

# The eNanoMapper Ontology

An application ontology to enable data integration for nanomaterial risk assessment

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# Overview

## 1 Introduction

- eNanoMapper
- Engineered nanomaterials and safety
- Ontologies
- Ontologies - example of a class

## 2 eNanoMapper Ontology

- Design of the eNanoMapper ontology

## 3 Migrating the eNM ontology CI/CD

- Migrating the ontology development
- A. Migrating current setup to GitHub actions
- B. Ontology Development Kit
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# Overview

## ① Introduction

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- Ontologies - example of a class

## ② eNanoMapper Ontology

## ③ Migrating the eNM ontology CI/CD

# eNanoMapper

- ▶ **eNanoMapper** is a broader project which aims to address data and model interoperability challenges for engineered nanomaterial safety. [1]
- ▶ **The eNanoMapper ontology** is an application ontology and reuses parts of several ontologies to describe the full domain of nanomaterial safety assessment. [2]. It was continued by NanoCommons and NanoSolveIT.

Figure 1: The eNanoMapper logo



# Engineered nanomaterials and safety

- ▶ Engineered nanomaterials (eNMs) are broadly defined as compounds that exist on a scale of 1–100 nm. in at least one of their dimensions. [2]
- ▶ Their safety assessment must cover the identification of:
  - Physicochemical properties
  - Biological properties and activity [1]
  - Diffusion behaviour into the natural environment [2]

# Ontologies

- ▶ A formal representation of a set of concepts and the rules constraining how they become structured within a knowledge domain. [3]
- ▶ They can be used to provide metadata (a computer-readable set of information that describes other data), or to reason over a domain.
- ▶ It consists of three syntactical categories: **Entities**, **Expressions** and **Axioms**, which can be given annotations for further description. [4]

# Ontologies

- All entities (classes, object properties, named individuals...) are uniquely identified by a sequence of characters called IRI (International Resource Identifier, an extension of URI).

Figure 2: An owl:Class in an ontology text file (the ontology document). IRIs in blue.

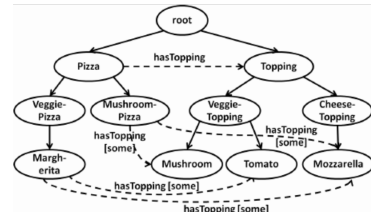
```
<!-- http://purl.obolibrary.org/obo/PATO_0001464 ->

<owl:Class rdf:about="http://purl.obolibrary.org/obo/PATO_0001464">
  <rdfs:subClassOf rdf:resource="http://purl.obolibrary.org/obo/PATO_0001018"/><rdfs:subClassOf
  rdf:resource="http://purl.obolibrary.org/obo/BFO_0000016"/>
  <obo:IAO_0000115 rdf:datatype="http://www.w3.org/2001/XMLSchema#string">A quality that is equal to the
  potential energy per unit charge associated with a static (time-invariant) electric field, also called the electrostatic
  potential.</obo:IAO_0000115>
  <oboInOwl:hasOBONamespace rdf:datatype="http://www.w3.org/2001/XMLSchema#string
  ">quality</oboInOwl:hasOBONamespace>
  <oboInOwl:id rdf:datatype="http://www.w3.org/2001/XMLSchema#string">PATO:0001464</oboInOwl:id>
  <oboInOwl:inSubset rdf:resource="http://purl.obolibrary.org/obo/pato#attribute_slim"/><oboInOwl:inSubset
  rdf:resource="http://purl.obolibrary.org/obo/pato#scalar_slim"/>
  <rdfs:label rdf:datatype="http://www.w3.org/2001/XMLSchema#string">electric potential</rdfs:label>
</owl:Class>
```

# Ontologies

- ▶ Most ontologies use the W3C standard language for ontologies, Web Ontology Language **OWL**.
- ▶ OWL ontologies are mainly stored in .owl files, which are a sort of **RDF** document.
- ▶ **RDF** (Resource Description Framework) is a standard for data exchange. It defines **triples** of (**subject**, **predicate**, **object**). [5]
- ▶ These triples form labeled graphs where the edge (predicate) represents the link between two resources (subject and object)

Figure 3: The pizza ontology, visualized as a graph [6]





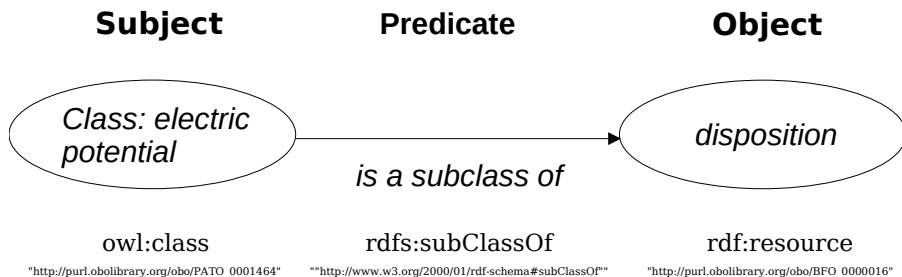
# Ontologies - example of a class

Figure 4: A view of the eNanoMapper ontology class hierarchy, showing the class "electric potential"

The screenshot displays the eNanoMapper ontology interface. The left pane shows the class hierarchy under 'owl:Thing', with 'entity (BFO:0000001)' as the parent. Under 'entity', there are several subclasses, including 'disposition', 'amphiphilic', 'bioavailability', 'biocompatible', 'biodegradable', 'electric potential' (highlighted), 'zeta potential', 'electronegativity', 'function', 'Oxidation-Reduction Potential', 'role', 'surface property', 'information content entity', 'material entity', 'process', and 'quality'. The right pane shows the details for the 'electric potential' class, including its label, ID (PATO:0001464), OBO namespace, quality, definition, and database cross-reference (Wikipedia: [http://en.wikipedia.org/wiki/Electric\\_potential](http://en.wikipedia.org/wiki/Electric_potential)). It also shows 'in\_subset' relationships with 'attribute\_slim' and 'scalar\_slim'.

# Ontologies - example of a class

Figure 5: A graph with two nodes (Subject and Object) and a triple connecting them (Predicate): the class with class description "electric potential" is a subclass of disposition.



# Ontologies - example of a class

Figure 6: The triple in the previous figure as expressed in the .owl document file of the ontology it is contained in.

```
<!-- http://purl.obolibrary.org/obo/PATO_0001464 -->  
  
<owl:Class rdf:about="http://purl.obolibrary.org/obo/PATO_0001464">  
<rdfs:subClassOf rdf:resource="http://purl.obolibrary.org/obo/BFO_0000016"/>  
  
(...)  
</owl:Class>
```

# Ontologies - example of a class

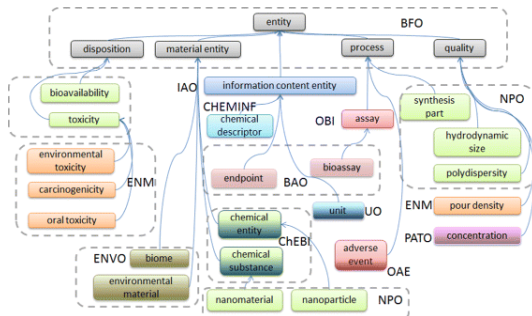
- ▶ **Foundation/Upper/Top-level ontologies**
- ▶ **Application and domain ontologies**

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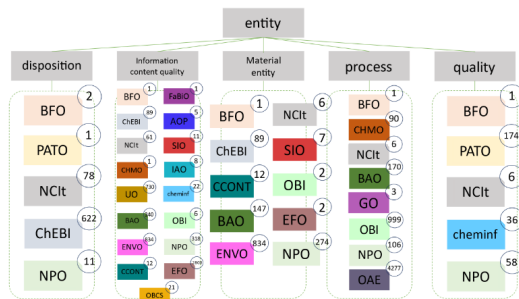
# Design of the eNanoMapper ontology

Figure 7: An overview of the upper levels, imports and manually annotated content going into the eNanoMapper ontology [2]



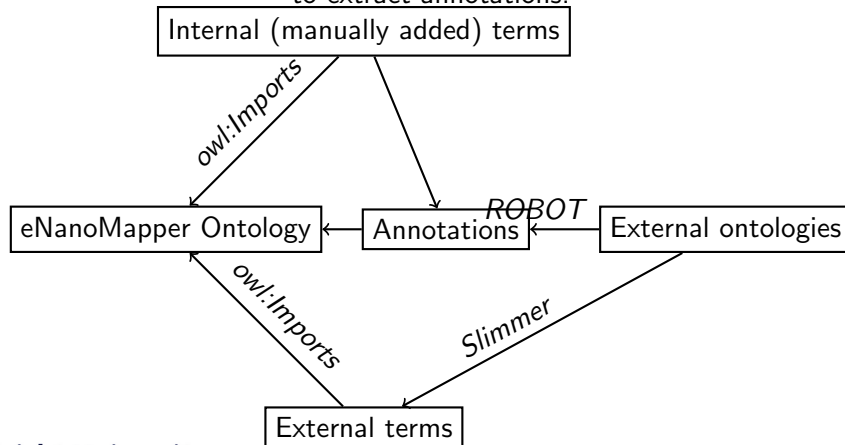
# Design of the eNanoMapper ontology

Figure 8: Distribution of the number of classes imported from each ontology into eNM [7]



# Design of the eNanoMapper ontology

The current approach uses in-house software (*Slimmer*) to slim imports, and ROBOT to extract annotations.





# Design of the eNanoMapper ontology

- ▶ **SLIMMER** Uses the OWLAPI to create slims of the ontologies to be imported, and determine the class hierarchy of the imports. [2]
- ▶ **ROBOT** is a wrapper for the OWL API used by developers in the Open Biomedical Ontologies (among others) for development and quality control. [8]

# Design of the eNanoMapper ontology

Figure 9: The current Jenkins setup for eNanoMapper ontology CI/CD

The screenshot shows the Jenkins web interface for the 'Project eNanoMapper Ontology'. The top navigation bar includes 'Dashboard', 'eNanoMapper', and 'eNanoMapper Ontology'. A search bar and a 'Log in' link are also present. The left sidebar contains links to 'Back to Dashboard', 'Status', 'Changes', 'Git Polling Log', 'GitHub', 'Open Blue Ocean', and 'Embeddable Build Status'. The main content area is titled 'Project eNanoMapper Ontology' and features a 'Recent Changes' section with a code icon. Below this is a 'Downstream Projects' section listing various projects, each with a status icon (green circle with a checkmark or red circle with an 'X') and a name. The list includes: Wikipathways - CHEBI, eNanoMapper - AOP, eNanoMapper - BAO, eNanoMapper - BAO - properties, eNanoMapper - BFO, eNanoMapper - BFO, eNanoMapper - CCONT, eNanoMapper - CHEBI, eNanoMapper - CHEMINF, eNanoMapper - CHMO, eNanoMapper - CLO, eNanoMapper - CTO - properties, eNanoMapper - GFO, eNanoMapper - ENVO, eNanoMapper - FAIR, eNanoMapper - GO, eNanoMapper - HUPSON, eNanoMapper - IAO, eNanoMapper - NFO, eNanoMapper - NFO - properties, eNanoMapper - OAE, eNanoMapper - OBCS, eNanoMapper - OH, eNanoMapper - PATO, eNanoMapper - RO - properties, eNanoMapper - SDO, eNanoMapper - SDO - properties, eNanoMapper - UBERON, eNanoMapper - UO, and eNanoMapper - CHEMINF - properties. The bottom of the page shows a 'Build History' section with a search bar and a list of builds, each with a status icon, a number, and a timestamp.

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# Migrating the ontology development

- ▶ Current setup: Jenkins for CI in own server (to be dropped soon)
- ▶ Alternatives:
  - Literal migration from Jenkins to GitHub actions
  - Migration to the Ontology Development Kit-ROBOT - rethinking the whole process

# A. Migrating current setup to GitHub actions

Figure 10: A test run for the GitHub actions-based setup for eNanoMapper ontology CI

jmilianacosta / eNM-ontology-build Private

<> Code Issues 2 Pull requests Actions Projects Security Insights Settings

workflow slim ontologies workflow slim ontologies #13

Summary

Jobs

- get-scripts
- slim-bao
- slim-efo
- slim-ecct
- slim-cheminf
- slim-chmo
- slim-efo
- slim-dabio
- slim-go
- slim-ao
- slim-ncit
- slim-mpo
- slim-nae
- slim-obcs
- slim-obi

Manually triggered 2 months ago Status Success Total duration 2m 12s Artifacts 17

slim-ontologies.yml on: workflow\_dispatch

```

graph LR
    get-scripts --> slim-bao
    slim-bao --> slim-efo
    slim-efo --> slim-ecct
    slim-ecct --> slim-cheminf
    slim-cheminf --> slim-chmo
    slim-chmo --> slim-efo
    slim-efo --> slim-dabio
    slim-dabio --> slim-go
    slim-go --> slim-ao
    slim-ao --> slim-ncit
    slim-ncit --> slim-mpo
    slim-mpo --> slim-nae
    slim-nae --> slim-obcs
    slim-obcs --> slim-obi
  
```

## A. Migrating current setup to GitHub actions

- ▶ A toolkit developed for the development of biomedical ontologies following the software development approach (a method preceded by the eNanoMapper/jenkins/Slimmer setup)
- ▶ It is delivered in a docker image and uses ROBOT (a wrapper for the OWL API) and a series of scripts to standardize and automate steps like preparing ontology releases, continuous quality control checking, and dependency management. [9]

Figure 11: The ODK logo

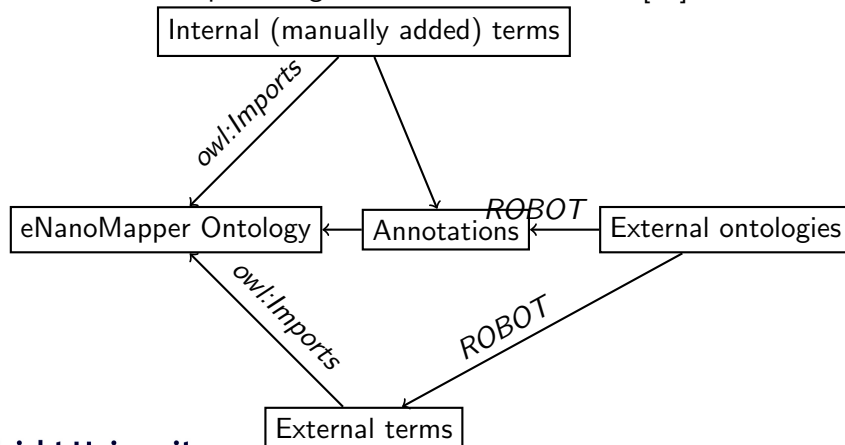


## A. Migrating current setup to GitHub actions

The challenge: replicating the same class hierarchies after substituting Slimmer with ROBOT.

## B. Ontology Development Kit

The ODK approach relies on ROBOT to handle ontologies and serialization of several steps through standardized Makefiles. [10]





# Comparison

Table 1: Comparison of the two alternatives for eNanoMapper CI/CD

	<b>GitHub actions</b>	<b>Ontology Development Kit</b>
<b>Uses tools</b>	GH actions, slimmer, maybe ROBOT for annotations, in-house scripts	ROBOT and ODK scripts
<b>Keeps current class hierarchy</b>	Yes	Perhaps, but will be a challenge
<b>Sticks to OBO recommendations</b>	Yes, after minimal adaptations if Slimmer is kept	Yes
<b>Hosted on</b>	GitHub	GitHub
<b>Time to develop</b>	Ready at any point	Not optimal for migrating existing ontologies [10]

# References I

- [1] Nina Jeliaskova, Charalampos Chomenidis, Philip Doganis, Bengt Fadeel, Roland Grafström, Barry Hardy, Janna Hastings, Markus Hegi, Vedrin Jeliaskov, Nikolay Kochev, Pekka Kohonen, Cristian R Munteanu, Haralambos Sarimveis, Bart Smeets, Pantelis Sopasakis, Georgia Tsiliki, David Vorgrimmmler, and Egon Willighagen. The eNanoMapper database for nanomaterial safety information. *Beilstein Journal of Nanotechnology*, 6:1609–1634, July 2015. ISSN 2190-4286. doi: 10.3762/bjnano.6.165. URL <https://www.beilstein-journals.org/bjnano/articles/6/165>.
- [2] Janna Hastings, Nina Jeliaskova, Gareth Owen, Georgia Tsiliki, Cristian R Munteanu, Christoph Steinbeck, and Egon Willighagen. eNanoMapper: harnessing ontologies to enable data integration for nanomaterial risk assessment. *Journal of Biomedical Semantics*, 6(1):10, December 2015. ISSN 2041-1480. doi: 10.1186/s13326-015-0005-5. URL <http://www.jbiomedsem.com/content/6/1/10>.

## References II

- [3] Markus Krötzsch, Frantisek Simancik, and Ian Horrocks. A Description Logic Primer, June 2013. URL <http://arxiv.org/abs/1201.4089>. arXiv:1201.4089 [cs].
- [4] OWL - Semantic Web Standards, . URL <https://www.w3.org/OWL/>.
- [5] RDF - Semantic Web Standards, . URL <https://www.w3.org/RDF/>.
- [6] Nick Drummond and Alan Rector. Pizza Ontology. URL <https://protege.stanford.edu/ontologies/pizza/pizza.owl>.
- [7] Laurent Winckers, Chris Evelo, and Egon Willighagen. Expanding the enanomapper ontology - icbo2020, September 2020. URL <https://doi.org/10.5281/zenodo.4032809>.

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- [8] Rebecca C. Jackson, James P. Balhoff, Eric Douglass, Nomi L. Harris, Christopher J. Mungall, and James A. Overton. ROBOT: A Tool for Automating Ontology Workflows. *BMC Bioinformatics*, 20(1):407, December 2019. ISSN 1471-2105. doi: 10.1186/s12859-019-3002-3. URL <https://bmcbioinformatics.biomedcentral.com/articles/10.1186/s12859-019-3002-3>.
- [9] Nicolas Matentzoglou, Damien Goutte-Gattat, Shawn Tan, James Balhoff, Seth Carbon, Anita Caron, William Duncan, Joe Flack, Melissa Haendel, Nomi Harris, William Hogan, Charles Hoyt, Rebecca Jackson, HyeongSik Kim, Huseyin Kir, Martin Larralde, Julie McMurry, James Overton, Bjoern Peters, and David Osumi-Sutherland. Ontology development kit: a toolkit for building, maintaining, and standardising biomedical ontologies, 07 2022.

## References IV

- [10] Nicolas Matentzoglou, Chris Mungall, and Damien Goutte-Gattat. Ontology Development Kit, July 2021. URL <https://github.com/INCATools/ontology-development-kit/issues/298>.