metabolite2Label ?mass1 ?mass2 WITH {

```
  SELECT ?metabolite1 ?metabolite2 WHERE {  
    ?pathwayItem wdt:P2410 "WP706";  
                 wdt:P2888 ?pwIri.
```

Wikidata

```
  SERVICE <http://sparql.wikipathways.org/> {  
    ?pathway dc:identifier ?pwIri.  
    ?interaction rdf:type wp:Interaction;  
                 wp:participants ?wpmb1, ?wpmb2;  
                 dcterms:isPartOf ?pathway.  
    FILTER (?wpmb1 != ?wpmb2)  
    ?wpmb1 wp:bdbWikidata ?metabolite1.  
    ?wpmb2 wp:bdbWikidata ?metabolite2.  
  }  
}
```

Wikipathways

```
} AS %metabolites WHERE {  
  INCLUDE %metabolites.  
  ?metabolite1 wdt:P2067 ?mass1.  
  ?metabolite2 wdt:P2067 ?mass2.  
  SERVICE wikibase:label { bd:serviceParam wikibase:language "[AUTO_LANGUAGE],en". }
```

Wikidata

From a remote SPARQL endpoint to Wikidata

The screenshot shows the UniProt SPARQL endpoint interface. At the top, there are links for "SPARQL", "Downloads", and "Documentation/Help". Below this, a section titled "Your query" contains a SPARQL query. The query is divided into two main parts by a green box:

```
20 SELECT DISTINCT ?wd_item ?physically_interacts_with ?interactswithLabel ?type ?iri ?uniprot ?text WHERE {
21   {SELECT * WHERE { ?iri a up:Protein ;
22     up:organism taxon:9606 ;
23     up:annotation ?annotation .
24     ?annotation a up:Natural_Variant_Annotation ;
25     rdfs:comment ?text .
26     FILTER (CONTAINS(?text, 'loss of function'))
27   }
28   SERVICE <https://query.wikidata.org/bigdata/namespace/wdq/sparql> {
29     VALUES ?use {wd:Q427492}
30     ?wd_item wdt:P352 ?uniprot ;
31       wdt:P129 ?physically_interacts_with ;
32       wdt:P2888 ?iri ;
33       wdt:P703 wd:Q15978631 .
34     ?wd_item p:P129 ?phys_interacts_with_node .
35     ?phys_interacts_with_node ps:P129 ?physically_interacts_with ;
36       pq:P366 ?use .
37     ?physically_interacts_with wdt:P31 ?type ;
38       rdfs:label ?interactswithLabel .
39     FILTER (lang(?interactswithLabel) = "en")
40   }
```

The first part of the query (lines 20-27) is annotated with the word "UniProt" in green. The second part (lines 28-40) is annotated with "Wikidata" in blue.

At the bottom left are buttons for "Submit Query" and "Cancel".

We use ShEx to describe contents of a remote endpoint (<http://shex.io>)

- Draft Shape Expression for the Europeana Sparql endpoint:

```
PREFIX rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>
PREFIX edm: <http://www.europeana.eu/schemas/edm/>
PREFIX rdvocab: <http://rdvocab.info/ElementsGr2/>
PREFIX dce: <http://purl.org/dc/elements/1.1/>
PREFIX dct: <http://purl.org/dc/terms/>
PREFIX skos: <http://www.w3.org/2004/02/skos/core#>
PREFIX dbp: <http://dbpedia.org/resource/>
PREFIX oaterms: <http://www.openarchives.org/ore/terms/>
PREFIX bibgalicia: <http://www.galiciiana.bibliotecadegalicia.xunta.es/aut/>
PREFIX gtaa: <http://data.beeldengeluid.nl/gtaa/>
PREFIX owl: <http://www.w3.org/2002/07/owl#>

<aggregation> {
    edm:dataProvider LITERAL* ;
    edm:rights <licenses>? ;
    edm:isShownBy IRI* ;
}

<concept> {
    rdf:type skos:Concept ;
    skos:prefLabel LITERAL* ;
    skos:broader [dbp:~]* ;
    skos:exactMatch IRI* ;
    skos:closeMatch IRI* ;
    skos:related [dbp:~ bibgalicia:~]* ;
    skos:notation LITERAL? ;
```