Applying Ontology Design Patterns in bio-ontologies

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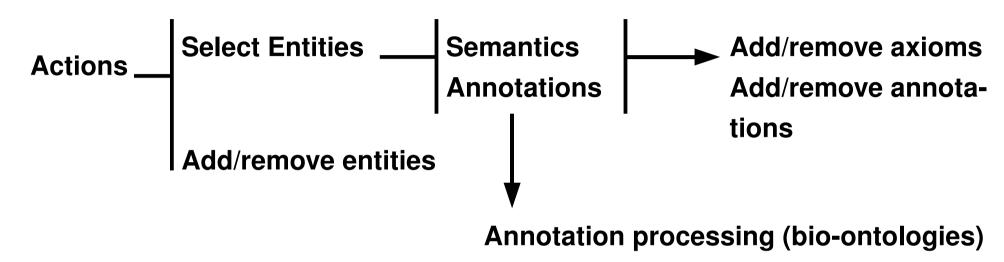
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ONTOLOGY PREPROCESSOR LANGUAGE (OPPL)

High level scripting language for OWL.



Asserted/inferred mode (Pellet, FaCT++, any DIG reasoner).

Currently two versions:

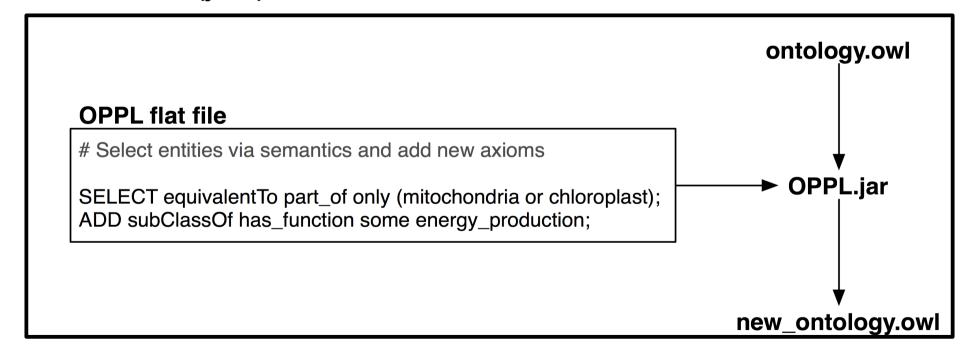
OPPL 1 (http://oppl.sf.net/)
OPPL 2 (http://www.cs.man.ac.uk/~iannonel/oppl/)

ONTOLOGY PREPROCESSOR LANGUAGE (OPPL)

OPPL syntax (Manchester OWL Syntax + OPPL keywords)

```
SELECT equivalentTo part_of only (mitochondria or chloroplast);
ADD subClassOf has_function some energy_production;
```

OPPL software (java)



ONTOLOGY PREPROCESSOR LANGUAGE (OPPL)

Store and share complex modelling for consistent application:

by different ontologists at different stages in different parts of the ontology (via queries)

Documented and explicit modelling: trace modelling.

Try complex modelling easily, then decide: prototypes.

Ontology cleansing/enrichment.

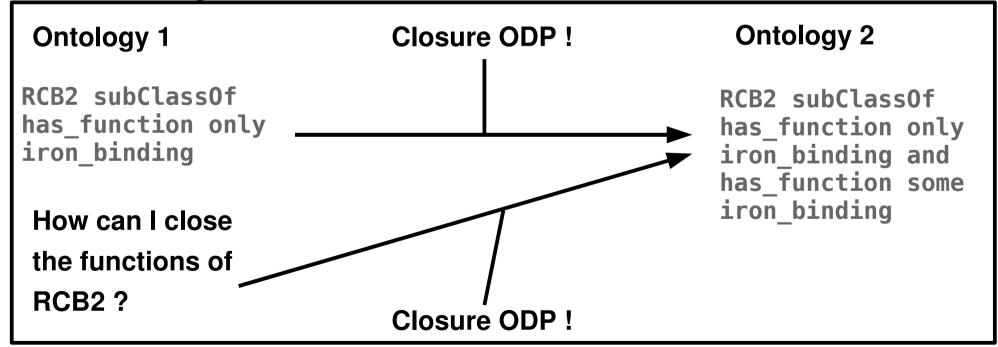
Ontology cleansing/enrichment in pipelines (e.g. CCO http://www.cellcycleontology.org/).

Automated modification/querying of big ontologies.

Ontology Design Patterns (ODPs): encapsulate complex semantics, easier modelling.

e.g. Closure ODP: prop only filler and prop some filler

Bio-ontologies: lean axiomisation



OPPL: store (flat files) and apply ODPs in OWL ontologies.

ODPs for modifiers

Entity-Quality ODP (E-Q ODP)
Entity-Property-Quality ODP (E-P-Q ODP)
Entity-Feature-Value ODP (E-F-V ODP)

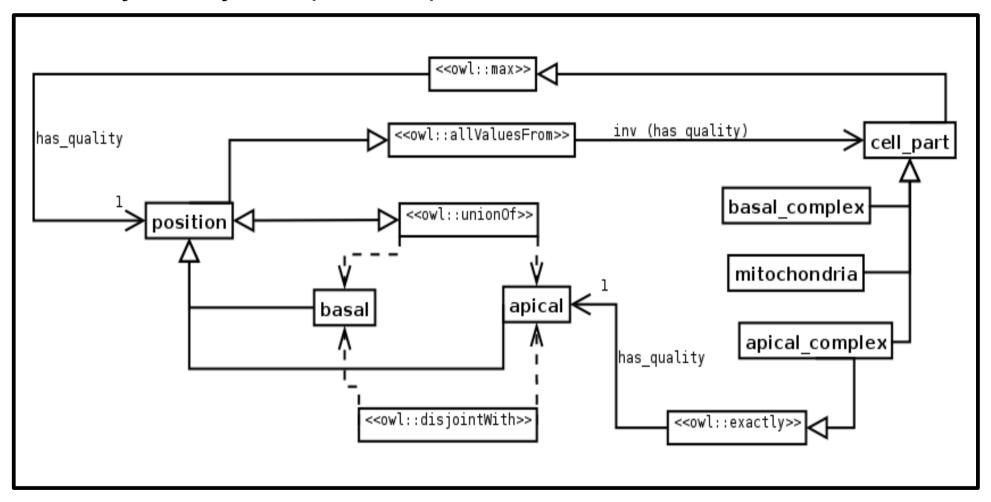
Try E-Q, E-P-Q, E-F-V, and then decide.

E-Q ODP in the Gene Ontology (GO): position of cell parts (e.g. the position of "apical complex" is the apical side of the cell).

Apply E-Q in GO via annotation query and processing with OPPL: 24/20,000.

Local vs global ODPs.

Entity-Quality ODP (E-Q ODP)



E-Q ODP applied in GO (OWL version) via OPPL script (flat file)

```
# Quality values
ADD Class: modifier:
ADD ObjectProperty: has_quality;
ADD Class: position; ADD subClassOf modifier; REMOVE subClassOf Thing;
ADD Class: apical; ADD subClassOf position; REMOVE subClassOf Thing;
ADD Class: basal; ADD subClassOf position; ADD disjointWith apical;
SELECT Class: position; ADD equivalent To apical or basal;
# Constrain the quality values to the entities (CCO_C0001882 = cell part)
SELECT Class: position; ADD subClassOf inv (has quality) only CCO C0001882;
# Not having a position is legal
SELECT Class: CCO_C0001882; ADD subClassOf has_quality max 1 position;
# Add position values to actual cell parts
SELECT label "(basallapical) (.+?)";ADD subClassOf has_quality exactly 1 <1>;
```

ONTOLOGY PREPROCESSOR LANGUAGE 2

Developed by Luigi lannone (BioHealth Informatics Group, University of Manchester).

Axiom centric, not entity centric: closer to OWL semantics.

Protégé plugin (autocomplete, ...).

Variables (e.g. Closure ODP)

?x:CLASS, ?z:CLASS SELECT ?x SubClassOf has_function only ?z
BEGIN ADD ?x SubClassOf has_function some ?z END;

Decidability: variables only to be bound by named entities, not expressions (Class, ObjectProperty, DataProperty, Individual, Constant).

CONCLUSION

OPPL: easy "programmatic" manipulation of OWL ontologies.

ODPs: semantic encapsulation; ease modelling.

OPPL for efficiently and consistently applying ODPs.

ODPs successfully applied in the CCO with OPPL:

Mikel Egaña Aranguren, Erick Antezana, Martin Kuiper, Robert Stevens.

Ontology Design Patterns for bio-ontologies: a case study on the

Cell Cycle Ontology. BMC bioinformatics 2008, 9(Suppl 5):S1.

http://www.biomedcentral.com/1471-2105/9/S5/S1



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