

FAIR Data APIs in the FISH (FAIR in Vivo Data Sharing) Platform

SEMANTICS Vienna 2022 - Onto4FAIR Workshop

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F.A.I.R - what does it mean ?

FAIR: Stands for

- Findable: To be Findable any Data Object should be uniquely and persistently identifiable
- Accessible: Data is Accessible in that it can be always obtained by machines and humans
- Interoperable: To be machine readable and based on shared metadata
- Reusable: To be Reusable Data Objects should be well described to be automatically linked with other data and refer to their sources

taken from GUIDING PRINCIPLES FOR FINDABLE, ACCESSIBLE, INTEROPERABLE AND RE-USABLE DATA PUBLISHING VERSION B1.0: https://www.force11.org/fairprinciples



Why do we care about FAIR

- Efficient organization, storage, retrieval and access of data is crucial to our business success as a Pharma and Diagnostics company
 - Enabler for generation of new insights from large data sets
 - Estimated [1] €10.2 billion is lost every year in research alone due to insufficient data management
- As a globally operating organization, we created or internalized a large amount of data in the past
- Every year our data output has been growing exponentially in terms of volume and variety.
 - Capability to properly master our value-driven data management processes is necessary for our future business success evolving towards a data-centric company.

Expected cost savings and gains in productivity enabled by proper data management are crucial to realize Roche's 10-years-ambition - to create at least
 11 Costwice the benefits for patients while reducing costs to society by 50% or more ³

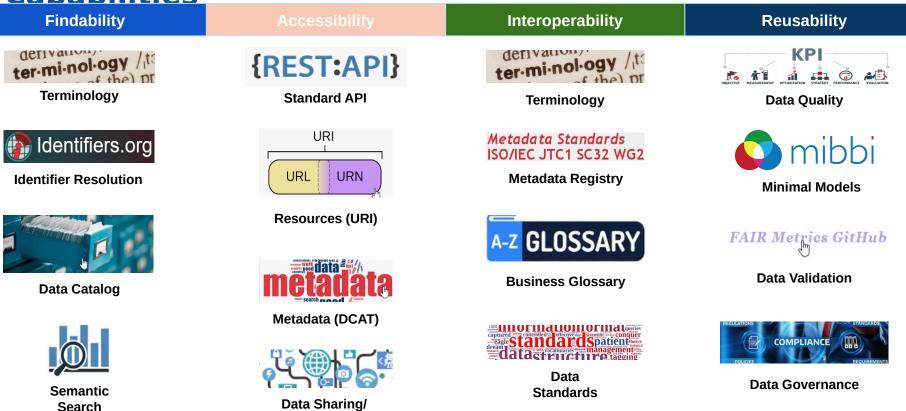


Approach

- Apply best data practices in order to build FAIR(er) applications: example FISH
- Implement best practices in a tool (RTS) that
 - Integrates them by leveraging on dependencies and existing information
 - Hosts reference data
 - Hides technical details behind a user interface → lowers the hurdle for practical implementation of FAIR practices
 - Main functionalities
 - URI namespace service
 - Terminology management
 - Reference Model management
 - Application-specific model management
 - ISON_ID interface specification for FAID data ADIS

RTS FAIR - Informatics & Organizational <u>Capabilities</u>

Data Protection

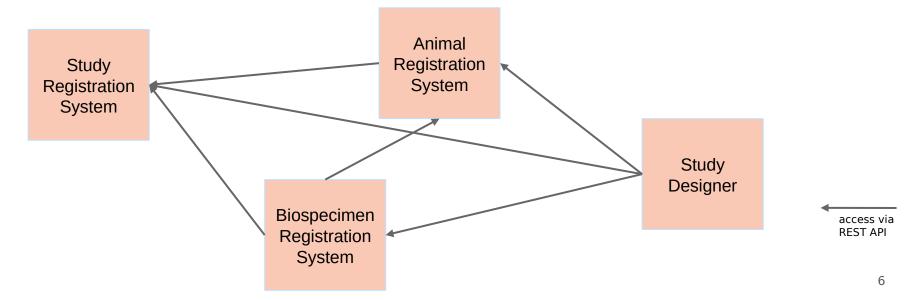


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FAIR in vivo data sharing platform (FISH) System topology

- Component-based Architecture
- RESTful APIs
- Relational DBs per component





Use Open Standards

Recommendations for content and code

- Open and well-defined standards for file formats and knowledge representation
- Eliminates risk of a vendor lock-in
- Secures future reusability and interoperability
- Notable Standards
 - HTTPS as secure content transfer protocol and RESTful API
 - OpenAPI for API documentation
 - JSON / JSON-LD for API file format
 - RDF/OWL/SHACL for metadata model definition
 - SPARQL as query language
 - Basic Formal Ontology (BFO)
 - Dublin core vocabulary for generic metadata
 - RTS for terminology management and as model repository
 - ...



Examples for FAIR URIs Taken from pREDi FISH Study Registration System

Digital Object	Scheme	Example		
Study	https://id.roche.com/a2/{studyId}	https://id.roche.com/a2/32		
Study Document Protocol	https://id.roche.com/a2/{studyId}/ document	https://id.roche.com/a2/32/document		
Local Descriptor	https://id.roche.com/a2/{studyId}/ localDescriptor/{descriptorId}	https://id.roche.com/a2/32/ localDescriptor/2		



Registration System: id.roche.com

- Stable URIs by decoupling URIs from application
- Implementation of best practises and learnings from past projects
- Rolled out Q3 2021
- short URIs
- uses opaque id part as combination of two characters/ digits from Base 33
- Self registration for solution architects and software engineers
- lean registration process due to no need to governan speaking ids → no need to align between all similar systems on a

pREDi FISH Component	Namespace			
Study Registration System	https://id.roche.com/a2/			
Animal Registration System	https://id.roche.com/a3/			
Biospecimen Registration System	https://id.roche.com/a4/			
Formulation Registration System	https://id.roche.com/a5/			
Study Designer	https://id.roche.com/a6/			



RTS as registration system for terminologies

Recommendation for Metadata guidelines and conventions

- Inventory of harmonized terminologies
- Supports frictionless data integration
- Is prerequisite for a fully harmonized and interoperable data landscape
- Prospectively align application specific terminologies with existing terminologies and concepts
- Formal and machine-readable terminology representation scales better than documentation only in Github or Confluence



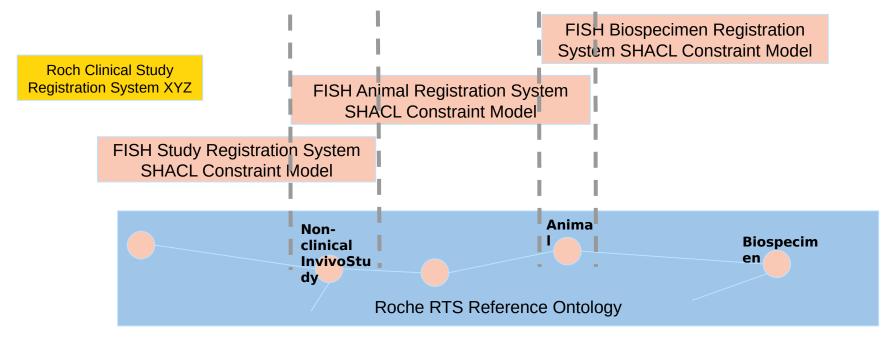
Terminology Management Application RTS Curation Client

	Application Navigator	+ Q છ	Application Concept	Entity Properties		
	 FISH Study Registration System Animal welfare authority App 1 	Termi	nology _{matio}	n	GUPRI	C Copy labels 🖊 Edit 🗘
	 Data record source system 4, , , , , , , , , , , , , , , , , , ,		Label:	Published		~ C
	ArchivedDrafted	_	Identifier: Local Status:	ROX1446033494572		*
	Not available Published		Status:	Active		
Cor	Cept	stem App Terr	Definition		Comment	
	Soci identifier for App form Soci identifier for App form Social identifier for App form Social identifier for App form Social identifier for App form	ory App Term	Data record that is i community (RTS).	eleased and available to a large	Mapping Lab	el
	E Roche portfolio project data so E Roche portfolio project stage A E Sponsoring organization App Tr E Study site App Term Q	pp Term Q	References Rel	ations Mapping Label Lan	nding Page	
	Flywheel		Concept		Terminology	Link
			Published data reco	vrd	Qualifier Descriptor	P
	▶ □ GEAR					



Model alignment between Applications

through the Reference Ontology



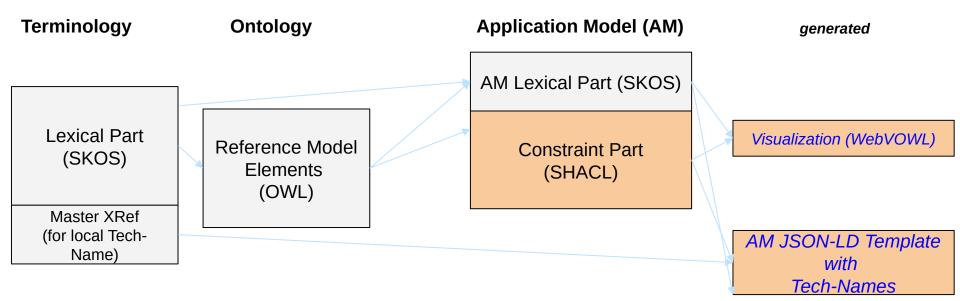


Ontology and Constraints Model in integrated RTS Tool

Roche Terminology System v3.9.0 PRODUCTION FAIR Metrics Roche Terminology Application III Model Information robert.trypuz@roche.com U Logout Variable Curation Administration Model Navigator + 9 0 Application Model Property Entity View Search Application Q Type search text C Productive Properties Under development SHACL S r: ROX38338272444006757 **Reference Model Navigator** FISH Biospecime EISH Formulation v e el is planned for Constraint Reference Model FISH Study Desig el: Class **OWL Ontology** Cell and Virus ObjectProperty e: * * Entity Models FISH SD Execu 🝷 🏶 Continuant el: is planned for 6 ▼ III FISH SD Mode # Generically dependent continuant Collected I ~ d * * Information content entity Reference URI: Formulation Collection Hereit Contract State S Inverse of: 🕨 🏶 Data item Model animal Decision Local Usage Properties Holecular entity type Directive information entity * * Nominal agent dose ObjectProperty Type: Formulation actual numeric set condition Herein Study * * Nominal agent dose Used at class: Study design * Person Mominal numeric set condition Plan of activity 8 🚍 🗙 Target class: Non clinical in vivo study Plan specification Plan of study activity sequence Algorithm Multiplicity: 1..1 🗸 Produced unmixed substance Plan of activity Produced unmixed substance type Local Technical plannedFor ~ Protocol Key: roject * Study design Study design ected biospecimen group specification Properties substance mixture description contains plan of study activity sequence Portion of substance description has comment Definition Comment Bergensent has number of first study day Bocument part 📑 is planned for indicates a planned process that is to be executed Hereit Representation is primary topic of according to the plan Harrister is reviewed by Study theme specifies study subject group # Independent continuant Study subject Specifically dependent continuant Study subject group * * Occurrent Substance mixture type

Relationships between layers

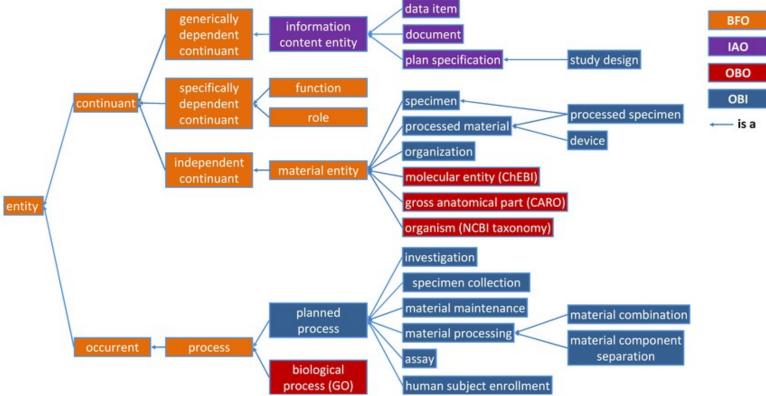






Foundational Ontologies

Ontology of Biomedical Investigations (OBI) & Basic Formal



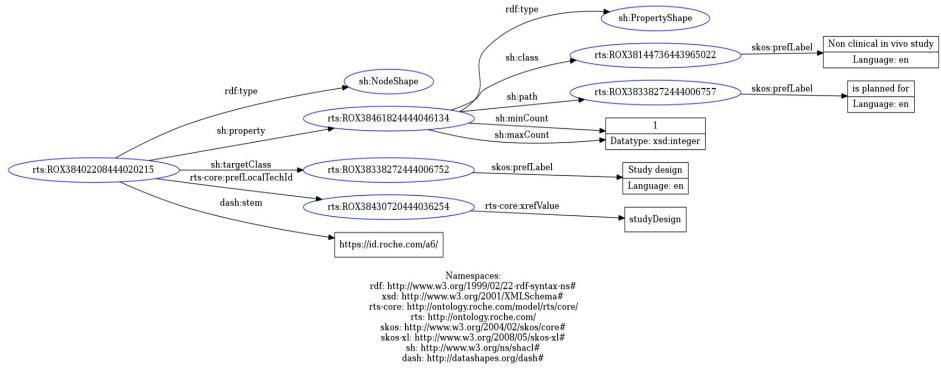


Application Specific Model in SHACL RTS Curation Client

	Model Navigator	Model ntity View	,		
	✓ □ FISH Study Registration System ✓ Ⅲ FISH SRS Model	Model Global Prope	erties	GUPRI	🖍 Edit 🛛 😂
	 * Animal welfare and use * Animal welfare authorit 	Waster Concept	ROX37664352443822935		S
	 * Document * Local descriptor 	Preferred Label Identifier:	has study scientific title		~
	* Person * Resource record	Local Technical Key:	studyScientificTitle		~
	* Roche portfolio project * * Study	Preferred Reference URI:			*
	has local descriptor	project			
lodeS	Shape	Used at class: Target class:	Study	Property	§ 🖆 🗙
	has study scientifies has study site			Constraints	🖆 ×
	ape		String		~
	FMI CRM Gatekeeper gBioPortal	Properties Definition		Comment	
	 ▶ □ GEAR ▶ □ George ▶ □ GOMAP 		ehensive summary of study design med at scientific audience. Scientific		

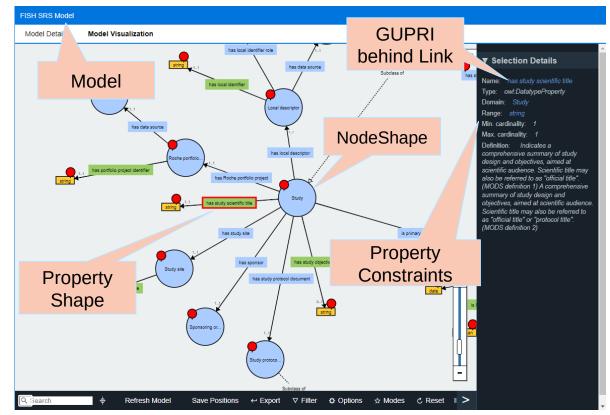


SHACL Constraint Model Example





Visualizing the SHACL Model - like an Ontology RTS Model Browser





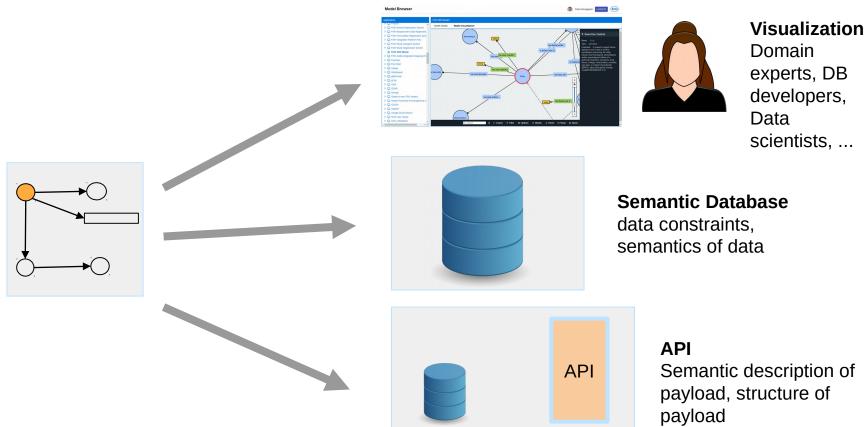
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Programmatic Access to Application Model Public REST API

Application Model RTS Application Model RTS Application Model RTS Application Models/	Todel API Export Application Model by given id.	~
Parameters		Try it out
Name	Description	
X-Auth-Token string (header)	"type": "Pro "datatype": "maxCount": "minCount": "path": { "id": " ROX "type": "C	"1", (37664352443822935",

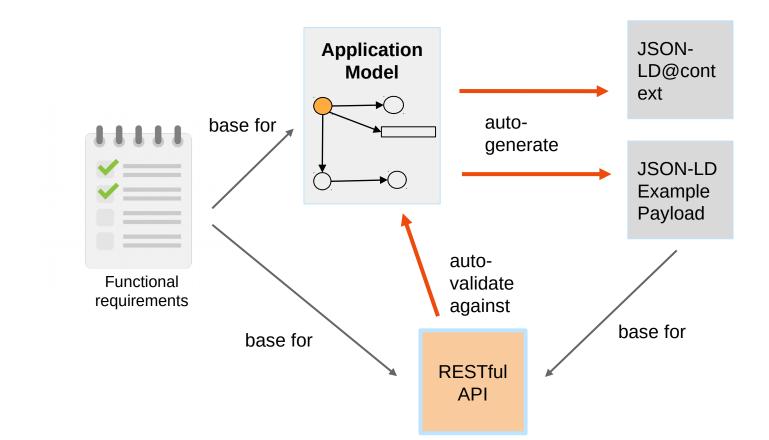
Usages for the Reference Model + Application SHACL Model







Model-supported JSON-LD REST API Design



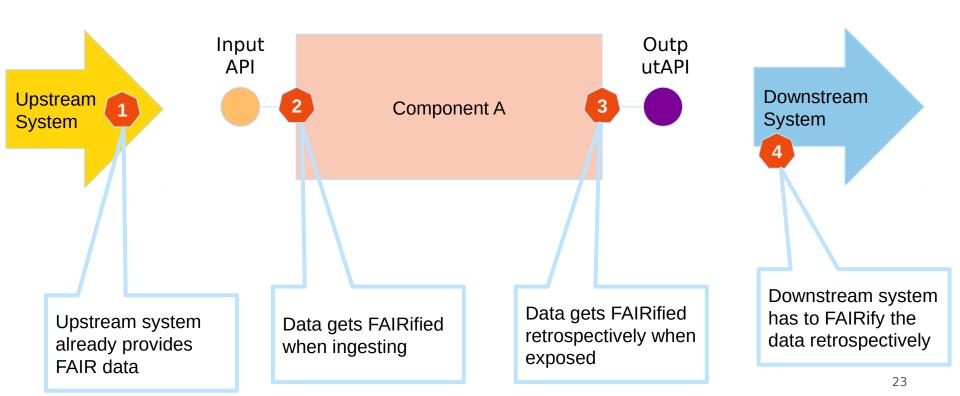


Apply a FAIR API Design with JSON-LD as payload Formation for Standards and code

- JSON (JavaScript Object Notation)
 de facto standard for Web Services based on REST
- JSON-LD (JavaScript Object Notation for Linked Data)
 - Valid JSON !
 - Keys can be still defined locally as part of API definition
 - Standardized (W3C Recommendation)
 - Adds semantic annotations
 - \rightarrow is also a valid semantic graph (RDF)
 - → maps payload to registered model
 - Reference Implementation of JSON-LD processor in Java: Apache Jena



Alternative places where to FAIRify data For the goal of having FAIR data in the downstream system





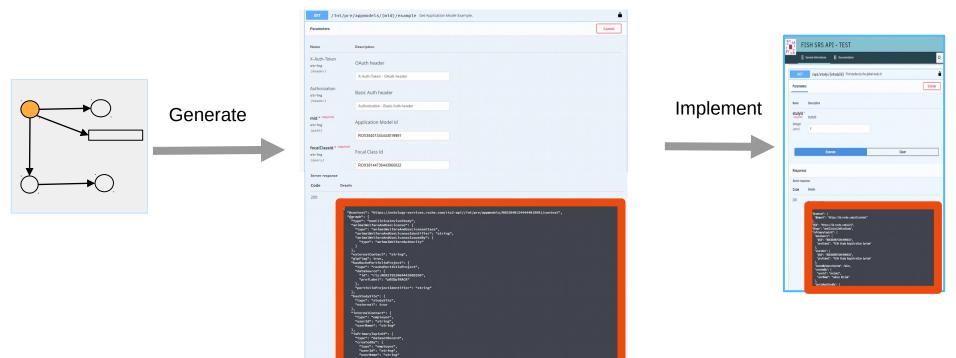
Example: Study Design Object as JSON-LD

```
{
   "@context":{
    "@import":"https://ontology-service.roche.com/rts2-api/v3/appmodels/
R0X38389248444017485/context?version=2022-06-08T09%3A05%3A30.000Z"
   },
   "@id":"https://id.roche.com/a6/1",
   "@type":"StudyDesign",
   "plannedFor":{
      "@id":"https://id.roche.com/a2/1",
      "@type":"NonClinicalInVivoStudy"
   },
   "numberOfFirstStudyDay":1,
   "comment": "some comment",
   "reviewedBy":{
      "@id":"https://id.roche.com/xyz/fishcur1",
      "@type":"Employee",
      "userName": "GLO FISH CURATOR"
   },
```



Generate API payload format from the model

RTS





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Generation of JSON-LD Example Payload

RTS Model Browser PROD 3.15.0

Applications	Merged Model				
~ Applications	Model Details Model version: 2022-08-24T16:13:01Z - Model Visualization Context Payload Example JSON-LD Validation				
✓ □ Productive	Focal Class / Shape				
	Study design +				
> 📮					
~ Ţ	Expand to prefLabels				
Cell and Virus exposure					
FISH SD Execution Model FISH SD FIS	TS as Base				
III Merged Model	Avoid special characters				
proposed exposure activity					
SD Substance-Molecule model	2				
Substance exposure activity Substance reference model					
> 🖵	GENERATE				
> _	Payload Example 🛅 COPY TO CLIPBOARD 👤 DOWNLOAD				
> _	{ "@context": {				
$\rightarrow \Box$	<pre>@context - \ "@import". "https://ontology-services.roche.com/rts2-apl/v3/appmodels/ROX38521440444104515/context?version=2022-08-24T16:13:012"</pre>				
> 🖵	3.				
	"@graph": { "@id": "https://id.roche.com/a6/studyDesign/1",				
	"@type": "studyDesign",				
	"comment": "string", "containsCollectedBiospecimenGroupSpecification": [
	(
	"@id": "https://id.roche.com/a6/collectedBiospecimenGroupSpecification/1", "@type": "collectedBiospecimenGroupSpecification",				
	"biospecimenType": (
	"@id": "rts:ROX1308059566380", "@type": "collectedBiospecimenTypeATEnum",				
	"gripe - to reacted by contrast y performing - "prefit abel", "Thymus"				
	}, "denotes": {				
× · · · · · · · · · · · · · · · · · · ·	"@di": "https://id.roche.com/a6/collectedBiospecimenGroup/1",				



Example: Study Design Object as Generated from SHACL

```
"@context":"https://ontology-services.roche.com/rts2-api/v3/appmodels/
ROX38389248444017485/context?version=2022-04-07T14%3A46%3A15.000Z",
   "@graph":{
      "@id":"https://id.roche.com/a6/1",
      "@type":"StudyDesign",
      "comment":"string",
       "plannedFor":{
         "@type":"NonClinicalInvivoStudy",
         "hasParticipant":[
               "@type":"Subject",
               "@id":"https://id.roche.com/a6/1/subjects/1"
            },
            . . .
```



Example: JSON-LD Context for Study Design Object

```
Look up "title" in RTS for detailed semantics
                                      http://ontology.roche.com/ROX37664352443
                                      822935
  "@context": {
                                      → "Study scientific title" : "Indicates a
    "id": "@id",
                                      comprehensive summary [...]"
    "type": "@type",
    "rts": "http://ontology.roche.com/",
    "prefLabel":
"http://www.w3.org/2004/02/skos/core#pref
    "StudyDesign":"rts:R0X38338272444006
      "plannedFor":{
          "@id":"rts:R0X3833827244400675
          "@type":"@id"
      },
    "title": "rts:R0X37664352443822935",
    "sponsoringOrganization": "rts:ROX38092032443955602"}
...
```



Validation of JSON-LD Payload

RTS Model Browser PROD 3.15.0

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Applications	Merged Model					
 Applications 	Model Details	Model version: 2022-08-24T16:13:01Z 👻	Model Visualization	Context	Payload Example	JSON-LD Validation
✓ □ Productive						
	Select JSON-LD	file to upload				
> 🛱	Datei auswähler	n Keine ausgewählt				
	Paste JSON-LD	content to validate				
✓ ↓ FISH Study Designer System						
Cell and Virus exposure						
FISH SD Execution Model						
FISH SD Model						
Merged Model						
proposed exposure activity						
SD Substance-Molecule model						
Substance exposure activity Substance reference model						
> 🖵						
	JSON-LD Valida	tion Result				(REPORT TO CLIPBOARD 👱 DOWNLOAD REPORT
> _	2. The JSON-L	D payload contains Classes not defined in the	ApplicationModel: [collecte	edBiospecimer	TypeATEnum, recordSt	tatusATEnum, studySubjectGroupCategoryATEnum]
> 🖵						
> 📮		dStatusATEnum" contains properties that are no				
> _		ctedBiospecimenTypeATEnum" contains proper				
		3. Class "collectedBiospecimenTypeATEnum" contains properties that are not defined in Application Model Class: [prefLabe]				
		 Class "studySubjectGroupCategoryATEnum" contains properties that are not defined in Application Model Class: [prefLabel] Class "studySubjectGroupCategoryATEnum" contains properties that are not defined in Application Model Class: [prefLabel] 				
$\mathbf{\nabla}$	5. Class sludy	SubjectoroupcategoryArEnum contains prope	crues that are not defined	III Application	would class. [preitabe	l)
	4. The range o	f the "biospecimenType" property is below the n	ninimal count of the said r	ange defined t	y the application model	
		f the "membersDerivedFromGroup" property is				
		f the "localIdentifier" property is below the minin				
\mathbf{x}		f the "memberOf" property is below the minimal				
\mathbf{Q}	-	f the "assignedFrom" property is below the mini				
$\overline{\mathbf{Q}}$		f the "participatesIn" property is below the minin f the "name" property is below the minimal cour	-			
\square	4. The fallye o	i the name property is below the minimal cour	it of the salu fallge deline	u by the applic		
	-					

F.A.I.R Metrics Assessment



Findability

A Accessibili

- ★ F1. Metadata are assigned globally unique and persistent identifiers:
- ★ F2. Data are described with rich metadata (structured and grounded)
- ★ F3. Metadata clearly and explicitly include the identifier of the data they describe
- ★ F4. Metadata are registered or indexed in a searchable resource

- ★ A1. Metadata are retrievable by their identifier using a standardised communication protocol
 - A1.1. The protocol is open, free and universally implementable
 - A1.2. The protocol allows for an authentication and authorisation where necessary
- ★ A2. Metadata is accessible even when the data is no longer available

Required for FAIR Data APIs



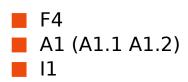
Interoperability

- ★ I1. Metadata use a formal, accessible, shared, and broadly applicable language for knowledge representation
- ★ I2. Metadata use vocabularies that follow the FAIR principles
- ★ I3. Metadata include qualified references to other metadata

Reusability

- ★ R1. Metadata are richly described with a plurality of accurate and relevant attributes
- ★ R1.2. Metadata are associated with detailed provenance
- ★ R1.3. Metadata meet domain-relevant community standards

Supported by FAIR Data APIs





Acknowledgements



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Doing now what patients need next



Subtitle goes here but is not mandatory