

Tools

Options ▾

[Get Data](#)
[Send Data](#)
[ENCODE Tools](#)
[Lift-Over](#)
[Text Manipulation](#)
[Filter and Sort](#)
[Join, Subtract and Group](#)
[Convert Formats](#)
[Extract Features](#)
[Fetch Sequences](#)
[Fetch Alignments](#)
[Get Genomic](#)

Execute an OPPL file against an ontology

Input ontology file:

6: gene_ontology.1_2..s_fixed.owl ▾

Choose reasoner:

Pellet ▾

OPPL file:

5: Disjoint_localisation.oppl ▾

Choose ontology output format:

OWL ▾

Execute

History



Unnamed history

[21: Compare two Datasets](#)[20: Get the terms that are data 19](#)[19: Execute an OPPL file a](#)[18: Hepatocyte part of re](#)[14: Execute an OPPL file a](#)