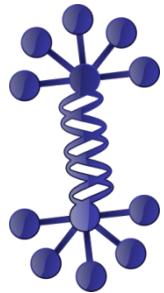


Analysis and visualisation of RDF resources in Ondex

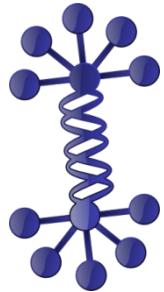
Catherine Canevet

catherine.canevet@bbsrc.ac.uk



Motivation

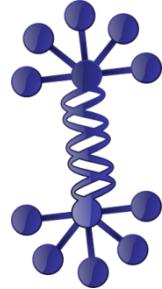
- Biomedical resources provide info on the Semantic Web
- Potential to advance biomedical research
- Need for tools
 - Data reduction
 - Data visualization
 - Interactive analysis capabilities



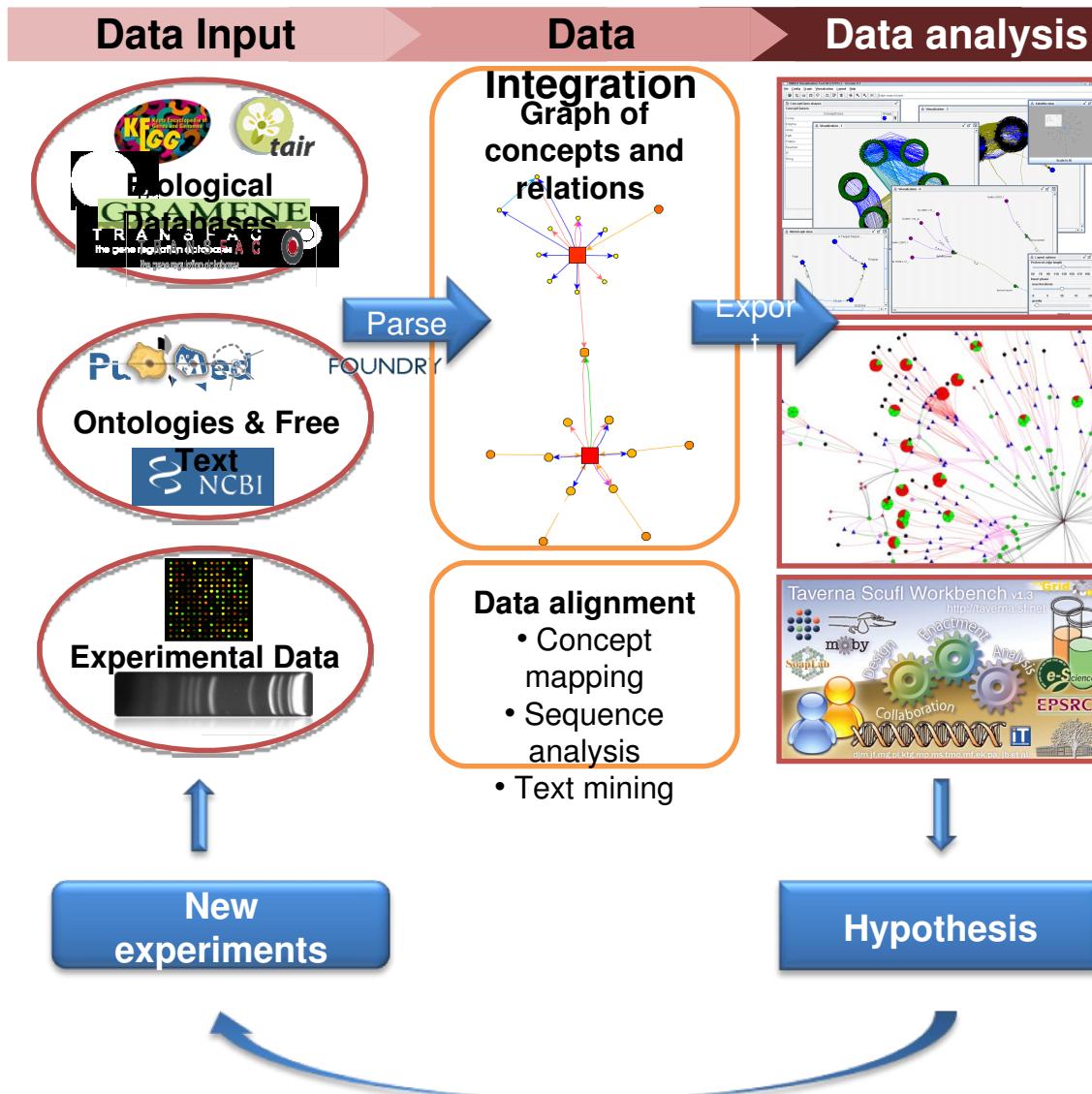
Ondex: a platform to exploit the Semantic Web

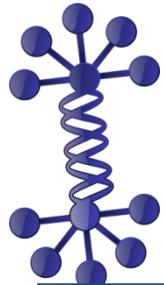
- 1) all information can be represented as a graph
- 2) all elements of the graph can be annotated with ontologies
- 3) data model conformant to Semantic Web framework, in particular to RDF

→ Ondex ideally positioned

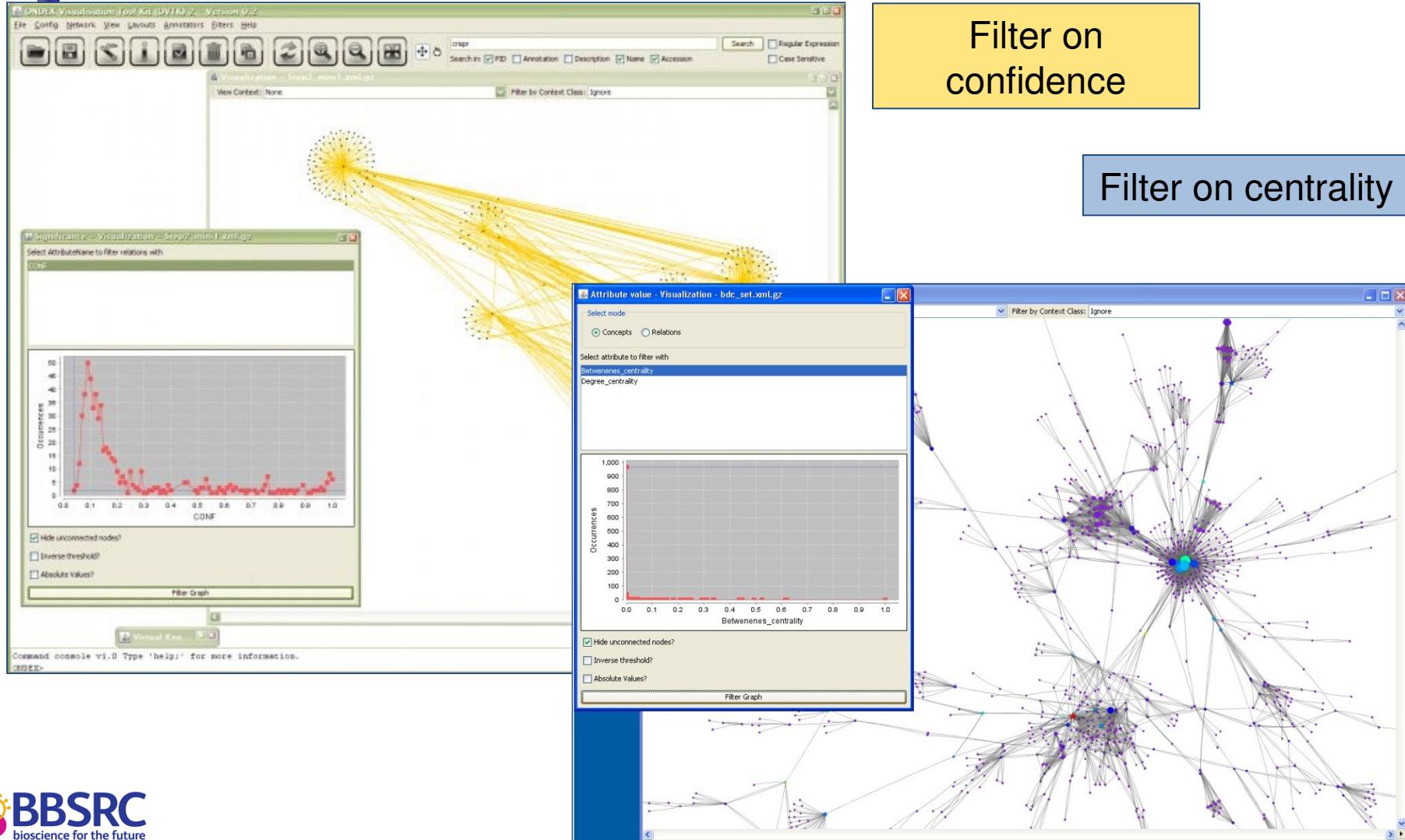


Data integration and visualisation in Ondex





Data reduction and representation methods



Nature Precedings : doi:10.1016/j.npre.2010.5430.1 : Postscript Dec 23, 2010

File Configure Help

- Export**
 - Experimental
 - Discontinued
 - Stable
 - OXL Export
 - Report writer
- Filter**
 - Discontinued
 - Stable
 - Experimental**
- Graph**
- Mapping**
 - Stable
 - Concept accession-based mapping (Memory-efficient)
 - Inparanoid
 - Experimental**
- Parser**
 - Stable
 - Aracyc2 (Release 6, October 14, 2009)
 - BCSDB (pre-release BCSDB-3 gamma version)
 - Brenda specific SBML Parser (BRENDA release online since 2nd July 2010)
 - EXPASY ENZYME (4/10/2009)
 - FASTA file parser
 - KEGG parser (Release 52.1, December 1, 2009)
 - KEGG parser, latest (Release 53.0, January 1, 2010)
 - OXL Parser
 - UniProt (UniProt release July2010)
 - Experimental**
- Statistics**
- Transformer**
 - Stable
 - Relation Collapser

New |

New memory graph

GraphName	default
Graph id	default

TAIR 9

Annotation	<input type="checkbox"/>
InputDir	<input type="button" value="Browse"/>
Graph id	default

UniProt

InputDir	<input type="checkbox"/>
TaxId	<input type="checkbox"/>
DbRefAcc	<input type="checkbox"/>
Accessions	<input type="checkbox"/>
AccessionsFile	<input type="checkbox"/>
ContextInformation	<input checked="" type="checkbox"/>
HideLargeScaleRef	<input checked="" type="checkbox"/>
GoFile	<input type="checkbox"/>
Graph id	default

Concept accession-based mapping (Memory-efficient)

EquivalentCC	<input type="checkbox"/>
EquivalentCV	<input type="checkbox"/>
GDSEqualsConstraint	<input type="checkbox"/>
IgnoreAmbiguity	<input type="checkbox"/>
RelationType	equ
WithinCVMapping	<input type="checkbox"/>
ConceptClassRestriction	<input type="checkbox"/>
CVRestriction	<input type="checkbox"/>
Graph id	default

Relation Collapser

RelationType	<input type="checkbox"/>
ConceptClassRestriction	<input type="checkbox"/>
CVRestriction	<input type="checkbox"/>
CloneGDS	<input checked="" type="checkbox"/>
CreateGDSSets	<input type="checkbox"/>
Graph id	default

OXL Export

ExcludeConceptsOfConceptClass	<input type="checkbox"/>
ExcludeRelationsOfRelationType	<input type="checkbox"/>
ExcludeGDSWithAttribute	<input type="checkbox"/>
IncludeOnlyGDSAtributte	<input type="checkbox"/>
IncludeOnlyConceptClass	<input type="checkbox"/>
IncludeOnlyRelationType	<input type="checkbox"/>
Pretty	<input checked="" type="checkbox"/>
ExportIsolatedConcepts	<input checked="" type="checkbox"/>
GZip	<input type="checkbox"/>
ForceTTL	<input type="checkbox"/>

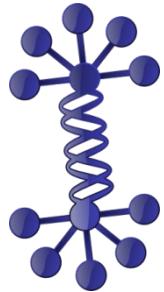
Arguments | Documentation | Required data | Comments |

a subset of the TAIR9 database parsing genes, proteins, domains

Annotation If true also GO annotations, protein domain information and UniProt accessions will be parsed

InputDir Absolute path to input directory

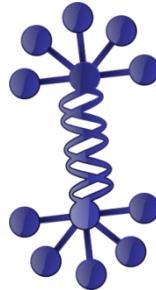
Graph id Graph that will be operated on by this plugin.



Demo

- Interactive importer based on SPARQL
- Query-driven construction of datasets
- Bring information from different RDF data resources into Ondex

- Welcome feedback on our prototype



Data used in demo integrated by Keywan Hassani-Pak

1. Input

45778 predicted poplar genes

2. Data Integration

PoplarCyc, Poplar TFDB

3. Comparative Genomics

BLAST against UniProt Plants

HMMer against Pfam

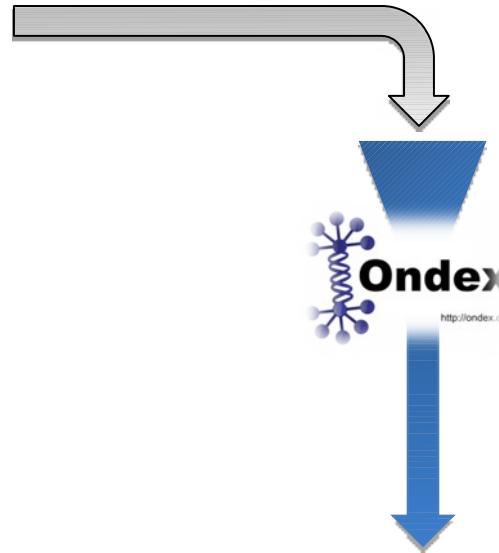
Inparanoid to Arabidopsis

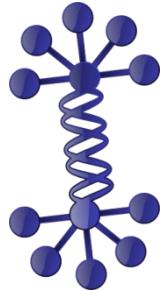
4. Autom. Function Prediction

Gene, Plant and Trait Ontology

5. Text Mining

Co-occurrence of genes and traits in Medline abstracts





Connecting and querying

The screenshot shows the Ondex software interface, version 0.3 (build null), running on a Windows operating system. The window title is "Ondex - Version 0.3 (build null)".

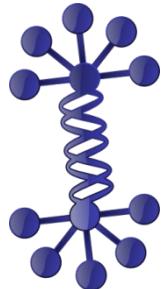
Item Information window:

- General Information**:
 - Name: <http://www.ondex.org/concept/313>
 - Parser ID: <http://www.ondex.org/concept/313>
 - Type: TO
 - Data Source: <http://babvs03.rothamsted.bbsrc.ac.uk:8050/sparql>
 - Evidence: IMPD
- Tags**
- Synonyms**:
 - <http://www.ondex.org/concept/313>
 - shoot branching
- Accessions**
- URI**:
 - <http://www.ondex.org/concept/313>

Visualization window: This window is currently empty.

Console window:

```
Command console v1.0 Type 'help;' for more information.  
ONDEX>sparql_mode  
Changing configuration - please wait...  
SPARQL interpreter loaded successfully  
ONDEX>connect("http://babvs03.rothamsted.bbsrc.ac.uk:8050/sparql")  
ONDEX>construct (?x ?y "shoot branching")  
where {graph ?g {?x ?y "shoot branching"}  
ONDEX>
```



Right-click pre-defined queries

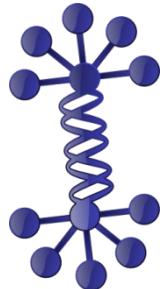
Screenshot of the Ondex Version 0.3 interface showing the right-click context menu for pre-defined queries.

The interface includes:

- File Edit View Appearance Tools Help** menu bar.
- Toolbar** with various icons for file operations and search.
- Search Bar** with fields for "Enter search here...", "Restrict by: Concept Class", "Data Source", "Tag", and "Case Sensitive".
- Item Information** window showing General Information, Tags, Synonyms, Accessions, and URI.
- Visualization - 0** window (empty).
- Console** window displaying SPARQL commands and their execution results.
- Context Menu** (highlighted with a yellow arrow) for the visualization area, listing options like Link, Hide, Show, Show / Edit Properties, Query (selected), Find relations subjects of, Find relations objects of, Find subjects of, Fill attributes, Find objects of, and Resolve URL.

```

Command console v1.0 Type 'help;' for more information.
ONDEX>sparql_mode
Changing configuration - please wait...
SPARQL interpreter loaded successfully
ONDEX>connect("http://babvs03.rothamsted.bbsrc.ac.uk:8050/sparql")
ONDEX>construct (?x ?y "shoot branching")
where {graph ?g (?x ?y "shoot branching")}
ONDEX>
  
```



Filling attributes

Ondex - Version 0.3 (build null)

File Edit View Appearance Tools Help

Enter search here... Restrict by: Concept Class Data Source Tag Case Sensitive

Item Information

General Information

Name: <http://www.onDEX.org/concept/292>
Parser ID: 13130077
Type: Publication
DataSource: <http://babv03.rothamsted.bbsrc.ac.uk:8050/sparql>
Evidence: IMPD

Annotation

[NUCLEOTIDE SEQUENCE [mRNA], FUNCTION, TISSUE SPECIFICITY, DEVELOPMENTAL STAGE, MUTAGENESIS OF ALA-49 AND ARG-50]

Tags

Synonyms

<http://www.onDEX.org/concept/292>
PMID:13130077

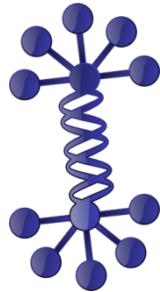
Accessions

<http://babv03.rothamsted.bbsrc.ac.uk:8050/sparql:10.1073/PNAS.1932414100> 13130077

Publication

LAX and SPA: major regulators of shoot branching in rice.
Keishi Komatsu; Masahiko Maekawa; Shin Ujiiie; Yuzuki Satake; Ikuo Furutani; Hiromoto Okamoto; Ko Shimamoto; Junko Kyozuka
Year: 2003
Published in: Proc. Natl. Acad. Sci. U.S.A.
The aerial architecture of plants is determined primarily by the pattern of shoot branching. Although shoot apical meristem initiation during embryogenesis has been extensively studied by molecular genetic approaches using *Arabidopsis*, little is known about the genetic mechanisms controlling axillary meristem initiation, mainly because of the insufficient number of mutants that specifically alter it. We identified the LAX PANICLE (LAX) and SMALL PANICLE (SPA) genes as the main regulators of axillary meristem formation in rice. LAX encodes a basic helix-loop-helix transcription factor and is expressed in the boundary between the shoot apical meristem and the region of new meristem formation. This pattern of LAX expression was repeatedly observed in every axillary meristem, consistent with our observation that LAX is involved in the formation of all types of axillary meristems throughout the ontogeny of a rice plant. Ectopic LAX expression in rice caused pleiotropic effects, including dwarfing, an altered pattern of stem elongation, darker color, bending of the lamina joint, absence of the midribs of leaves, and severe sterility.

Visualization - 0



Proteins related to shoot branching

Ondex - Version 0.3 (build null)

File Edit View Appearance Tools Help

Enter search here... Regular Expr. Restrict by: Concept Class Data Source Tag Case Sensitive

Item Information

General Information

Name: <http://www.onDEX.org/concept/159>
Parser ID: LAX
Type: Protein
DataSource: <http://babv03.rothamsted.bbsrc.ac.uk:8050/sparql>
Evidence: IMPD

Tags

Synonyms

<http://www.onDEX.org/concept/159>
LAX
LAX1
LOC_Os01g61480
Os01g0831000
P0446G04.27
Transcription factor LAX PANICLE

Accessions

<http://babv03.rothamsted.bbsrc.ac.uk:8050/sparql>: AP003252
Q7XQ6 PS50888 IPR001092 AB115668 4327431
LOC_Os01g0831000 OSA:4327431 OS_38423 PF000010
NP_001044699.1 Os01g0831000

Taxonomy

39947

Aminoacid sequence (AA):

```
MDIDPRGFPI KFQPYHLMPT AGGLGEGRMR
GGGERRRPAK LSTDPOSVAA RERIRRISDR
F RULRSILUP GGSKMDTUSM LEQAINIVKTF
LKAQVTLHQIA ALVQHDESGCQ HADUAAAATSA
AD ADIALEL MHRHGGAGDE DAGMTTEMA
PMEAVNGYGD GPANQMMQQA LDPAQQLMIG
```

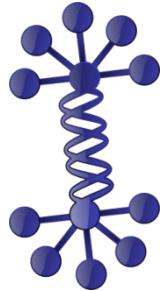
Metadata Legend

Concept Classes	Data Sources	RelationTypes	EvidenceTypes
	Publication	12 (12)	
	TO	7 (7)	
	Protein	8 (8)	
Total Concepts 27 (27)			
Copy to clipboard			

Visualization - 0

Console

```
Command console v1.0 Type 'help;' for more information.
ONDEX>sparql_mode
Changing configuration - please wait...
SPARQL interpreter loaded successfully
ONDEX>connect ("http://babv03.rothamsted.bbsrc.ac.uk:8050/sparql")
ONDEX>construct (?x ?y "shoot branching")
where (graph ?g (?x ?y "shoot branching"))
ONDEX>
```



Colour proteins by taxonomy ID

Ondex - Version 0.3 (build null)

File Edit View Appearance Tools Help

Enter search here... Regular Expr. Case Sensitive

Select attribute to annotate concepts with

- DOI
- JOURNAL_REF
- JOURNAL_REF:0
- MeSH
- PUB_TYPE
- TAXID
- URI
- YEAR

Set blanks to white?

Exclusive use of attribute?

Annotate Graph

Legend Colour Concepts

Color	
3694	purple
39947	green

Accessions

<http://babv03.rothamsted.bbsrc.ac.uk:8050/sparql>: AP003252
Q7XAQ6 PS50888 IPR001092 AB115668 4327431
LOC_Os01g0831000 OSA4327431 OS_38423 PF00010
NP_001044699.1 Os01g0831000

Taxonomy

39947

Aminoacid sequence (AA):

```
MDIDPGRGTFP I KFQPYHMLPTT AGGLGEGRMR
GGGERRRPAK LSTDPQSVAA RERIRRISDR
F RULRSILUP GGSKMDTVSM LEQAINIVKTF
LKAQVTLHQIA ALVQHDESGCQ HADUAAAATPSA
AD ADIALEL MHRHGGAGDE DAGMTTELEMA
PMQEAVNGYGD GPANQMMQQA LDPAQQLMIG
```

Metadata Legend

Concept Classes Data Sources RelationTypes EvidenceTypes

Icon	Type	Count
Orange square	Publication	12 (12)
Blue square	TO	7 (7)
Red square	Protein	8 (8)

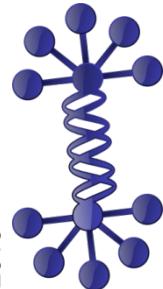
Total Concepts 27 (27)

Copy to clipboard Refresh Metadata

Visualization - 0

Console

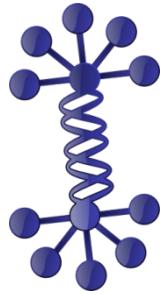
```
Command console v1.0 Type 'help;' for more information.
ONDEX>sparql_mode
Changing configuration - please wait...
SPARQL interpreter loaded successfully
ONDEX>connect ("http://babv03.rothamsted.bbsrc.ac.uk:8050/sparql")
ONDEX>construct (?x ?y "shoot branching")
where (graph ?g (?x ?y "shoot branching"))
ONDEX>
```



Scale relation based on BLAST

The figure shows the ONDEX version 0.3 interface with several windows open:

- Scale/Colour Relations by N...**: A configuration window for scaling relations. It includes fields for "Min Relation Size" (1), "Max Relation Size" (10), and checkboxes for "inverse scaling", "colour edges", and "No attribute colour".
- Legend Colour Con...**: A legend window showing color swatches for concept IDs 3694 (purple) and 39947 (green).
- Visualization - 0**: A main window displaying a network graph of concepts. Nodes are represented by colored shapes (triangles, squares, circles) and are labeled with URLs such as <http://www.ondex.org/concept/211>. Edges connect nodes, representing relationships between concepts.
- Metadata Legend**: A window showing the distribution of concept classes. It lists "Publication" (12 (12)), "TO" (7 (7)), and "Protein" (8 (8)). The total number of concepts is 27 (27). A "Copy to clipboard" button is available.
- Console**: A terminal window showing command-line interactions. The session starts with "Command console v1.0 Type 'help;' for more information.", followed by "ONDEX>sparql_mode", "Changing configuration - please wait...", "SPARQL interpreter loaded successfully", and a series of SPARQL queries related to "shoot branching".



Acknowledgements

- Artem Lysenko
- Andrea Splendiani
- Matthew Pocock
- Chris Rawlings