

Towards FAIR metadata standards in FAIRsharing.org

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Inaugural Workshop Metadata 4 Machines (M4M)
GO-FAIR initiative

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FAIRsharing.org

A curated, informative and educational resource on data and metadata *standards*, inter-related to *databases* and data *policies*

We guide **consumers** to **find**, **select** and **use** these resources with confidence, and **producers** to make their resource more **discoverable**, more widely **adopted** and **cited**

Researchers in academia, industry, government

Journal publishers or organizations with data policy

Learned societies, unions and associations



Developers and curators of resources

Research data facilitators, librarians, trainers

Funders and data policy makers

Pre-print at: https://doi.org/10.1101/245183

Authored by 68 stakeholders in academia, industry, funding agencies, standards organisations, infrastructure providers and scholarly publishers

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FAIRsharing and the FAIRmetrics

- FAIRsharing works with the FAIR metrics WG to serve as:
 - Registry to describe digital assets, such as databases/repositories, standards, policies, enhancing their discoverability (schema.org) and citability (DOIs)
 - Look up service for identifier schemas and standards (phase 1: now)
 - Validation service against metadata standards (phase 2: planned)

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A design framework and exemplar metrics for FAIRness

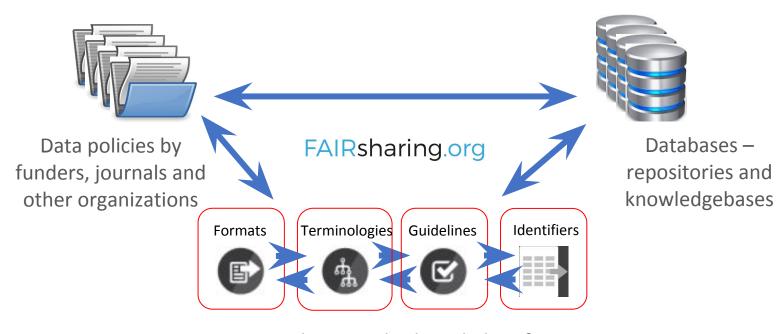
Mark D. Wilkinson [™], Susanna-Assunta Sansone [™], Erik Schultes, Peter Doorn, Luiz Olavo Bonino da Silva Santos & Michel Dumontier [™]

FAIRmetrics.org

also part of:



Mapping a complex and evolving landscape...

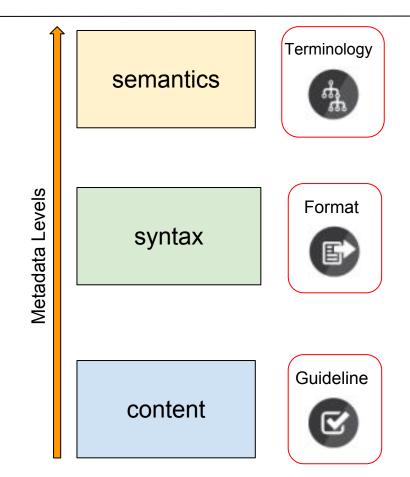


Metadata standards and identifiers

Medata standards in practice

Databases/ Repositories





Medata standards in practice Databases/ **GEO** ArrayExpress **Gene Expression Omnibus** Repositories DATABASE DATABASE **Terminology** semantics **Experimental Factor Ontology** TERMINOLOGY ARTIFACT Metadata Levels **Format** syntax MAGE-TAB MINIML MicroArray Gene Expression Tabular Format MIAME Notation in Markup Language MODEL/FORMAT MODEL/FORMAT Guideline content MIAME Minimum Information About a Microarray Experiment REPORTING GUIDELINE

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Reporting guidelines

a.k.a. Minimum Information Checklists

https://fairsharing.org/standards/?q=&selected_facets=type_exact:reporting%20guideline

- There are 151 reporting guidelines registered in FAIRsharing.org
 - The majority of them are described through manuscripts
 / documents (usually PDF or Word files)
 - Only some of those reporting guidelines have been described in a machine readable and verifiable way (e.g XML schema, YAML, JSON schema) e.g. MIBIG, MIAPTE, MICEE, MIDE, MIABIS, MIACA, MIACME, AIRR
 - Some guidelines have been implemented in different formats and those need to be taken into account

Towards FAIR metadata standards

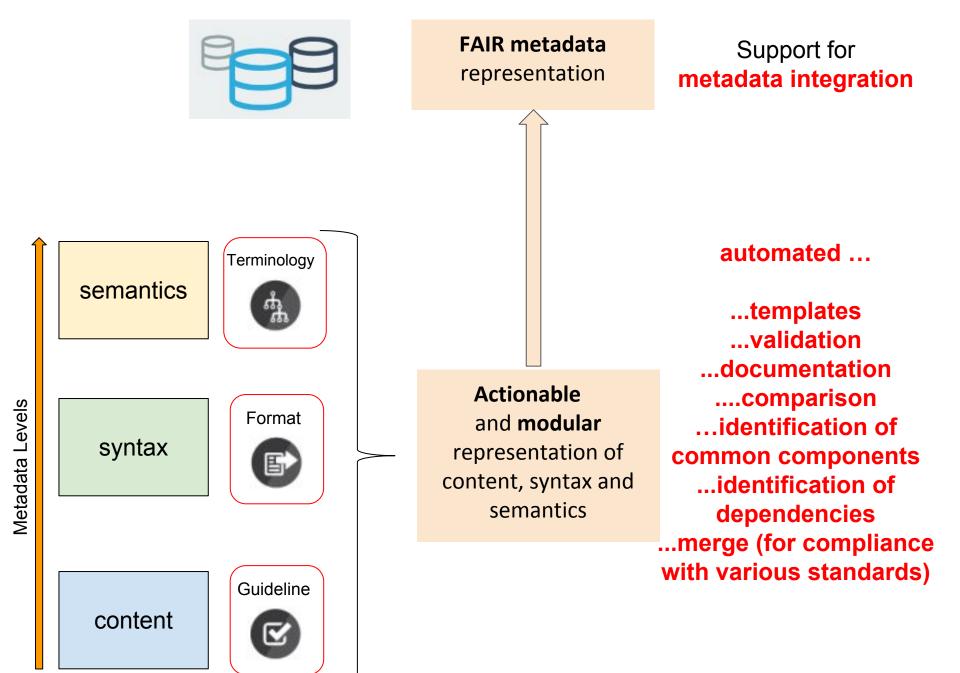
Machine actionable metadata standard components:

- Facilitate the creation of compliant metadata through automated templates
- Automate verification of compliance with the guidelines through automated validation (a.k.a. FAIR metric)
- Automated generation of documentation about the guidelines
- Automated syntactic and semantic comparison between reporting guidelines

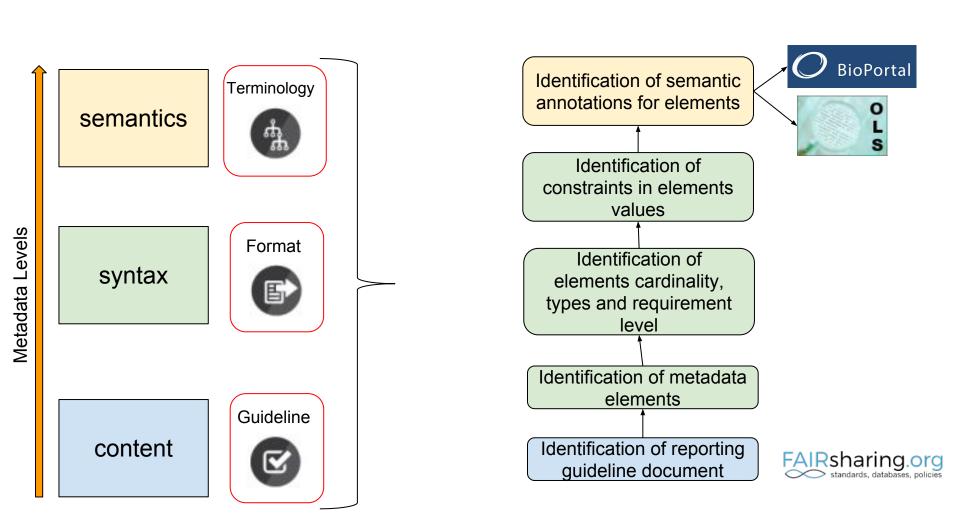
Reusable (modular) metadata standard components:

- Automatic identification of common reusable components and differences
- Automatic identification of dependencies between guidelines
- Automated creation of templates that comply with a two or more reporting guidelines (automated merge)

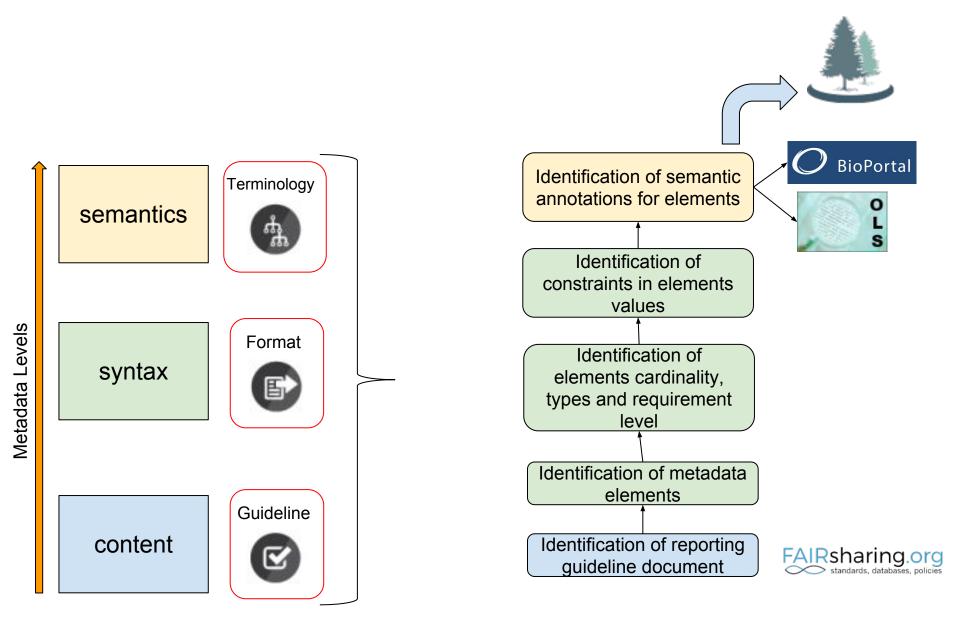
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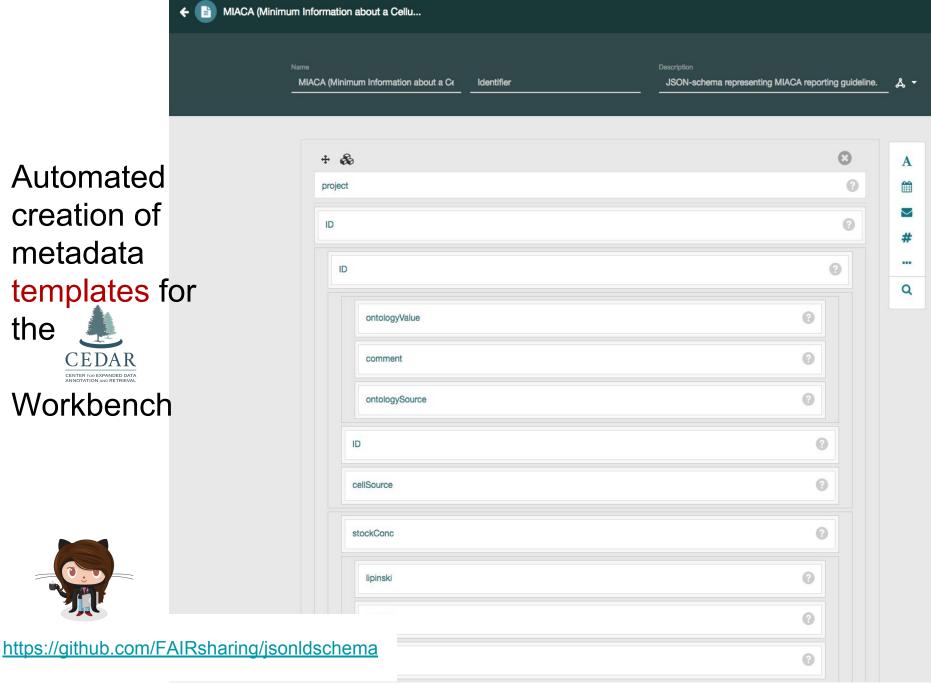
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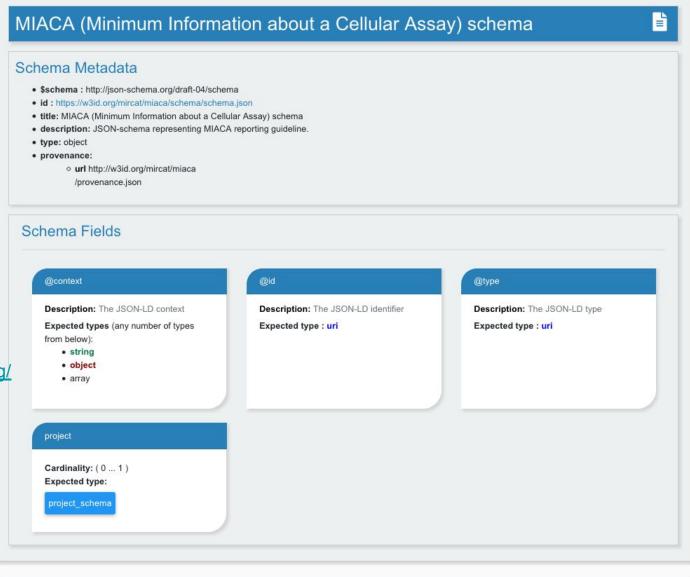
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Automating standards documentation

https://github.com/FAIRsharing/ JSONschema-documenter





MIACA (Minimum Information about a Cellular Assay) project schema

Schema Metadata

Automating the syntactic and semantic comparison between checklists



sdo:CreativeWork MIACA (Minimum Information about a Cellular Assay) schema MIACME (Minimum Information about a Cell Migration Experiments) schema sdo:Person MIACA Source - corresponding to a person and their organization MIACME person schema. A human being. sdo:addressCountry country X sdo:identifier ID identifier sdo:name name X sdo:address address address sdo:adstName X lastName sdo:dastName X fax sdo:faxNumber X fax sdo:givenName X firstName sdo:givenName X firstName sdo:description MIACA (Minimum Information about a Cellular Assay) project schema MIACME investigation schema sdo:description projectDescription description sdo:member source X sdo:instrument X contacts sdo:name X tittle		MIACA	MIACME	
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Take-home messages

- FAIRsharing.org is an important resource for helping in the analysis and convergence of metadata standards to implement the FAIR principles
- FAIRsharing is part of GO-FAIR and RDA
- FAIRsharing works in support of the development of FAIR metrics
 - Look up service
 - Validation service
- FAIRification of metadata standards
 - Actionable and modular representation of reporting guidelines, formats and their semantic requirements and constraints
 - Automated templates, validation, documentation, comparison, identification of commonalities and dependencies, merging







Team is funded by







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KINGDOM



Our team works to enabling reproducible research and FAIR science, driving science and discoveries



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Alejandra Gonzalez-Beltran, PhD Research Lecturer



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Delphine Dauga, PhD *Biocurator Contractor*



Ramon Granell Research Software and Knowledge Engineer



Dominique Batista Research Software and Knowledge Engineer

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- NIH CEDAR: https://metadatacenter.org
- NIH Data Commons Pilot: https://commonfund.nih.gov/commons

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Support for metadata integration

semantics



JSON-LD context files



syntax

Metadata Levels



{**x**}

JSON

line schemas (for JSON-LD)

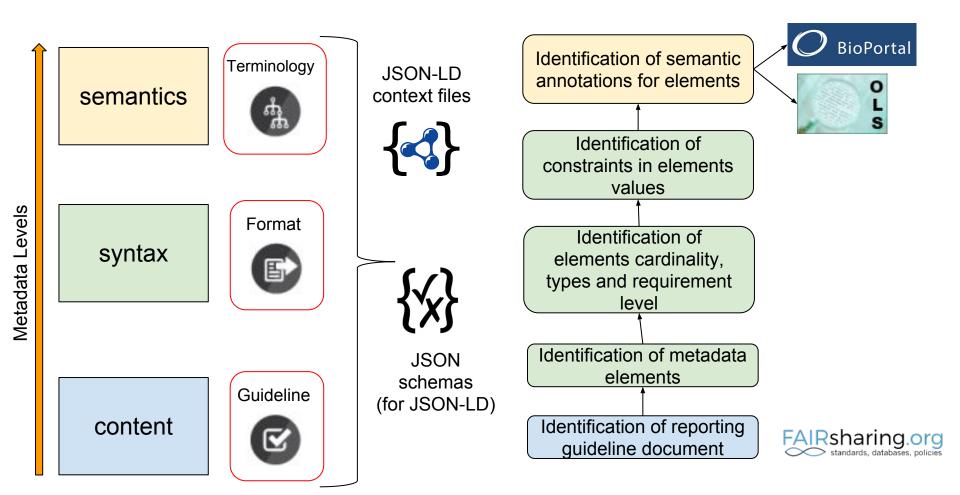
content



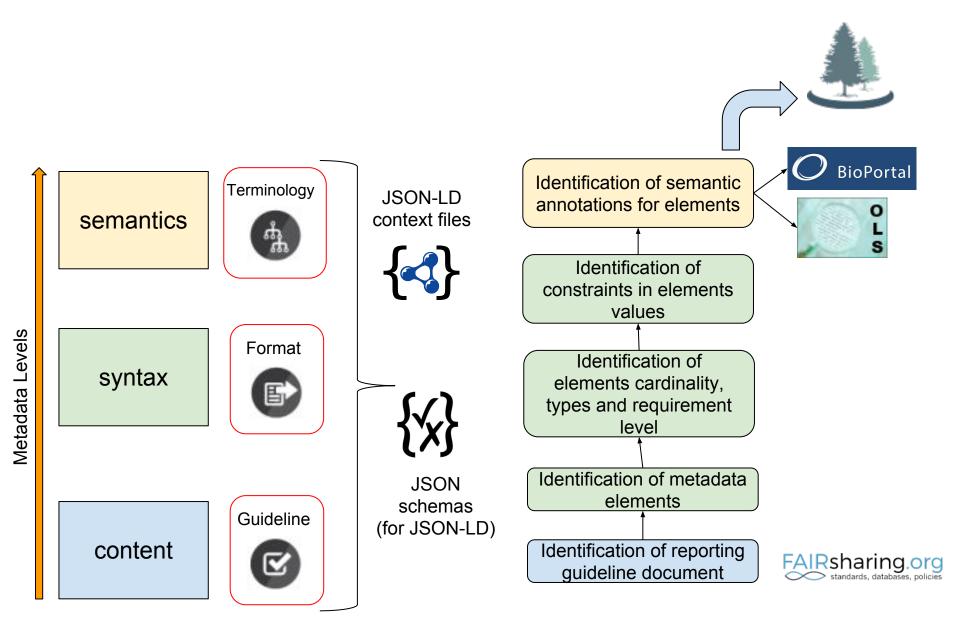
automated ...

...templates
...validation
...documentation
....comparison
...identification of
common components
...identification of
dependencies
...merge (for compliance
with various standards)

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