

Swiss Institute of
Bioinformatics

Federation: Beyond the home organization.

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Principal Software Engineer – Swiss-Prot group

(Con)Federation

FMI
Friedrich Miescher Institute
for Biomedical Research



Swiss TPH

ETH
Eidgenössische Technische Hochschule Zürich
Swiss Federal Institute of Technology Zurich



Agroscope

Zürich University
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u^b
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BERN

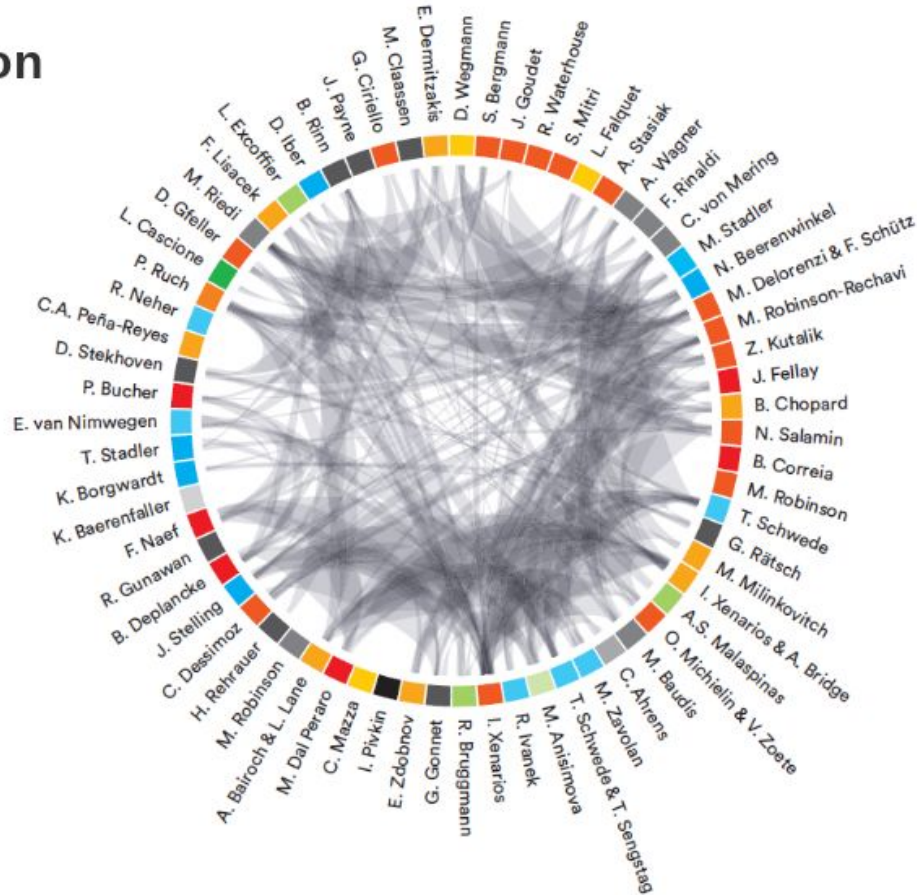
U^s
Università
della
Svizzera
italiana

IOR
Institute of Oncology Research

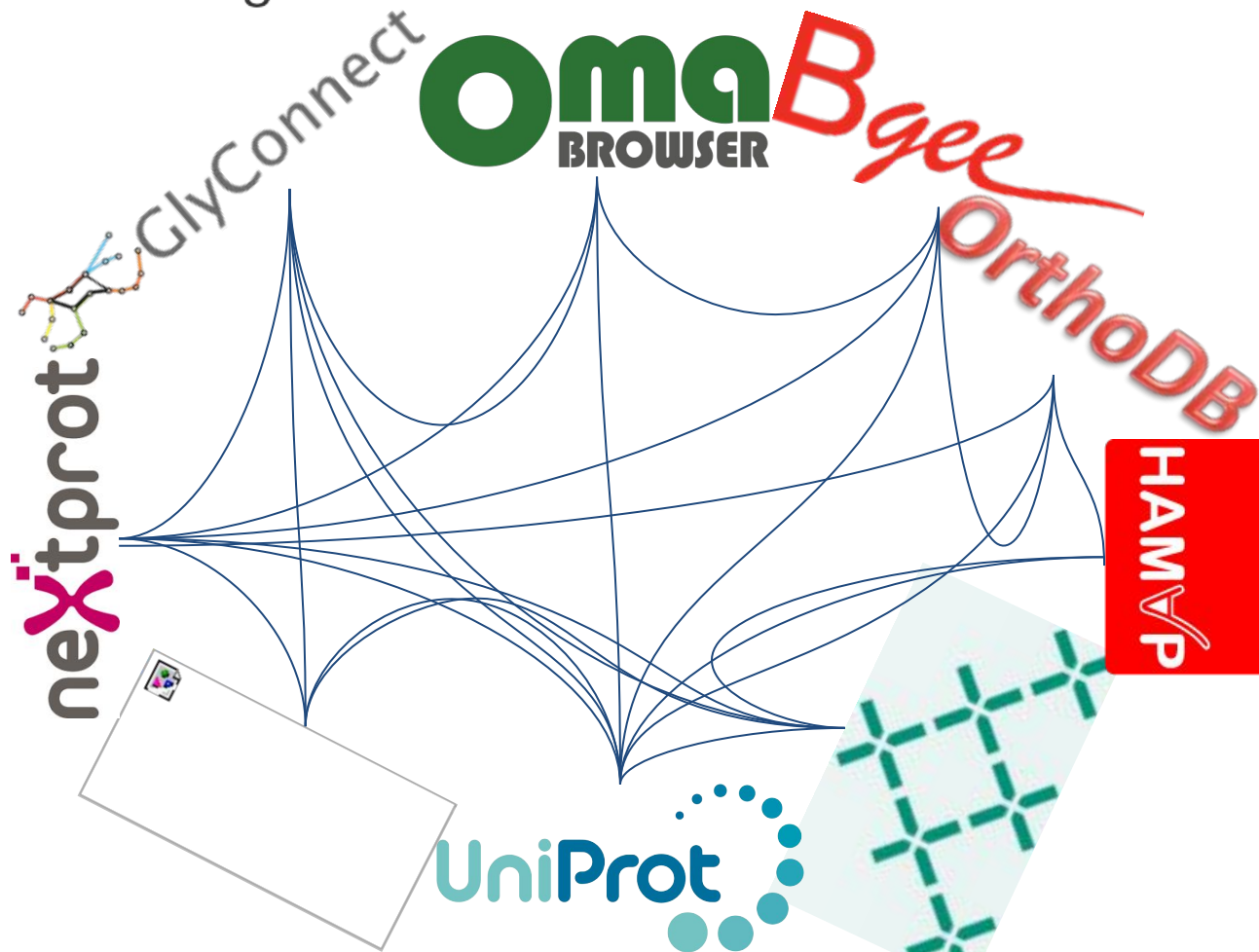
20 institutional
partners
all over
Switzerland

SIB collaboration network

254 publications
in 2018



Fostering scientific collaboration



Federation inside the home

The screenshot shows the Expasy website interface. At the top, there is a navigation bar with links for Home, About, SIB News, and Contact. Below this is a search bar with the text "SPARQL" and a red search button. Under the search bar, there is a hint: "e.g. BLAST, UniProt, MSH6, Albumin...". Below the search bar, there is a message: "You can also query 'SPARQL' into a selection of SIB databases in parallel" with a red plus icon. The main content area displays "10 resources found for SPARQL". There are ten cards, each representing a different SPARQL endpoint. Each card has a red icon, a title, and a brief description. The cards are arranged in two columns. At the bottom of the page, there is a footer with the Expasy logo, the text "Expasy is operated by the SIB Swiss Institute of Bioinformatics | Terms of Use", and a link "Back to the top".

Expasy

Home About SIB News Contact

SPARQL

e.g. BLAST, UniProt, MSH6, Albumin...

You can also query "SPARQL" into a selection of SIB databases in parallel

10 resources found for SPARQL

- OMA SPARQL endpoint**
Orthology inference among complete genomes
- Bgee SPARQL endpoint**
Gene expression expertise
- OrthoDB SPARQL endpoint**
Evolutionary and functional annotations of orthologs
- HAMAP SPARQL endpoint**
Use HAMAP + SPARQL to generate portable annotation pipelines
- UniProt SPARQL endpoint**
Query UniProt + related data in a powerful SQL-like language
- neXtProt SPARQL endpoint**
SPARQL endpoint for the neXtProt Human protein knowledgebase
- Rhea SPARQL endpoint**
SPARQL access to the Rhea DB
- SwissLipids SPARQL endpoint**
SPARQL access to the SwissLipids knowledgebase
- MetaNetX SPARQL endpoint**
SPARQL access to MetaNetX
- GlyConnect SPARQL endpoint**
SPARQL access to GlyConnect

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Publicly available SPARQL endpoints

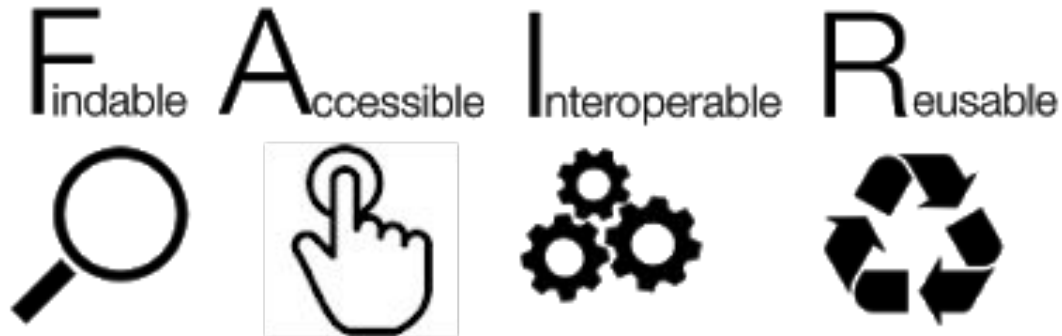
More internal scale

Thousands to multi billions

Lots of different implementations

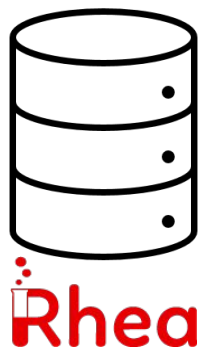
- RDF4j native
- Virtuoso 7.2 Open Source
- GraphDB
- Blazegraph (to be replaced)

The FAIRRest format of them all



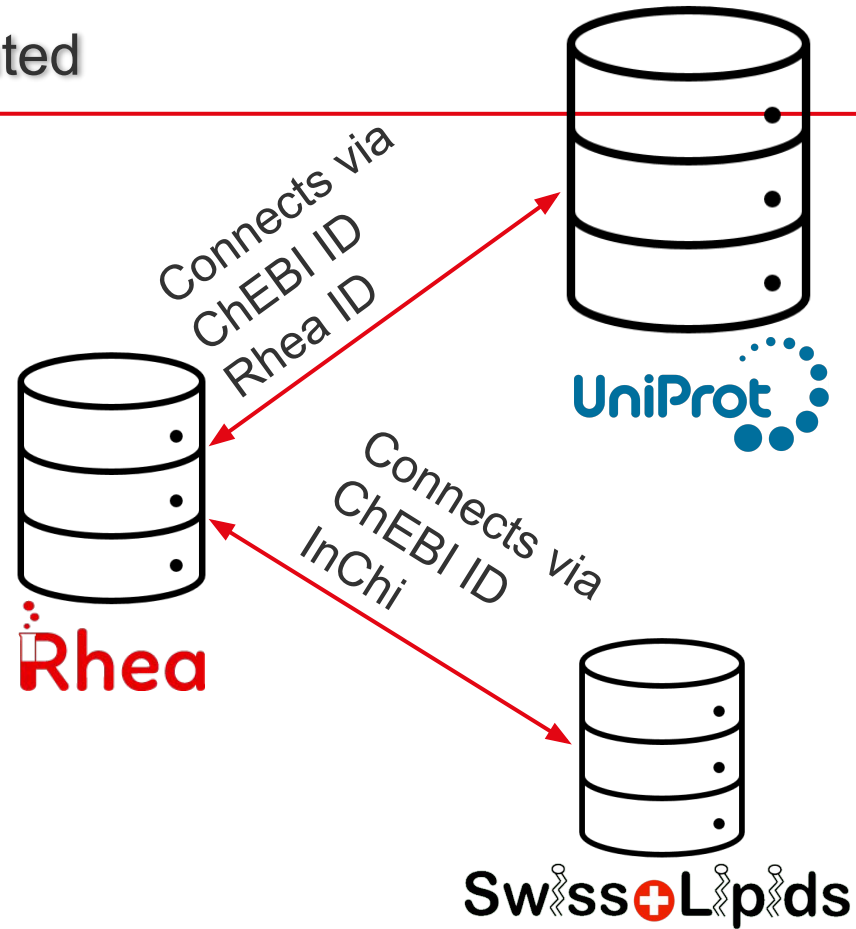
FAIR via SPARQL

Chemical reaction database

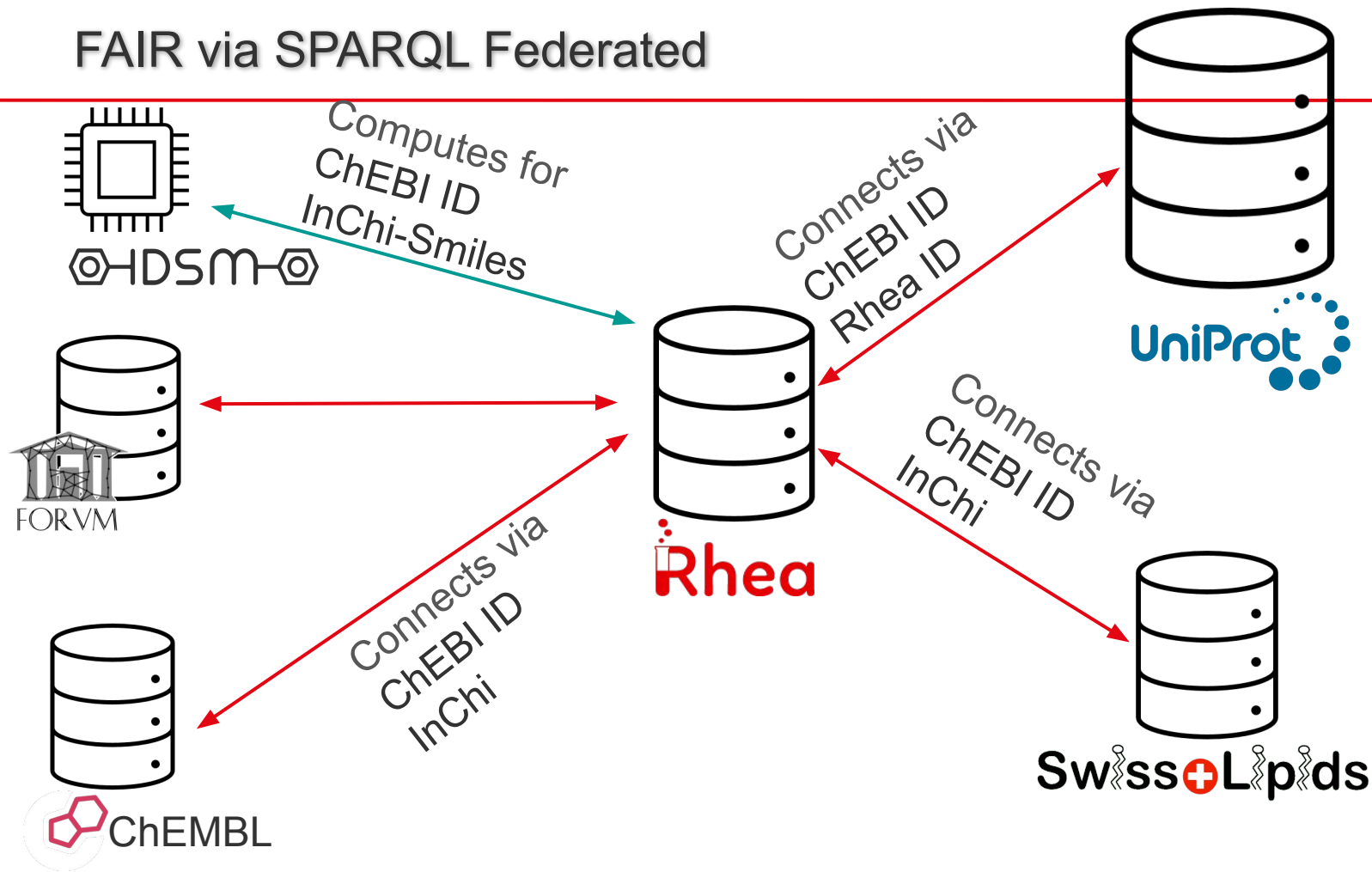


<https://github.com/sib-swiss/sparql-training>

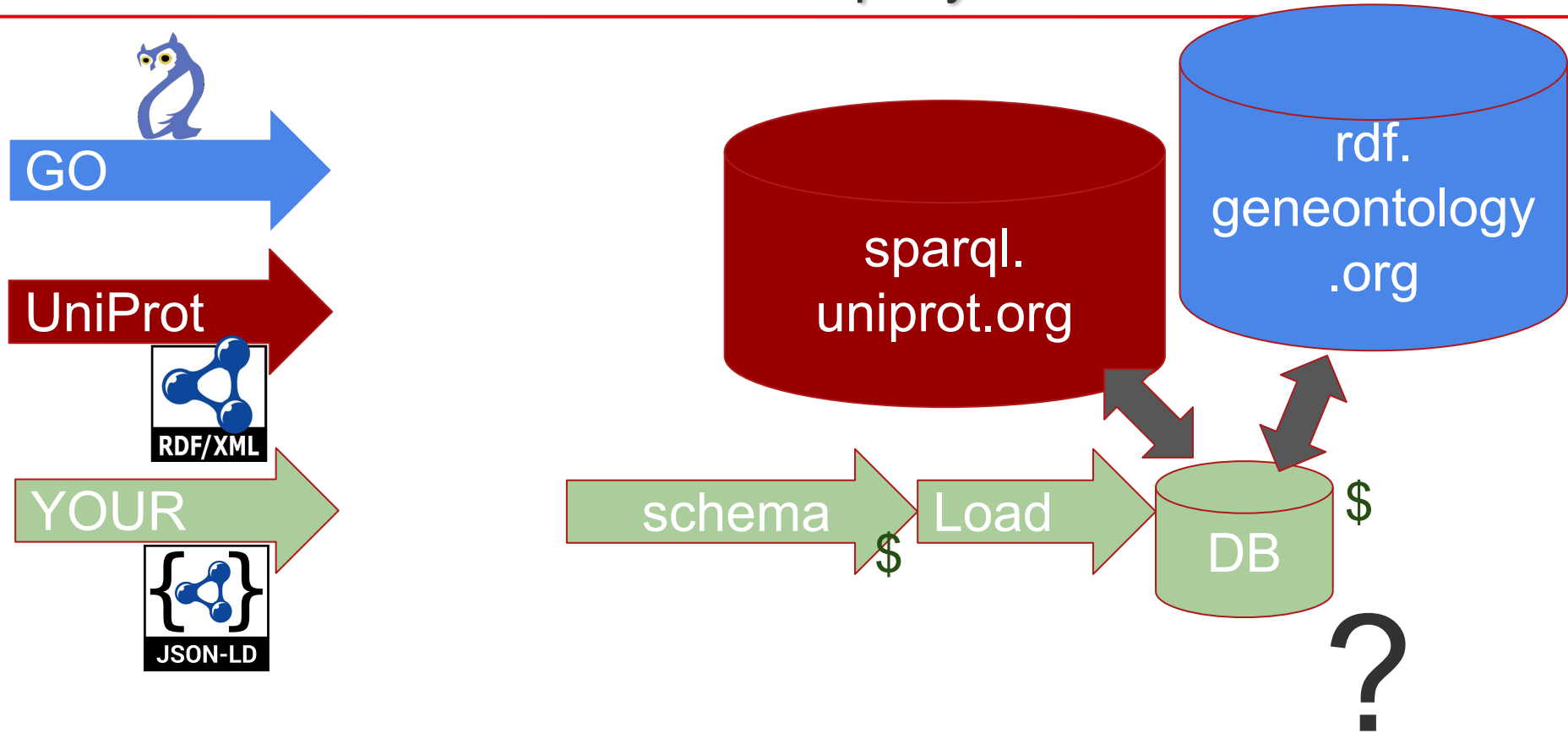
FAIR via SPARQL Federated



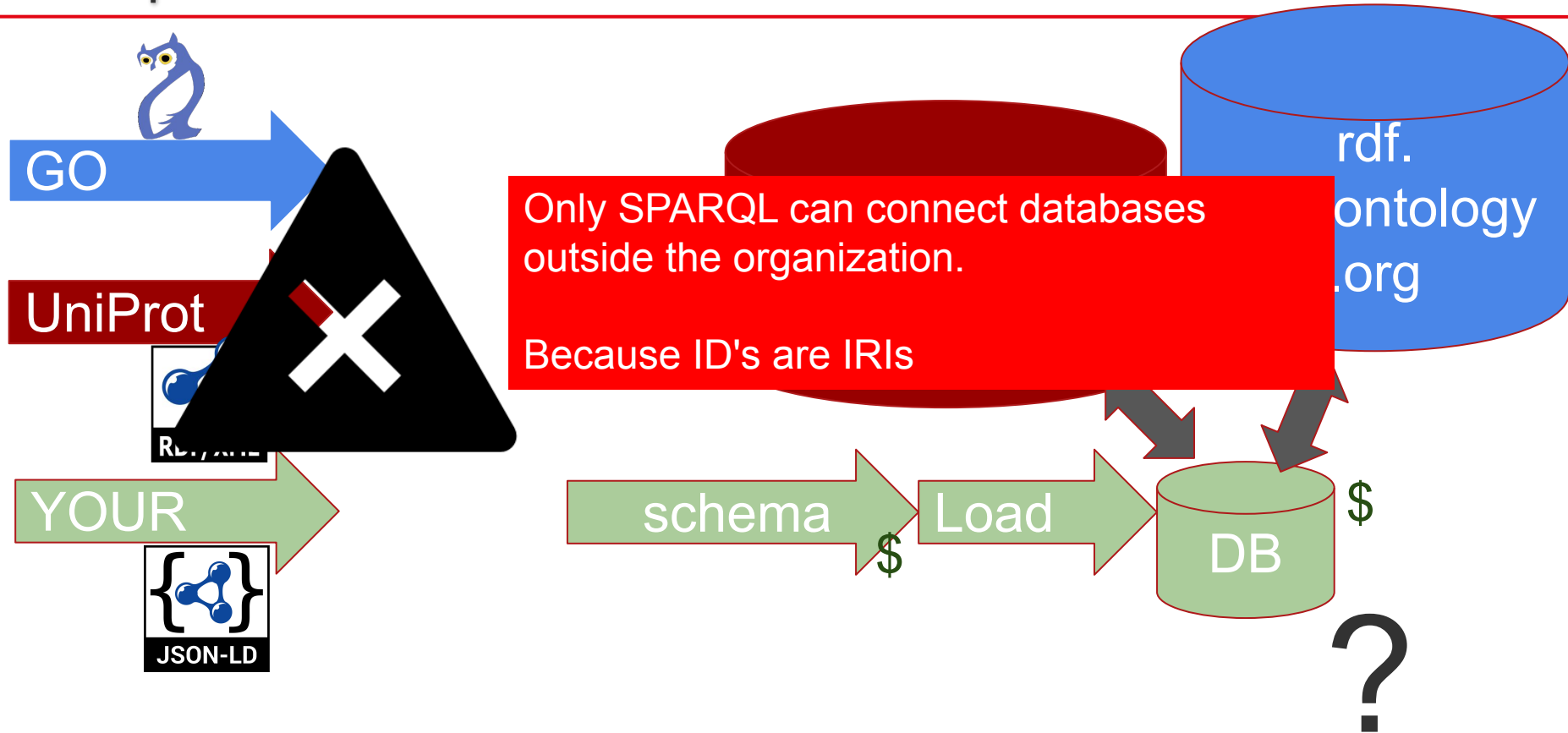
FAIR via SPARQL Federated



RDF + SPARQL 1.1 with build in query federation



Not possible with REST services



Ask DB 2 to be part of the solution



BASE <<http://purl.uniprot.org/uniprot/>>

PREFIX ...

SELECT

?protein ?rhea

WHERE {

SERVICE <<https://sparql.rhea-db.org/sparql>> {

?rhea rdfs:subClassOf rh:Reaction ;

rh:status rh:Approved ;

rh:side/rh:contains/rh:compound/rh:chebi/rdfs:subClassOf+

CHEBI:26739 .

}

?protein up:annotation/up:catalyticActivity/up:catalyzedReaction ?rhea .

}

Issue when federating

- Bugs add up
 - SPARQL 1.1 compliance
 - Downtime
- Optimizers are blind
 - Order
 - Distance
 - Endpoints change
 - Number and kind of triples
 - Implementation

Federation – UniProt to Japanese Medical Dictionary

```
SELECT
  ?protein ?englishLabelStr
WHERE {
  SERVICE <http://data.allie.dbcls.jp/sparql>{
    [] rdfs:label "アミロイド前駆体タンパク質"@ja ;
      rdfs:label ?englishLabel .
    FILTER(lang(?englishLabel) = "en")
  }
  BIND (STR(?englishLabel) AS ?englishLabelStr)
  ?protein
    a up:Protein ;
    (up:recommendedName|up:alternativeName) ?structuredName .
}
```

Federation – UniProt to EU Environmental Database

```
SELECT ?taxon ?ncbiTaxid ?eunisTaxon ?eunisname ?image
WHERE
{
  ?taxon a up:Taxon .
  ?taxon rdfs:subClassOf taxon:8835 .
  BIND(substr(str(?taxon), 35) AS ?ncbiTaxid)
  SERVICE <https://semantic.eea.europa.eu/sparql> {
    ?eunisTaxon a eunisSpecies:SpeciesSynonym ;
      eunisSpecies:binomialName ?eunisname ;
      eunisSpecies:sameSpeciesNCBI ?ncbiTaxid ;
      foaf:depiction ?image .
  }
}
```

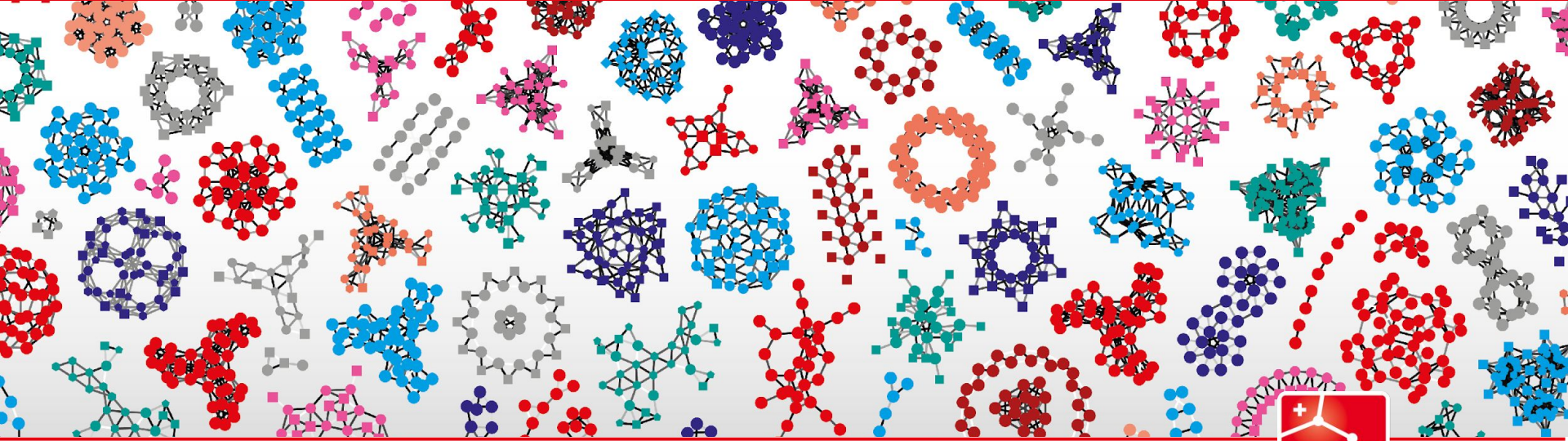
Federation – Rhea to Sachem (compute)

```
SELECT ?chebi ?name ?rhea
WHERE {
  SERVICE <https://idsm.elixir-czech.cz/sparql/endpoint/chebi>
  { ?ssc sachem:compound ?chebi ;
    sachem:score ?ligandSimilarityScore ;
    sachem:similaritySearch/sachem:query
    ""CC1=C(CCC([O-])=O)C2=...=c(C)c(CCC([O-])=O)c1=C2"";
    sachem:cutoff "8e-1"^^xsd:double ;
    sachem:aromaticityMode sachem:aromaticityDetect ;
  }
  ?rhea
  rh:side/rh:contains/rh:compound/(rh:chebi|(rh:reactivePart/rh:chebi)|(rh:underlyingChebi/rh:chebi)) ?chebi .
  ?chebi up:name ?name .
}
```

Federation – Query Against Programs

- Override 1 method
 - **RDF4j** - getStatementsInternal
 - **RDFlib** - triples
 - **Jena** - eval
 - **OxiGraph** - encoded_quads_for_pattern
 - **Ruby-RDF** - query-instance_method

Power of all SPARQL



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Knowledge should be shared !