OLSP Ontology working group

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OLSP Ontology Working Group (2020~)

Promoting the interoperability of life science data at RIKEN through the development of a "Shared Vocabulary"

- Conduct a survey on the usage and demand for shared vocabularies, such as ontologies and public IDs.
- Examine and propose actionable strategies to accelerate data interoperability for cross-domain integration.
 - BDR Itoga, H., Wang, FF., Onami, S.
 - CSRS Takahashi, M., Arita, M.
 - CBS Okumura, Y., Imaizumi, I.
 - BRC Kushida, T., Masuya, H.*
 - **R-IH** Kato, M., Yamagata, Y., Kobayashi, N.**
 - RAP Yokota, H.
 - IMS Kasukawa, T.**

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A survey of research topics covered by RIKEN's database (2021)



Summary of the survey

- Overall, it is rich in variety, but the topics are relatively concentrated within each center
- Many datasets contains genes, but they are not interconnected due to variations such as species differences
- To connect datasets from RIKEN, it is essential to introduce public data for the life sciences
- Ortholog data is considered useful for linking RIKEN data and demonstrating its utility in disease research and other applications
- OMA (<u>https://omabrowser.org/oma/</u>) developed in SIB, which is available as RDF, is a strong candidate of the ortholog data
- (We collaborate with SIB through RIKEN hackathons)



An example in RIKEN BRC (using RDF)



RIKEN has a data integration platform using RDF.

RIKEN Metadatabase (2016-) https://metadb.riken.jp/

RIKEN MetaDatabase This database portal site is used to provide information on RIKEN's various life science databases to order to help researchers around the world make full use of RIKEN's research results.		
Databases	🗎 Ontologies 🗕 About RIKEN MetaDatabase < Related MetaDatabase	•
Q	Search	Faceted Navigation (for databases with metadata)
Search option		C Species
123 Database(s) Found.		C Theme
1 2 3 4 5 6 7 (All)		Publisher
		Datalype
a <u> RIKEN Data</u>	base Catalog http://metadb.riken.jp/db/DBcatalog	Related Projects
& Classes	List of databases + RIKEN organizations or centers + Topics + Data types +	🕐 RIKEN Open Science Portal
Database		RIKEN HACKATHON
FANTOM5 MetaDatabase <u>http://metadb.riken.jp/db/fantom5</u>		2020
& Classes	FANTOM5 Data >	RIKEN Open Life
	Library Cage Library RNAseq Library SRNAseq Library Cagescan Library	Related MetaDatabase
	Sample Sample By Type Strain Quality Control Time Course	Aliken Microstructural Imaging Metadatabase
	Protocol RNA Extraction Protocol RNA Protocol Description File	Plant Metabolome MetaDatabase
S FANTOM5 CAGE Peak Annotation http://metadb.riken.jp/db/fantom5_cage_peak		
& Classes	CAGE Peak of Mouse + Annotation of Mouse + Transcript of Mouse + CAGE Peak of Human + Annotation of Human + Transcript of Human +	
refTSS MetaDatabase http://metadb.riken.jp/db/refTSS		
& Classes	Project Organization Project File Experiment	
	Human Human refTSS 1 + Human refTSS 2 + Human refTSS 3 + Human Gene Annotation 1 + Human Gene Annotation 2 + Human Gene Annotation 3 + Human Transcript 1 + Human Transcript 2 + Human Transcript 3 + Human Regulatory 1 + Human Regulatory 2 + Human TatAbox Annotation 1 + Human TAtAbox Annotation 2 + Human TATAbox Annotation 3 + Human TSS Source 1 + Human TSS Source 2 + Human TSS Source 3 + Human TSS Classification 1 + Human TSS Classification 2 + Human TSS Classification 3 + Human TSS Classification 3 + Human TSS Classification 2 + Human TSS Classification 3 +	
	Mouse Mouse refTSS 1 , Mouse refTSS 2 , Mouse Gene Annotation 1 , Mouse Gene Annotation 2 , Mouse Transcript 1 , Mouse Transcript 2 , Mouse Regulatory 1 , Mouse Regulatory 2 , Mouse TATAbox Annotation 1 , Mouse TATAbox Annotation 2 , Mouse TSS Source 1 , Mouse TSS Classification 1 , Mouse TSS Source 2 , Mouse TSS Classification 2 ,	

Report: How should we address URI inconsistencies in the RIKEN MetaDB? Masuya, H., Yamamot, Y., Kushida, T., Kato, M (2024) https://docs.google.com/document/d/1oMXUrNCK2oc5GuuJNbU_93jiOF-X0Ao87hOyLUt0tB8/edit?usp=sharing

A key issue in RDF-based data integration is the failure to connect data due to URI inconsistencies. This report proposes practical solutions for addressing URI inconsistencies in RIKEN MetaDB operations for the RIKEN OLSP.

Conclusion:

- Using the data registration system of the RIKEN MetaDB, correcting URIs during data registraiton is efficient considering cost and performance after the change.
- However, it will require rewriting existing data.
- The modification targets should be limited (e.g., genes, species, proteins, diseases).
- Overview of the modification algorithm: A conversion table for the modification targets will be prepared. After correcting typos with tools like RDFDoctor, the conversion table will be referenced to correct any inconsistencies.

Development of a system on the OLSP website to offer introductions on the RIKEN's databases (2023)



Development of a system on the OLSP website to offer introductions on the RIKEN's databases (2023-2025)



Ongoing (2024-2025)...

- Installing the system to the server
- Preparing metadata of databases



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