

https://disease-ontology.org/do-kb/

The DO-KB is expanding the disease information ecosystem

- expanding the representation of the DO diseaseome
- a comprehensive network of disease to disease relationships
- a disease feature similarity network for differential diagnosis exploration

DO-KB SPARQL Sandbox

- A data playground for querying the Human Disease Ontology SPARQL endpoint and to perform federated SPARQL queries across connected resources for disease knowledge discovery.
- A linked data representation (RDF triple store) of the Human Disease Ontology's diseases, disease features and mechanisms.

DO-KB Faceted Search Interface

- A new way to explore human disease features and mechanisms, represented here as data facets.
- The DO-KB Faceted Search Interface enables exploration of connectivity across diseases.

The DO-KB's SPARQL endpoint, explore DO data and query disease, genomic, proteomic resources through federated queries.

Mine the DO-KB disease-data connections through faceted query and retrieval of DO diseases sharing features or mechanisms, including phenotypes, environmental or genetic drivers, anatomy, variant type, and age of onset.



DO-KB SPARQL Sandbox

Select one of the provided SPARQL queries or paste in a novel query to retrieve bespoke ML-ready datasets. Searches may be performed against the DO's primary release file, doid.owl or the doid-merged.owl file for exploring annotations, e.g. ECO codes, and axiom relationships.

- The DO's SPARQL endpoint (https://sparql.disease-ontology.org) is available for building federated
 queries.
- Visit our SPARQL Resources page to learn more about using the SPARQL Sandbox and endpoint.
- Our SPARQL Sandbox and endpoint are currently in beta. If you discover any issues, please reach out
 and let us know!

Query

```
PREFIX rdfs: <a href="http://www.w3.org/2000/01/rdf-schema#">http://www.w3.org/2002/07/owl#</a>
PREFIX owt <a href="http://www.w3.org/2002/07/owl#">http://www.w3.org/2002/07/owl#</a>
PREFIX obc <a href="http://www.geneontology.org/formats/obolnOwl#">http://www.geneontology.org/formats/obolnOwl#</a>
SELECT ?class ?id ?label
WHERE {
    ?class a owl:Class ;
    obolnOwl.id ?id ;
    rdfs:label ?label ;
    rdfs:subClassOf* obo:DOID_225 .
}
```

Submit Query

Download Results 🕹

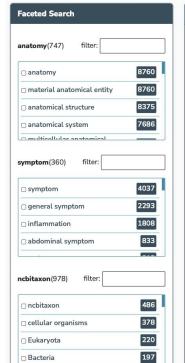
Choose a Query:

- All diseases with their parents (identified by label)
- 2. Count of diseases in each branch (diseases may be in multiple branches)
- Syndromes (branch search; returns IRI, ID and label)
- 4. Diseases with MeSH cross-references
- 5. Diseases with OMIM cross-references
- 6. Diseases with PubMed sources (limited to first 10 results)
- 7. Diseases in
 - DO_infectious_disease_slim with their parents and branches
- 8. Diseases with their definitions and count of logical axioms
- 9. Diseases with their Evidence and Conclusion Ontology (ECO) reference types (queries the doid-merged.owl graph)
- Count of classes imported from other ontologies (queries the doidmerged.owl graph)
- All diseases with equivalent class axioms (axioms are formatted for readability and may not be displayed in full; queries the doid-merged.owl graph)
- Extract the disease and supporting import class hierarchies using subClassOf relationships (executes a CONSTRUCT query that returns RDF triples)

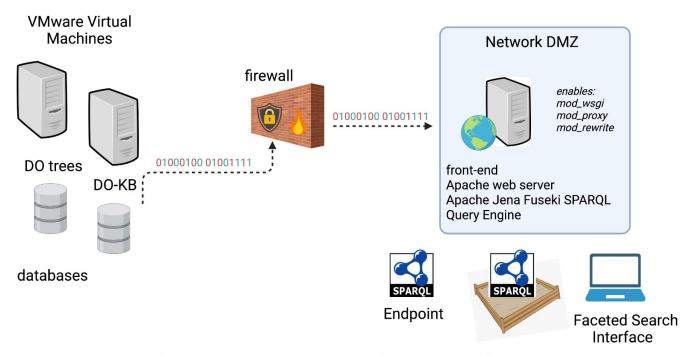
Request New SPARQL Queries

DO-KB Faceted Search Interface

Identify diseases with similar characteristics and explore relationships between diseases through common characteristics or features.



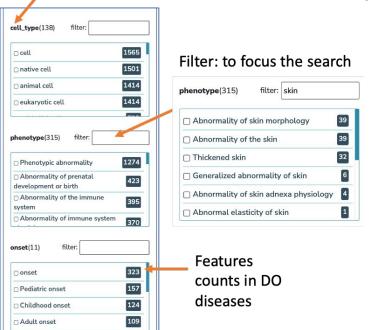


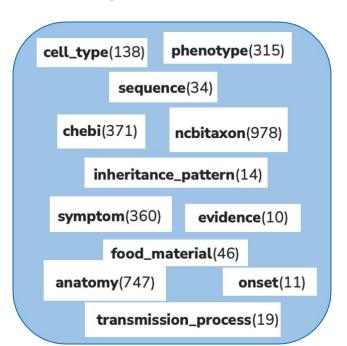


Augmenting DO technology to launch DO-KB. DO servers are VMware Virtual Machines running Ubuntu 22.04. DO server (Python-Django powered web application) and DO-KB serer driven by a Python-Flask framework. VMware vSphere client to monitor the health of our front-end server. Elasticsearch back-end service for the Faceted Search. DO-KB "OWL Flattener" program using Java and the Java OWL API package to convert the hierarchical DO OWL file into a flat list of documents

DO-KB Faceted Search Interface

- A Facet: a features or characteristic of a disease, defined in the DO via ontology imports
- Disease to Disease connections are defined by these features
- Facet boxes: (DO imports)
 - Select one or more features to retrieve DO diseases sharing those features or mechanisms
 - As facets are selected, the counts refresh, to show additional related features



















Identify diseases with similar characteristics and explore relationships between diseases through common characteristics or features

