

Enhancing the AOP-Wiki usability and accessibility with semantic web technologies

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Introduction

There is a need for faster and more efficient use of existing data to assemble effective assessment strategies for the ever growing number of chemicals. Therefore, a framework to organize existing mechanistic information, the Adverse Outcome Pathway (AOP), was introduced. [1] The main repository for such AOPs is the AOP-Wiki. However, it is challenging to automatically and systematically parse, filter, and use its captured knowledge.

Objective

We explored the use of semantic web technologies to link the AOP-Wiki with chemical and biological databases, allowing more detailed exploration of the database, thereby better supporting risk assessment workflows.

Results

This exercise resulted in a conversion of the complete AOP-Wiki into an RDF schema (Figure 1) that includes over 66,000 triples. 16 ontologies were used for the semantic annotations, and over 6,000 persistent identifiers of 20 types are included. Also, the APOPO was used to annotate over 4,000 components, and the addition of the AOP-DB RDF introduced over 5,000 diseases, 3,000 pathways, and 1,000 ToxCast assays related to relevant genes.

Conclusion

The created RDF and its accessibility through a SPARQL endpoint and Web API assist in the expansion and usability of the knowledge of the AOP-Wiki and AOP-DB. Furthermore, the use of ontologies and persistent identifiers allow new ways to explore the AOP knowledge, and makes the integration of this database in workflows possible. For example, federated SPARQL queries or integration in Jupyter notebooks can answer complex questions that require multiple information sources.

References

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Methods

- The AOP-Wiki XML was parsed and converted into Resource Description Framework (RDF) using a custom Python script.
- All parts of the AOP-Wiki, and the relationships between them, were semantically annotated using ontologies and standard vocabularies such as APOPO, CHEMINF, RDFS, and DC.
- The AOP, KE, KER and Stressor were registered in the MIRIAM Registry [2] and the resulting resolvable Uniform Resource Identifiers (URIs), along with the ones for chemicals and text-mapped genes, were integrated in the RDF.
- The identifier mapping service BridgeDb [3] was used to add matching identifiers for chemicals and genes of twelve additional databases.
- Similarly, parts of the AOP-DB [4,5] were converted in RDF as part of the OpenRiskNet implementation challenge.

