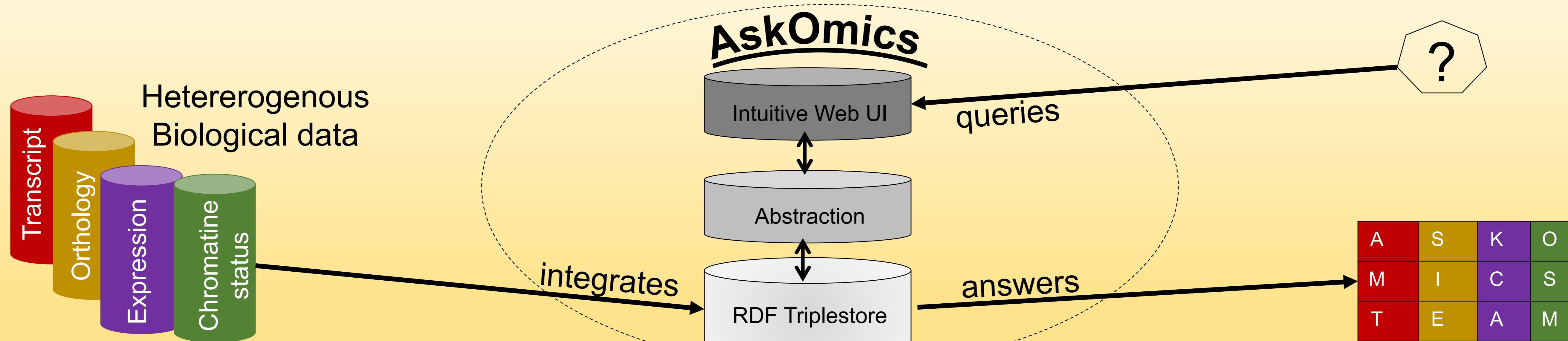


Integration and query of biological datasets with Semantic Web technologies : AskOmics

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AskOmics is a tool supporting both intuitive data integration and querying while shielding the user from most of the technical difficulties underlying RDF and SPARQL



3- AskOmics generates RDF triples {Entity, Relation, Value}

Conversion in RDF triples

```

@prefix : <http://www.w3.org/2000/01/rdf-schema#> .
@prefix rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#> .
@prefix rdfs: <http://www.w3.org/2000/01/rdf-schema#> .
@prefix dc: <http://purl.org/dc/terms/> .
@prefix g: <http://www.semanticsweb.org/iris/ontologies/2010/1/igepp-ontology#> .
@prefix o: <http://www.semanticsweb.org/iris/ontologies/2010/1/igepp-ontology#> .

[g:AT1G01010 rdf:type g:Transcript ; 
  rdfs:label "AT1G01010" ; 
  o:transcript_position "AT1" ; 
  o:chromosomeName "AT1" .]
  
```

Examples of RDF triples

Entity	Relation	Value
AT1G01010	isLocalisedOn	AT1*
BnaC09g15260	isLocalisedOn	BnaC09
AT1G05260	isLocalisedOn	AT1*
AT1G01010	hasExpressionValue	1.6
BnaC09g15260	hasExpressionValue	**
AT1G05260	hasExpressionValue	10.1

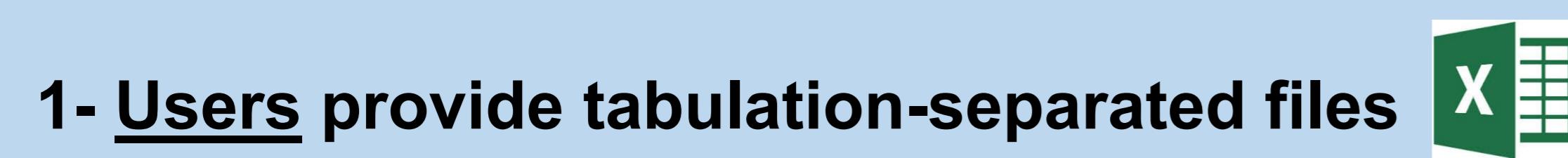
*Redundant value ok
**Empty value ok

4- AskOmics stores RDF in a Triplestore



2- Users upload files in the Web AskOmics interface

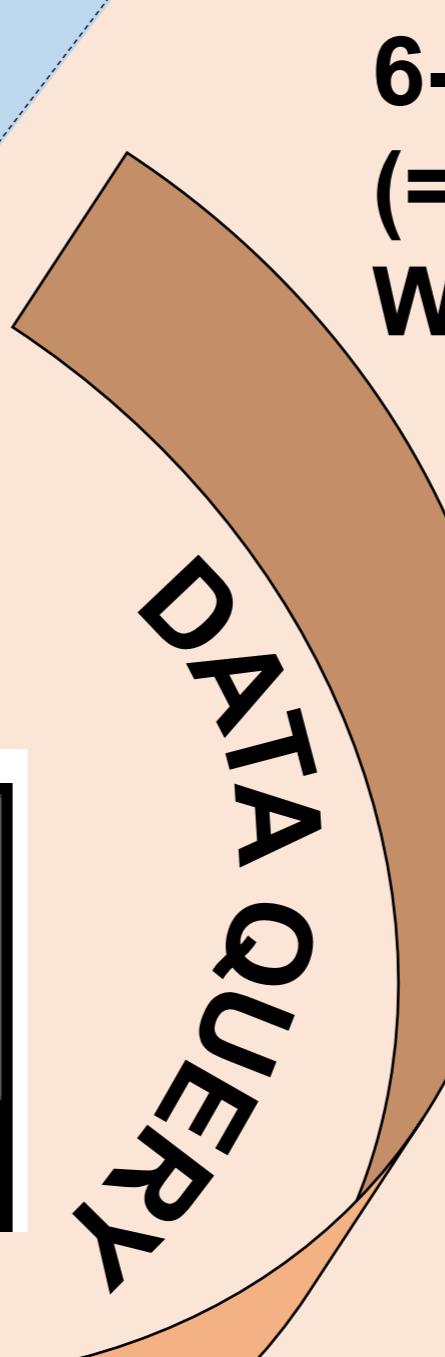
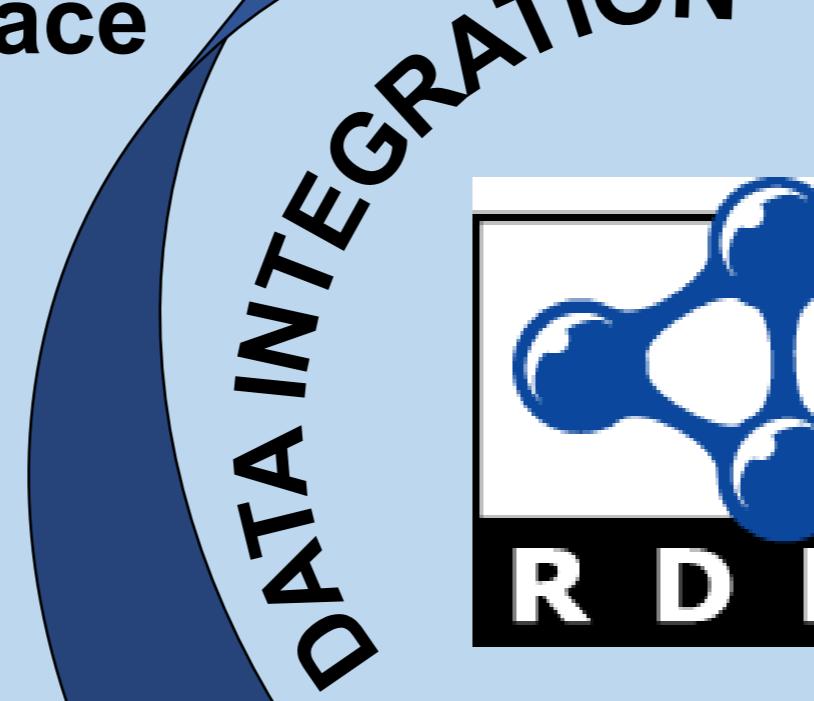
1- Users provide tabulation-separated files



F	G
Transcript	ExpressionValue
AT1G01010	1.6
AT1G05260	10.1
BnaC09g15260	

F	G
Transcript	ChromosomeName
AT1G01010	AT1
BnaC09g15260	BnaC09
AT1G05260	AT1

F	G
Transcript	ChromosomeName
AT1G01010	AT1
BnaC09g15260	BnaC09
AT1G05260	AT1



6- Users query their data by selecting nodes (=integrated Entities) on the left panel of the Web interface

=> AskOmics translates the user's selection into Sparql and queries the RDF Triplestore

7- Users restrict their biological results by applying specific filters from the right panel of AskOmics interface

8- Users download their results

Coming soon:

Integrate positional intervals for query

Support federated queries with external RDF endpoints

