



BrAPI2RDF - Converting between BrAPI and RDF

Cyril Pommier - Pierre Larmande

Use case Wheat Phenotyping dataset

Phenotypes

Winter wheat (*Triticum aestivum L.*) phenotypic data from the multi-year trials of the INRA Small Grain Cereals Network.

François-Xavier Oury, Emmanuel Heumez, Bernard Rolland, Jérôme Auzanneau, Pierre Xavier Charrier, Hubert Chiron, Camille Depatureaux, Laurent Falchetto, Olivier Gardet, Christophe Lecomte, Jean-Yves Morlais, Pierre Pluchard, Didier Tropée, Maxime Trotter, Michel Rousset, Gilles Charmet

[Query dataset as a semantic graph.](#)

[Or download the dataset as RDF archive.](#)

Abstract

Published 2018 by Portail Data Inra

[Back to Form](#)

Search parameter(s):

[Geolocation](#)



Trial list **Phenotypic data**

Data table view : Default

LEVEL: REPLICATION

1-10 of 81,434 replication | Display 10 results per page

Lot Number	GENOTYPE ID		Treatment	Trial Name	Trial Site
	Accession Number	Accession Name			
CF99016	29114	CF99016	treated	BTH_Dijon_2000_SetA	Dijon
RE99002	29191	RE99002	treated	BTH_Dijon_2000_SetA	Dijon
SOISSONS	6607	SOISSONS	low inputs	BTH_Dijon_2000_SetA	Dijon
RE9819	37778	RE9819	untreated, no fungicide	BTH_Le_Moulon_2000_SetA	Le Moulon
CF99016	29114	CF99016	treated	BTH_Le_Moulon_2000_SetA	Le Moulon
CF99031	22479	CF99031	low inputs	BTH_Le_Moulon_2000_SetA	Le Moulon

VARIABLE	VARIABLE	VARIABLE
rdt: Grain yield at 0% humidity	r: Yield	prec: Earliness
73,8	105,6	
66	120,5	
110,6	142,9	
106,5	102,4	
41,6	104,8	
39,9	75,5	141
64,6	115,4	139
66,0	88,0	142

Why RDF for BrAPI?

- Easier integration across sources
 - Germplasm linking at different levels (varieties, accession, hybrid)
 - Linking to literature data through traits
 - Linking/integrating and subsetting accross different sources/dataset
 - ...
- More complex “chained” queries are easier to perform on graphs
- Validation (model-, type- and value-level)
- One workflow/tool for all BrAPI compliant databases.
- → Let's lift BrAPI JSON and add JSON-LD context

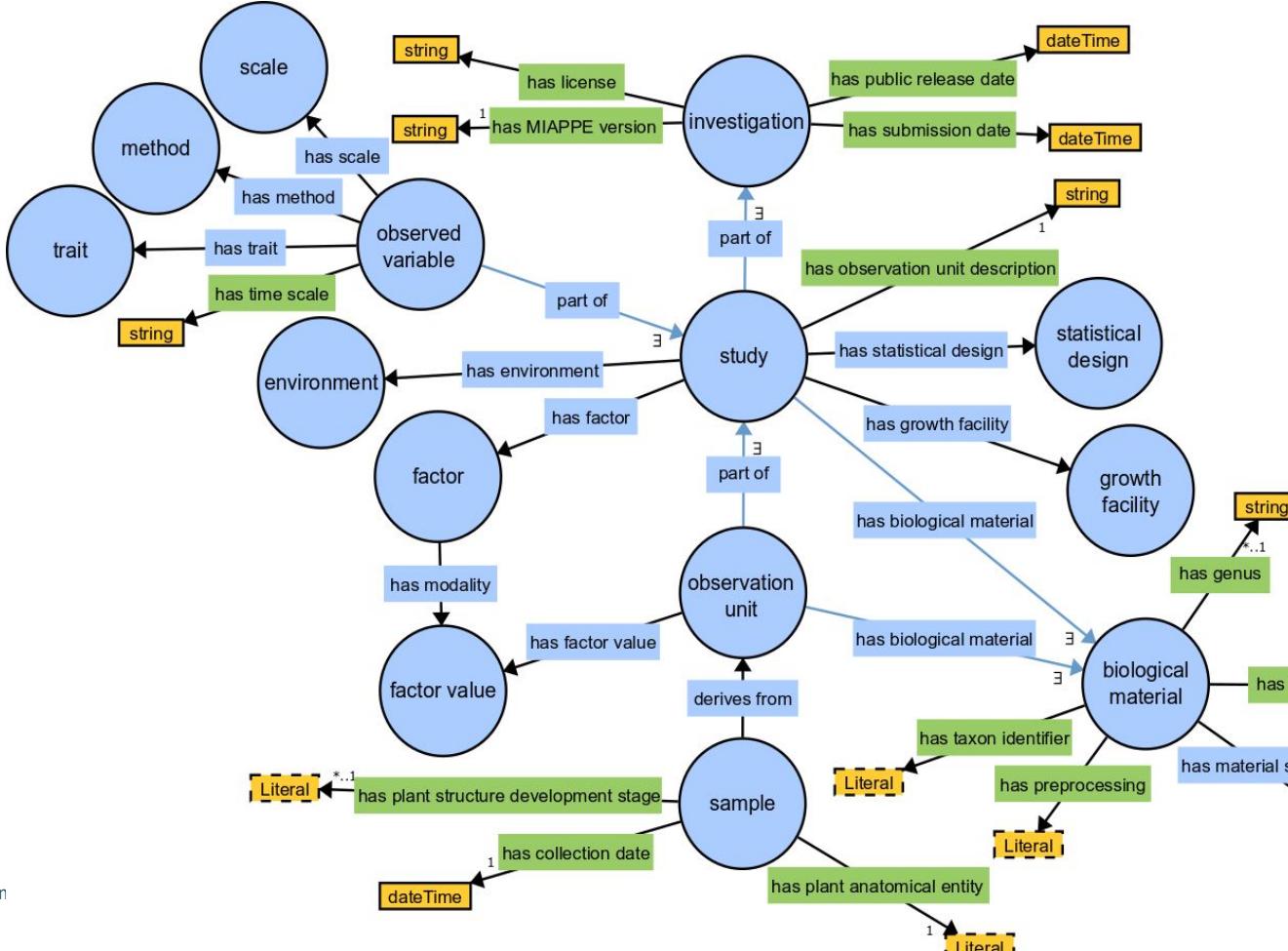
Ontologies

BrAPI types

- "<http://brapi.org/rdf/Study>"

MIAPPE Ontology

- Plant Phenotyping Experiment Ontology



JSON-LD basics

A JSON compatible format to map JSON data to RDF triplets

```
{ "@context":  
  "http://brapi.org/rdf/StudyGermplasm.jsonld",  
  
  "studyDbId": "123",  
  
  "trialName": "myBestTrial",  
  "germplasm": [ {  
      "germplasmPUI": "http://www.crop-dive  
rsity.org/mgis/accession/01BEL084609",  
      "germplasmDbId": "382",  
      "germplasmName": "Pahang",  
      "accessionNumber": "ITC0609",  
      ...  
    } ]  
}
```

INRAE

```
{  
  "@context":  
    "http://brapi.org/rdf/StudyGermplasm.jsonld",  
    "studyPUI": "http://.../studyABC",  
    "studyDbId": "123",  
    "trialPUI": "http://.../trials#trialX",  
    "trialName": "myBestTrial",  
    "data": [ {  
        "germplasmPUI": "http://www.crop-diversity.or  
g/mgis/accession/01BEL084609",  
        "germplasmDbId": "382",  
        "germplasmName": "Pahang",  
        "accessionNumber": "ITC0609",  
        ...  
      } ]  
}
```

JSON-LD basics - context

Mapping and result for Study

```
{  
  "@context":  
  "http://brapi.org/rdf/StudyGermplasm.jsonld",  
  "studyPUI":"http://.../studyABC",  
  "studyDbId": "123",  
  "trialPUI":"http://.../trials#trialX",  
  "trialName": "myBestTrial",  
  "germplasms": [ {  
      "germplasmPUI":"http://www.crop-dive  
rsity.org/mgis/acquisition/01BEL084609",  
      "germplasmDbId": "382",  
      "germplasmName": "Pahang",  
      "acquisitionNumber": "ITC0609",  
      ...  
    } ]  
}
```

```
{  
  "@context": {  
    "xsd": "http://www.w3.org/2001/XMLSchema#",  
    "brapi": "https://brapi.org/rdf/",  
  
    "Study": "brapi:Study",  
    "Trial": "brapi:Trial",  
    "Location": "brapi:Location",  
    "studyPUI": "@id",  
    "studyDbId": "brapi:hasDatabaseIdentifier",  
    "studyName": "brapi:hasName",  
    "trialPUI": {  
      "@id": "brapi:hasTrial",  
      "@type": "@id"  
    },  
    "germplasmPUI": {  
      "@id": "brapi:hasGermplasm",  
      "@type": "@id"  
    }  
  }  
}
```

BrAPI JSON-LD: ETL approach

Example: github.com/gnpis/BrAPI-extract-index-prototype

- 1) Extract BrAPI data
- 2) Transform
 - a) URI generation where missing (this creates objects link by URI)
 - b) Add JSON-LD context (mapping ontology properties)
 - c) Convert to RDF format
- 3) Load into a triplestore

URI generation

template: {brapi server}/brapi/v1/{brapi object}/{database id}

example URI: <https://urgi.versailles.inra.fr/GnplISCore-srv/brapi/v1/study/E012E0>

Data lifting

BrAPI calls for individual resources

OK With BrAPI V2

- Trial
- Study [/studies/{studyDbId}](#)
- Germplasm
- ObservationVariable
- ObservationUnits

Data Lifting

Links between resources

Studies → Germplasm & others

Not natively present in BrAPI JSON (anymore ?)

To be added to exchange a whole trial/dataset ?

→ File Archive (RO Crate) Use case

SPARQL Micro-Services

Querying Web APIs with SPARQL

Courtesy of Franck MICHEL



INRAE

BrAPI to Linked Data, Pierre Larmande, Cyril Pommier
27/04/2022

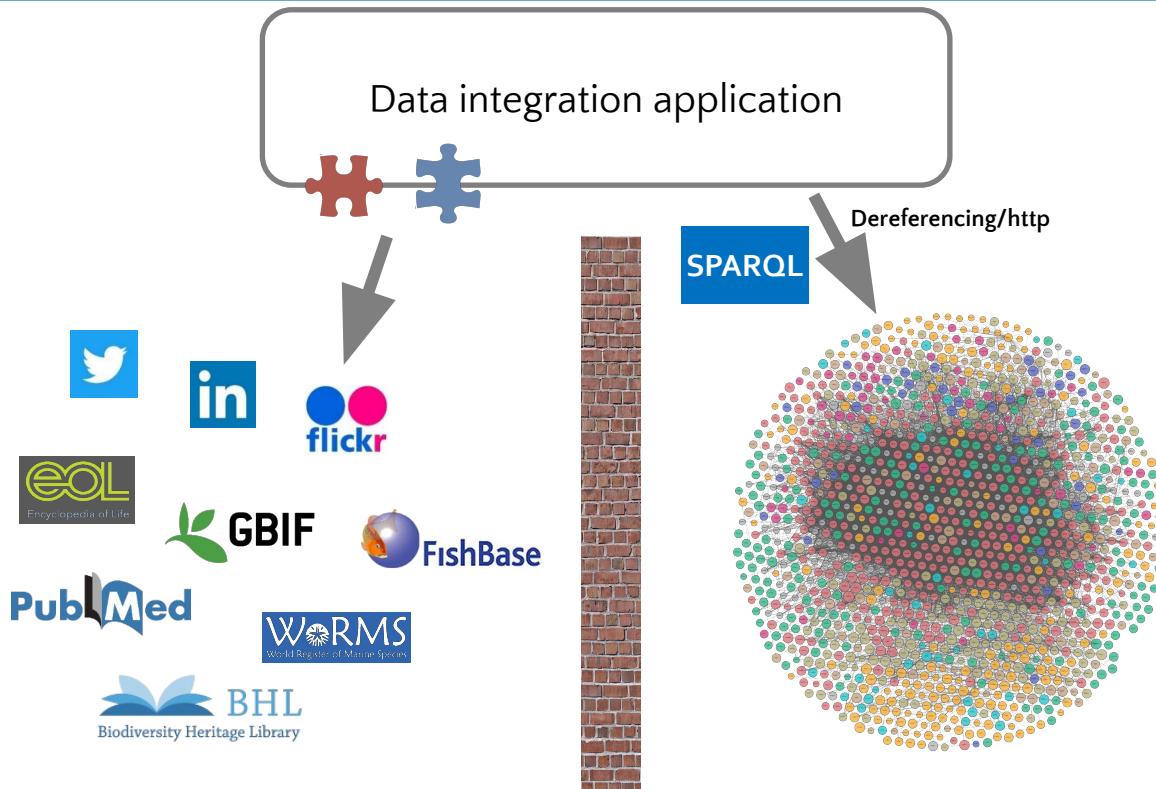
UNIVERSITÉ
CÔTE D'AZUR



inria

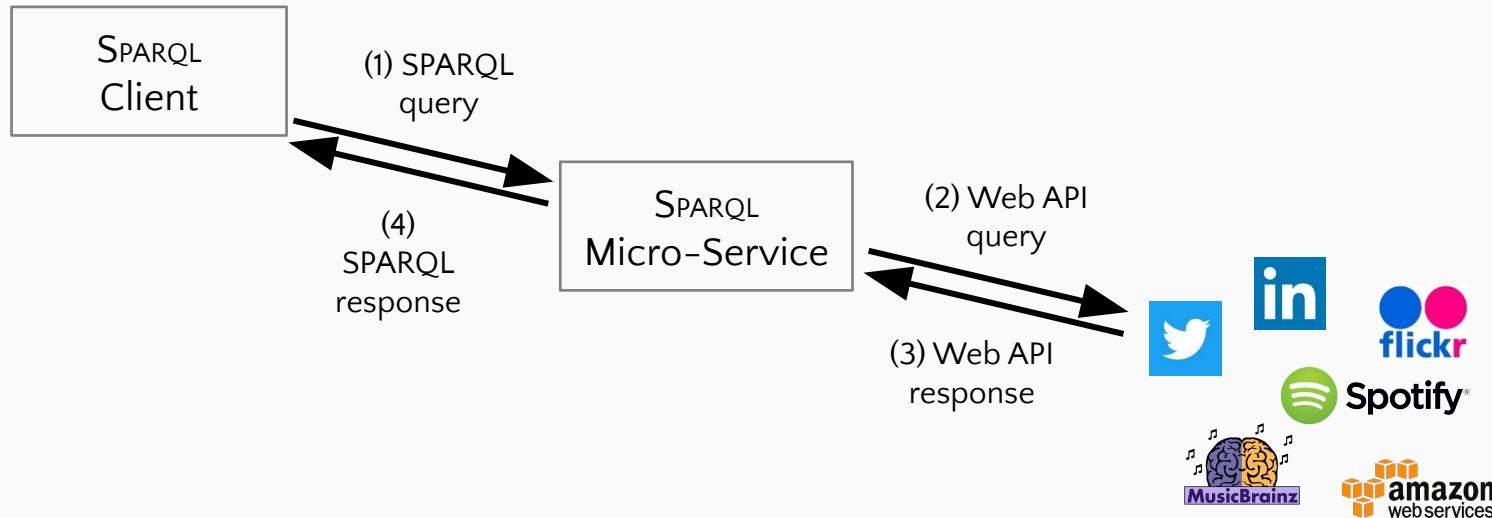
cnrs

How to leverage Web APIs and Linked Data at the same time?



The SPARQL Micro-Service Architecture

Lightweight method to **query a Web API with SPARQL**
(and assign dereferenceable URIs to Web API resources)



A SPARQL μ-service is a configurable SPARQL endpoint whose arguments delineate the graph being queried.

Endpoint: <http://example.org/flickr/getPhotosByTag?tag=bridge>

```
SELECT * WHERE {  
    ?photo a schema:Photograph;  
           schema:name ?title;  
           schema:contentUrl ?img.  
}
```

Arguments passed
as HTTP parameters

Endpoint: http://example.org/flickr/getPhotosByTag_sd

```
SELECT * WHERE {  
    ?photo a schema:Photograph;  
           schema:keywords "bridge";  
           schema:name ?title;  
           schema:contentUrl ?img.  
}
```

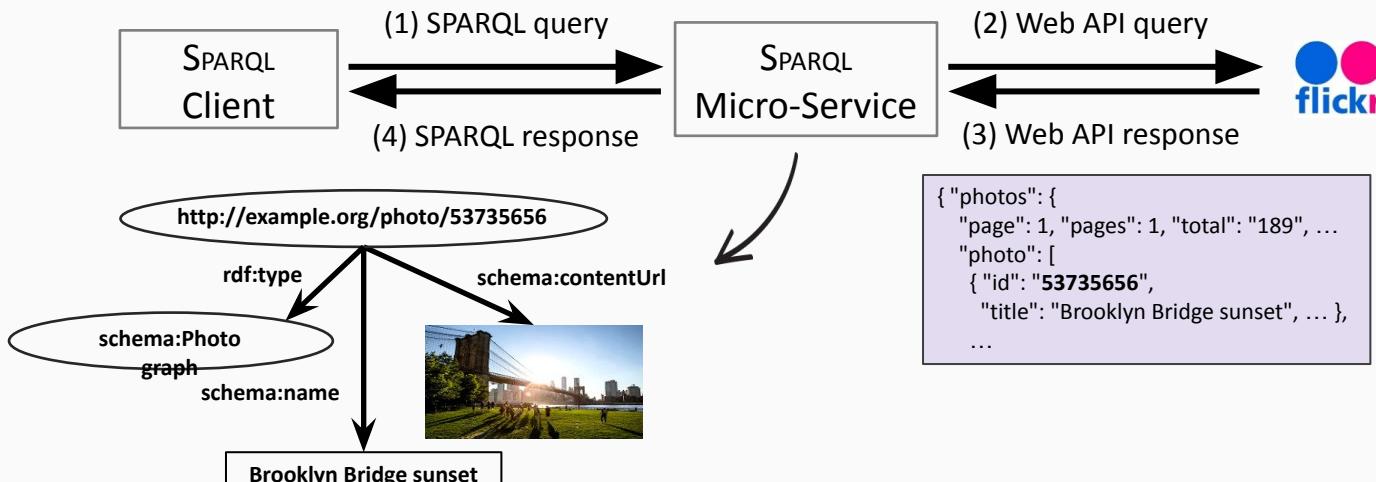
Arguments passed
in the graph pattern

Into the details - Arguments passed as HTTP parameters

```
SELECT * WHERE {  
    ?photo a schema:Photograph;  
    schema:name      ?title;  
    schema:contentUrl ?img.  
}
```

<http://example.org/flickr/getPhotosByTag?tag=bridge>

[https://api.flickr.com/services/rest/?method=flickr.photos.search&](https://api.flickr.com/services/rest/?method=flickr.photos.search&format=json&per_page=100&tags=bridge&)

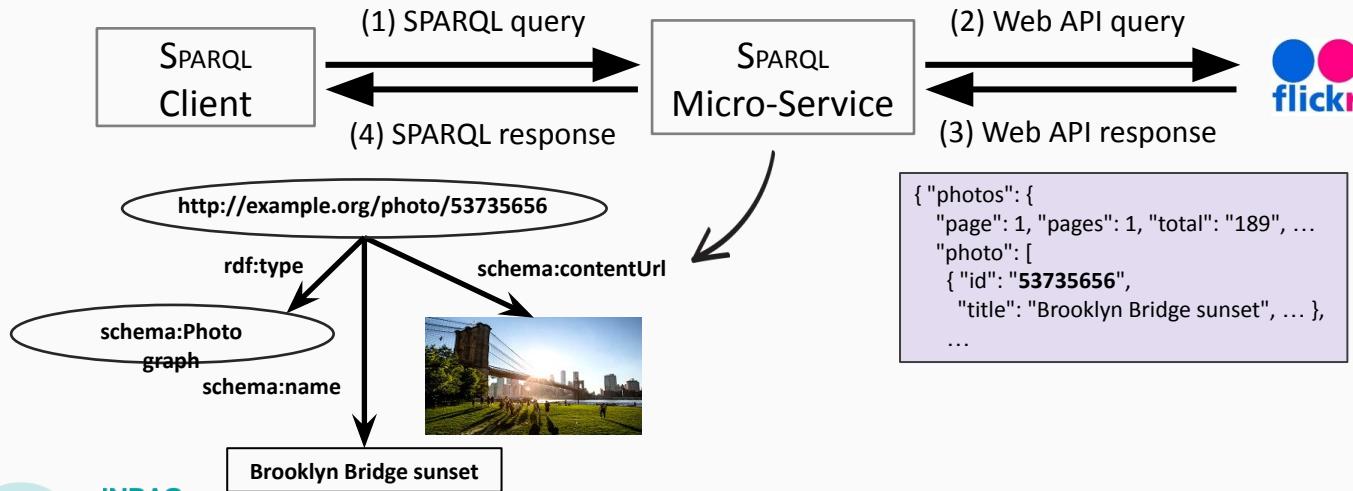


Into the details - Arguments passed in the graph pattern

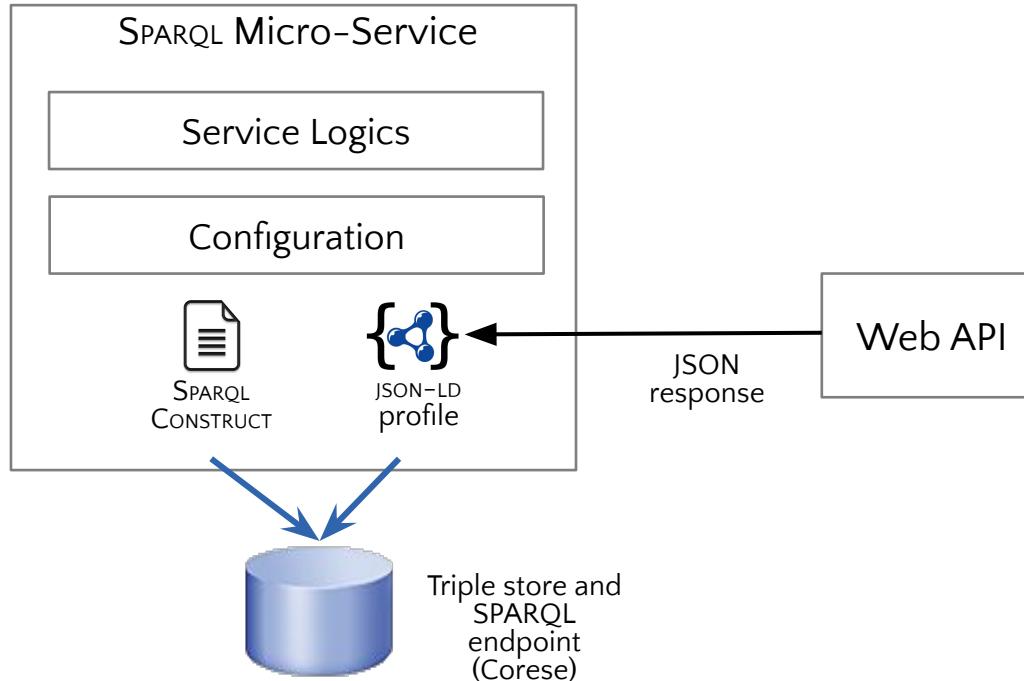
```
SELECT * WHERE {  
    ?photo a schema:Photograph;  
    schema:keywords "bridge";  
    schema:name ?title;  
    schema:contentUrl ?img.  
}
```

http://example.org/flickr/getPhotosByTag_sd

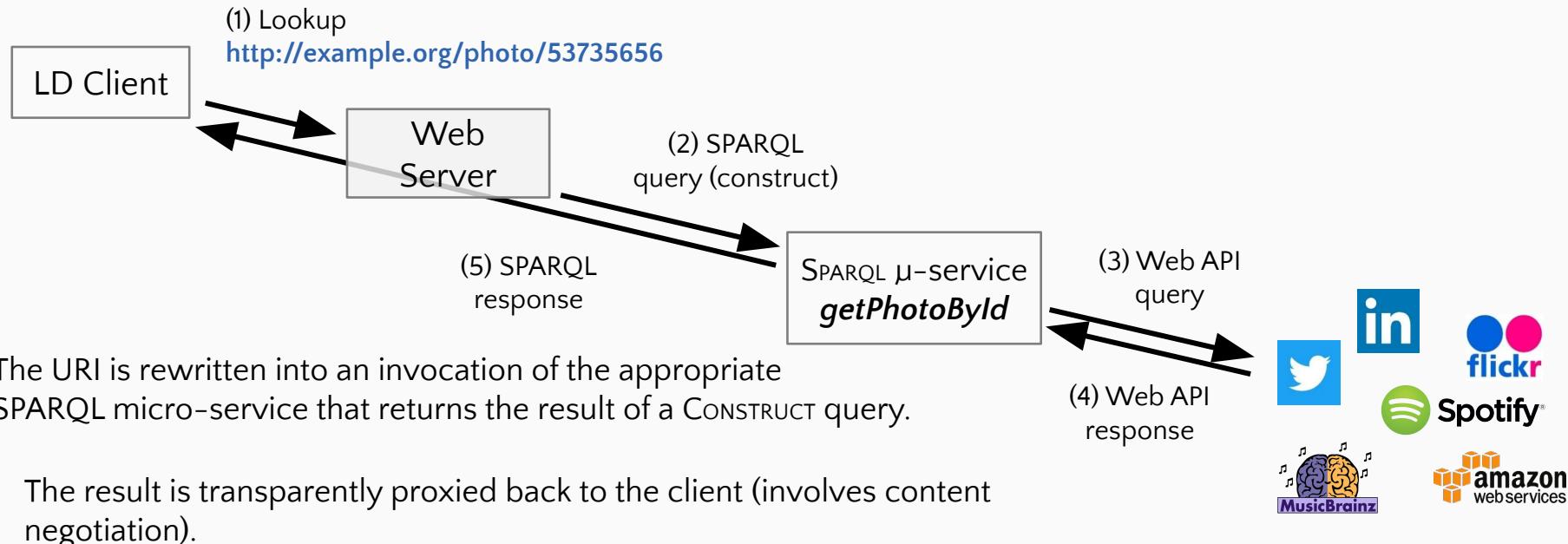
https://api.flickr.com/services/rest/?method=flickr.photos.search&format=json&per_page=100&tags=bridge&...



Translating the Web API response to RDF

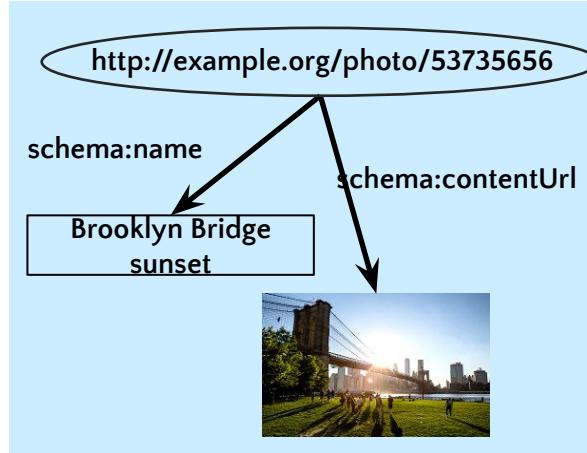


Bridging Web APIs and the Web of Data

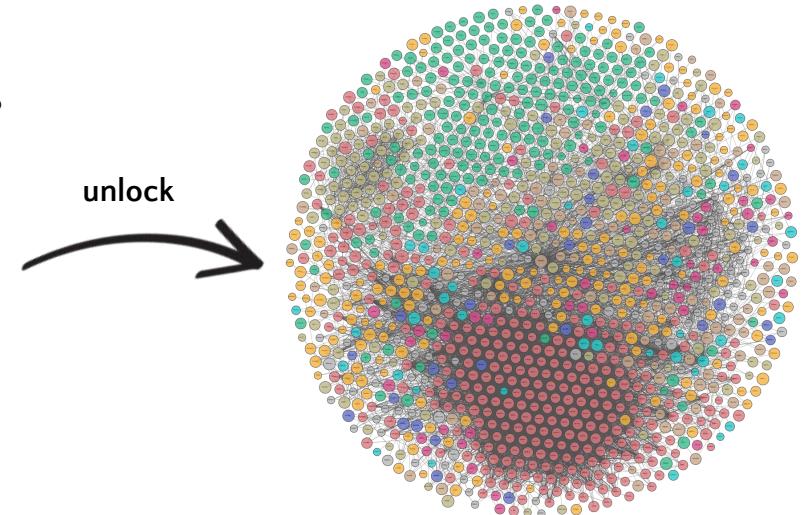


Bridging Web APIs and the Web of Data

SPARQL µ-service
getPhotosByTag

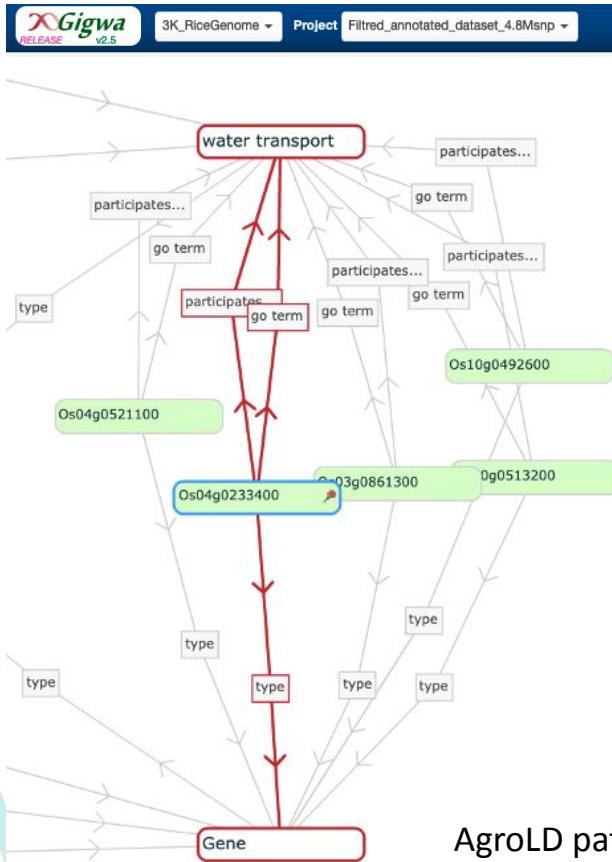


Assign dereferenceable URIs to
Web API resources



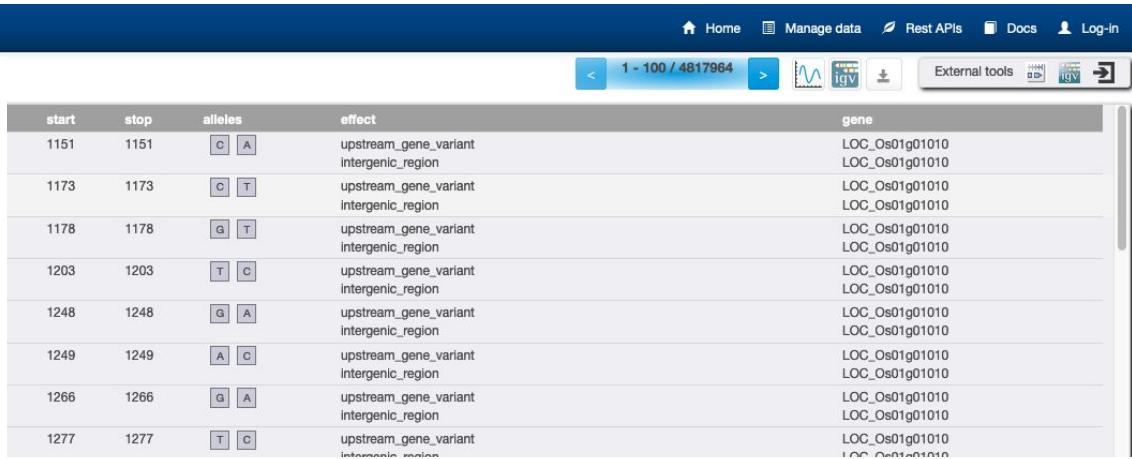
Expose into the Web of Data
resources previously locked in a
silo

Use case with Gigwa and AgroLD



The AgroLD pathway viewer displays a network of genes and their relationships. A central node, "water transport", is connected to several other nodes: "participates..." (multiple edges), "go term" (multiple edges), and "type" (multiple edges). One "type" edge points to a node labeled "Gene". Another "type" edge points to a node labeled "Os04g0233400", which is highlighted with a blue border. This node is also connected to "type" nodes for "03g0861300" and "0g0513200". A green box highlights the "Os04g0521100" node.

AgroLD pathway viewer



The Gigwa interface shows a table of variants:

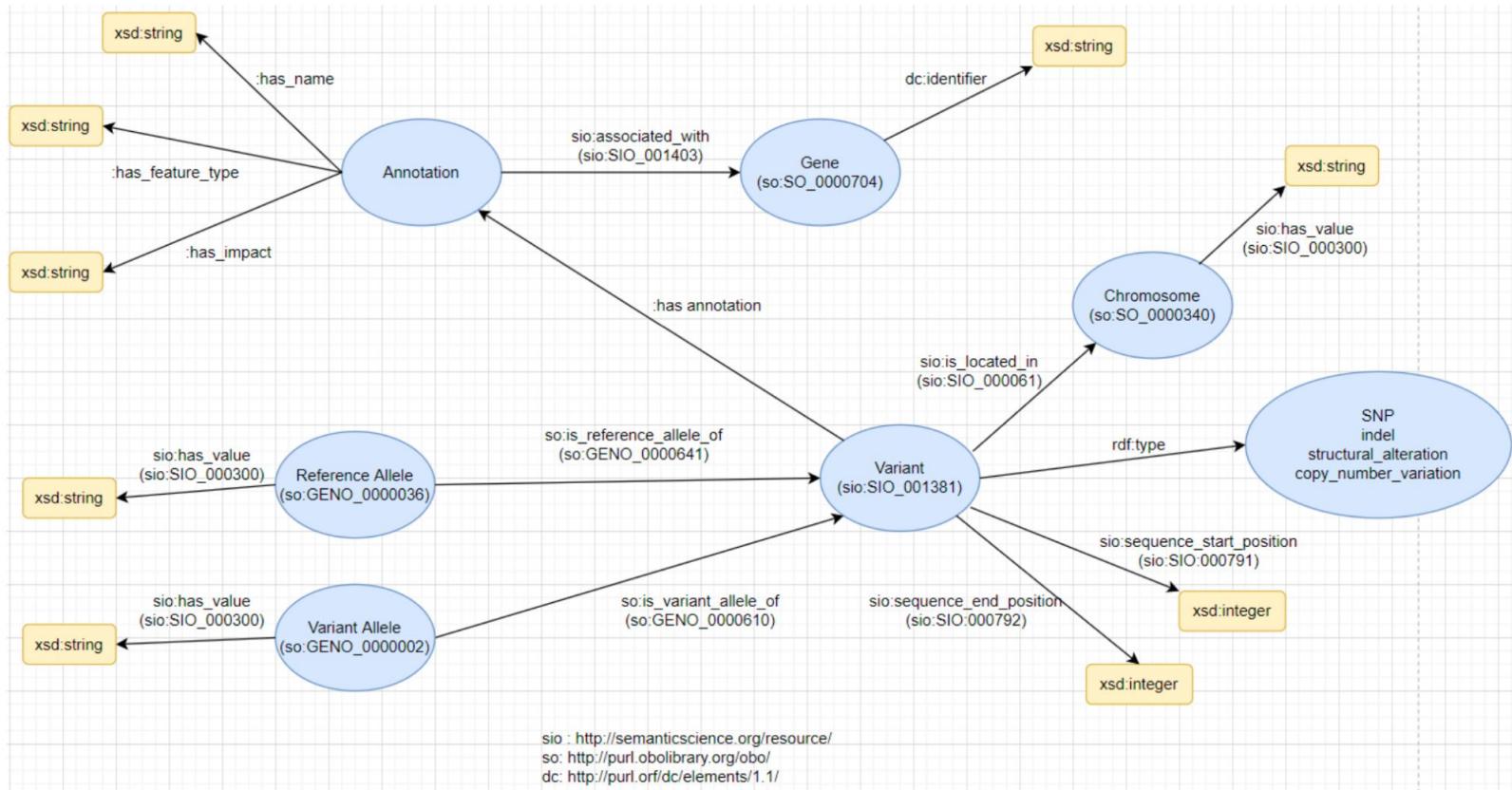
start	stop	alleles	effect	gene
1151	1151	C A	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010
1173	1173	C T	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010
1178	1178	G T	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010
1203	1203	T C	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010
1248	1248	G A	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010
1249	1249	A C	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010
1266	1266	G A	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010
1277	1277	T C	upstream_gene_variant intergenic_region	LOC_Os01g01010 LOC_Os01g01010

1 - 100 / 4817964 < > Home Manage data Rest APIs Docs Log-in External tools IGV IGV

Which genes involved in “water transport” pathway contain some variations with effects in translation or traduction

p. 19

RDF(s) model for Gigwa



Example retrieving variant from Gigwa

```
https://gigwa.ird.fr/gigwa/rest;brapi/v2/variants?end=1152&referenceDbId=3K_RiceGenome§1&start=1000
```

start position : 1000

end position : 1152

referenceDbId: 3K_RiceGenome§1

2 micro-services were created

getVariants : get all variants between start and end pos on a chr

getVariantsWithAnnotations : get annotations from a variant

```
▼ data:  
  ▼ 0:  
    ▼ additionalInfo:  
      ▼ transcriptEffects:  
        ▼ 0:  
          Allele: "A"  
          Annotation: "upstream_gene_variant"  
          Annotation_Impact: "MODIFIER"  
          Gene_Name: "LOC_Os01g01010"  
          Gene_ID: "LOC_Os01g01010"  
          Feature_Type: "transcript"  
          Feature_ID: "LOC_Os01g01010.1"  
          Transcript_BioType: "Coding"  
          HGVS.c: "c.-462C>A"  
          Distance: "1752"  
        ▼ 1:  
          Allele: "A"  
          Annotation: "intergenic_region"  
          Annotation_Impact: "MODIFIER"  
          Gene_Name: "LOC_Os01g01010"  
          Gene_ID: "LOC_Os01g01010"  
          Feature_Type: "intergenic_region"  
          Feature_ID: "LOC_Os01g01010"  
    ▼ alternate_bases:  
      0: "A"  
      end: 1151  
      referenceBases: "C"  
      referenceName: "1"  
      start: 1151  
      variantDbId: "3K_RiceGenome§1151"  
      variantType: "SNP"
```

Thanks for your attention