OWL Paths

A library for processing SPARQL-like property paths over OWL classes

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OWL Paths creation was motivated by our group’s need to validate consistency of modeling in a large ontology for human anatomy.

Anatomical structures repeated throughout the body (such as joints, muscles, and bones) should be modeled in similar and predictable ways.

We want to use path expressions to verify modeling consistency in our ontology.
Our work is about writing rules for authoring and auditing large biomedical ontologies

We call these rules “knowledge representation patterns”
A fictional medical record for an **individual** has these three triples:

```
demo:MedicalRecord_001
  demo:has_subsection
    demo:PersonalData_001
      demo:has_subsection
        demo:DemographicData_001
      demo:has_subsection
        demo:ContactInfo_001
```
A fictional medical record class has this modeling:
This fictional medical record for an individual can be queried by SPARQL for subsections:

demo:MedicalRecord_001 demo:has_subsection+ ?x
But – SPARQL does not provide a way to query for classes whose individuals would be connected by a recursive property pattern.
But – SPARQL does not provide a way to query for classes whose individuals would be connected by a recursive property pattern

• In practice this limitation is overcome by punning (a pre-processing step), and then using SPARQL

• The purpose of OWL Paths is to enable SPARQL-like path processing over the classes in an ontology as if the pun individuals were present
Representing anatomy
The anatomy ontology that catalyzed this work is the Foundational Model of Anatomy (FMA) ontology

- The FMA models adult anatomy using over 100,000 classes and 100 relations.
- In development for 20 years by the Structural Informatics Group at the University of Washington
The anatomy ontology that catalyzed this work is the Foundational Model of Anatomy (FMA) ontology

• I have recently been funded to create a derivative called the Foundational Model of Human anatomy (FMHA)

• The FMHA is intended to be more consistent and complete in modeling than the FMA, and eventually to replace the FMA

• We will develop and implement knowledge representation patterns to construct and audit the FMHA
An example modeling a muscle (biceps brachii) and its relation to bones (scapula, radius)
An example modeling a muscle (biceps brachii) and its relation to bones (scapula, radius)

Biceps brachii

has constitutional part

Tendon of long head of biceps brachii

attached to

Supraglenoid tubercle

attached to

Glenoidal rim of scapula

attached to

Head of scapula

attached to

Lateral angle region of scapula

Tendon of short head of biceps brachii

attached to

Area of origin of short head of biceps brachii

attached to

Apical part of coracoid process

attached to

Coracoid process

Muscle body of biceps brachii

attached to

Radial tuberosity

attached to

Proximal epiphysis of radius

attached to

Radius

Bicipital aponeurosis
Implementation
The OWL Paths is a Java library built on the OWL API for processing property paths over OWL classes, in a similar manner as SPARQL is used to query using paths over OWL individuals

- OWL Paths grammar is syntactically similar to SPARQL
- Many but not all SPARQL constructs are supported
- The expressivity of OWL Paths needn’t be limited to that of SPARQL
<table>
<thead>
<tr>
<th>Syntax form</th>
<th>Matches</th>
<th>SPARQL</th>
<th>OWL Paths</th>
</tr>
</thead>
<tbody>
<tr>
<td>uri</td>
<td>A URI or a prefixed name. A path of length one.</td>
<td>✓</td>
<td>✓</td>
</tr>
<tr>
<td>^elt</td>
<td>Inverse path (object to subject).</td>
<td>✓</td>
<td></td>
</tr>
<tr>
<td>(elt)</td>
<td>A group path elt, brackets control precedence.</td>
<td>✓</td>
<td>✓</td>
</tr>
<tr>
<td>elt1 / elt2</td>
<td>A sequence path of elt1, followed by elt2</td>
<td>✓</td>
<td>✓</td>
</tr>
<tr>
<td>elt1 ^ elt2</td>
<td>Shorthand for elt1 / ^elt2, that is elt1 followed by the inverse of elt2.</td>
<td>✓</td>
<td></td>
</tr>
<tr>
<td>elt1</td>
<td>elt2</td>
<td>A alternative path of elt1, or elt2 (all possibilities are tried).</td>
<td>✓</td>
</tr>
<tr>
<td>elt*</td>
<td>A path of zero or more occurrences of elt.</td>
<td>✓</td>
<td>✓</td>
</tr>
<tr>
<td>elt+</td>
<td>A path of one or more occurrences of elt.</td>
<td>✓</td>
<td>✓</td>
</tr>
<tr>
<td>elt?</td>
<td>A path of zero or one elt.</td>
<td>✓</td>
<td></td>
</tr>
<tr>
<td>elt{n,m}</td>
<td>A path between n and m occurrences of elt.</td>
<td>✓</td>
<td></td>
</tr>
<tr>
<td>elt{n}</td>
<td>Exactly n occurrences of elt. A fixed length path.</td>
<td>✓</td>
<td></td>
</tr>
<tr>
<td>elt{n,}</td>
<td>n or more occurrences of elt.</td>
<td>✓</td>
<td></td>
</tr>
</tbody>
</table>
We can use the library to query over classes whose individuals would include paths containing zero or more sequential occurrences of an object property.
This is the code to set up the query and return the classes for the three subsections

```java
// set up subjects
Set<OWLClassExpression> subjects = new HashSet<>();
OWLClass start = factory.getOWLClass("https://purl.org/owl/demo/medicalrecords.owl#MedicalRecord");
subjects.add(start);

// process OWL paths
String path = "demo:has_subsection*";
PathExpression pe = new PathExpression(structural_reasoner);
Set<OWLClassExpression> results = pe.processPath(path, subjects);
```
In summary – just as SPARQL enables property path exploration between *individuals*, this library is a utility that enables similar exploration between *classes*.

We are using this processing capability to validate *knowledge representation patterns* in a large biomedical ontology.

Source code at

[https://gitlab.com/endless-forms-studio/owl_paths/](https://gitlab.com/endless-forms-studio/owl_paths/)

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