

OpenBiodiv-O Ontology: Bridging the Gap Between Biodiversity Data and Biodiversity Publishing

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Motivation & Aims

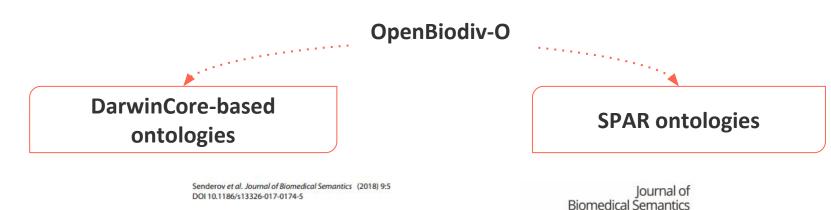
Open
Biodiversity
Knowledge
Management

Building a semantic knowledge graph from literature-extracted biodiversity data

Conceptual modelling of the biodiversity publishing domain



Introducing OpenBiodiv-O



RESEARCH Open Access

OpenBiodiv-O: ontology of the OpenBiodiv knowledge management system

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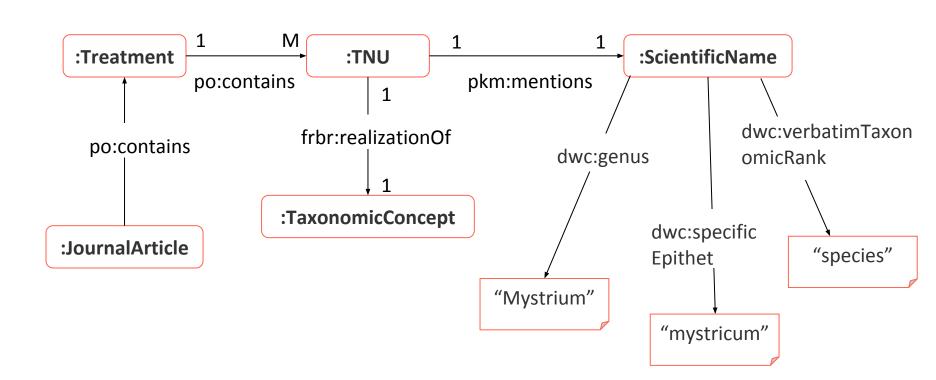


Biodiversity Publishing

- Taxonomic articles:
- Abstract, Introduction, Materials and Methods, Results, Conclusions
- Taxonomic treatment:
 - Nomenclature
 - Type material
 - Etymology
 - Diagnosis
 - Description
 - Distribution
- Metadata and various identifiers (DOI, ORCID, Zoobank ID)
- Taxonomic names



Modelling Taxonomic Names

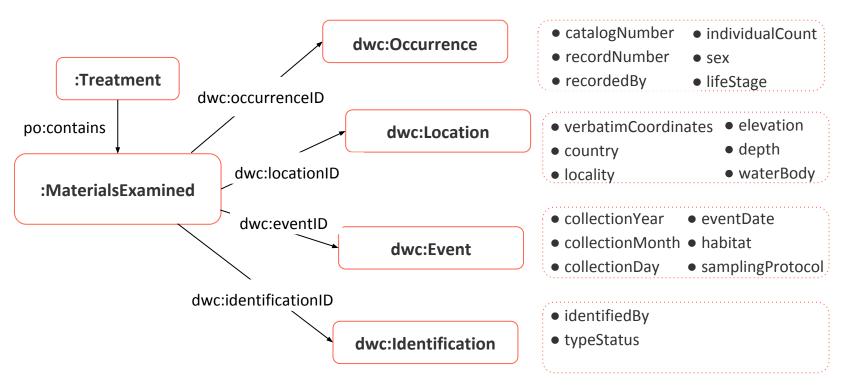




Bottom-up approach to modifying OpenBiodiv-O



1. Materials Examined







2. Institutional identifiers

GRSciColl: The Global Registry of Scientific Collections

http://biocol.org/urn:lsid:biocol.org:col:34985



dwc:institutionID	:Institu	:Institution	
:MaterialsExamined	:institutionName	dwc:inst	

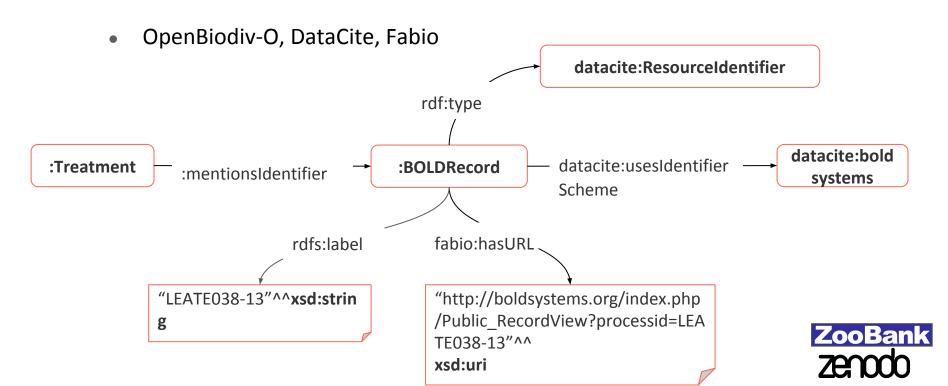
"KwaZulu-Natal

Museum"





3. Molecular resource identifiers





4. Personal identifiers



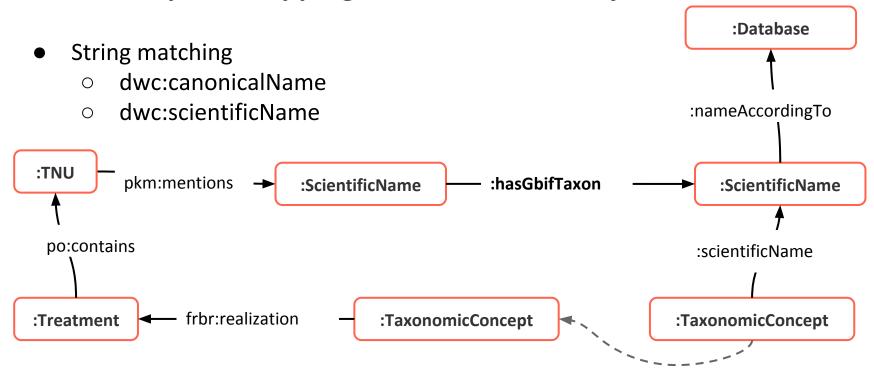
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openbiodiv:8B0A6890-3094-4431-8262-23748A86B071 rdf:type fabio:ResearchPaper;
      dcterms:creator openbiodiv:7F19D49E-B4CA-4E7A-808D-A57DAA7E02A3.
openbiodiv:7F19D49E-B4CA-4E7A-808D-A57DAA7E02A3 rdf:type foaf:Person;
      rdfs:label "Mengmeng Liu";
      openbiodiv:affiliation "College of Ecology, Lishui University, Lishui, Zhejiang, China";
      datacite:hasIdentifier orcid:0000-0002-0985-5852.
orcid:0000-0002-0985-5852 rdf:type datacite:PersonalIdentifier;
      datacite:usesIdentifierScheme datacite:orcid;
      rdfs:label "0000-0002-0985-5852".
```

ResearchPaper $\xrightarrow{1}$ \xrightarrow{M} Person $\xrightarrow{1}$ $\xrightarrow{1}$ PersonalIdentifier





5. Explicit mapping to GBIF's taxonomy







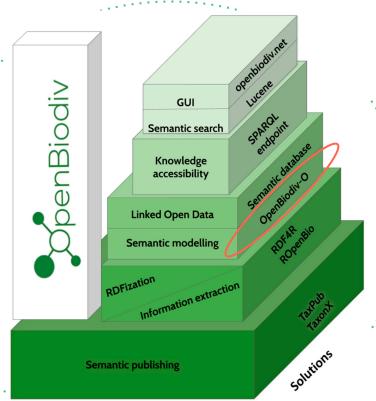
Use cases

- Modelling and linking resources across domains
 - Publishing
 - Taxonomy
 - Genomics
- Serving users from different groups
 - Taxonomists
 - Ecologists
 - Curators
 - Institutions



Application

PS 0101



Concepts









OpenBiodiv: Linking Type Materials, Institutions, Locations and Taxonomic Names Extracted From Scholarly Literature

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The OpenBiodiv project began in 2015 and set to establish a knowledge graph of biodiversity statements extracted from taxonomic articles, published by Pensoft and treatments, extracted by Plazi. Starting with a conceptualisation of the biodiversity publishing domain, several agreed vocabularies like DarwinCore and the SPAR ontologies were combined into a single ontology, OpenBiodiv-O. Semantic enhancement of articles, published in eXtensible Markup Language (XML) enabled their transformation into the machine-readable Resource Description Framework (RDF), which gave rise to the Linked Open Dataset. We also converted GBIF's backbone to RDF and mapped scientific names from taxonomic articles to it. Storing and managing linked statements in the OpenBiodiv knowledge graph allows easy traversal through the statements. This facilitates the answering of complex queries related to biodiversity and biodiversity publishing by institutions, taxonomists, curators, conservation experts and funding organisations.

Increasing accessibility of biodiversity knowledge to stimulate scientific research and conservation efforts.

Developing OpenBiodiv into a knowledge graph capable of answering complex biodiversity questions.

Figure 1: The OpenBiodiv architecture



Figure 2: Semantic relationships between resources in the graph database enable complex queries



Example use cases

Use Case	User Group	SPARQI
Find papers which describe a	Researcher	
taxon, the type material for which	Institution	
is held in the NHM	Funding body	
Find GenBank accession numbers associated with taxa with type material deposited in the NHM	Researcher	
Find the geographical coordinates, habitat and storing institutions, preserving specimens from the family Theraphosidae	Researcher Curator Institution Conservation expert Tarantula owner	
Find institutions storing type material specimens of the genus Prosopistoma from various	Researcher Curator Institution	

Table 1: Example use cases of OpenBiodiv and the user groups which might benefit from them. Answering these questions is done by executing the corresponding SPARQL queries at the OpenBiodiv SPARQL endpoint, available at: http://graph.openbiodiv.net/sparql. Here, you can execute the oueries corresponding to the last 2 questions yourself by scanning the QR codes and specifying the username and password to be both "biodiversity_next".

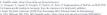
- Disambiguation of author names, taxon names, institution names and codes
- •GRSciColl institution list and Wikidata could provide data to disambiguate institutions
- · ORCID identification system could be used to disambiguate
- Information extraction (e.g. catalog numbers, institution abbreviations) from unstructured text
- · Different NLP techniques, such as gazetteer use, supervised machine learning and rule-based solutions, could be of help.

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Next steps

- Facilitate federated SPARQL queries by mapping external ontologies to OpenBiodiv-O.
 - Wikidata
- Expanding the ontology as the number of extracted entity types increases



Thank you!

Questions?



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