PSICQUIC

The PSI Common QUery Interface

Interesting facts
pronounced "psykick", but also known as "pisquick"
spelled in 40 different ways (PSIQUIC, PSICIQK, QPSICUI...)



About me

- Name: Bruno Aranda
- Affiliation: European Bioinformatics Institute (EBI)
- Role: Software Engineer at the IntAct Team
- IntAct provides Molecular Interaction data through its open source services and tools.
- One of such tools is PSICQUIC.

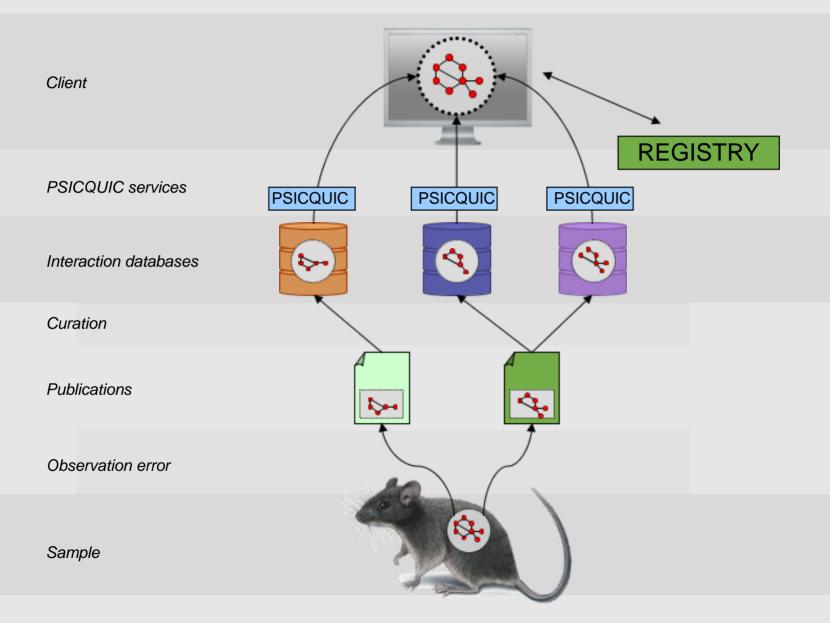


Take a deep breath

INTRODUCTION



What is PSICQUIC?





What is PSICQUIC?



- Proteomics Standards Initiative Common QUery InterfaCe.
- Community effort to standardise the way to access and retrieve data from Molecular Interaction databases.
- PSICQUIC is a specification of a web service.
- Resources already implementing PSICQUIC are listed in a registry.
- Based on the PSI standard formats (PSI-MI XML and MITAB)
- Documentation: http://psicquic.googlecode.com



Justification

• So basically, if all interaction databases provide data in the same way, we can have...





"The one Client to rule them all"





MatrixDB

845

228,262

STRING

12,231,763



404,453

MINT

124,473

More than 14 million binary interactions available using **PSICQUIC**



24,268





581,858



20,769



337,957

416,124

APID



What can I do?

METHODS



Web Service Methods

- getByInteraction
 Retrieves interactions by using an interaction AC.
- getByInteractionList
 Retrieves interactions by using a list of interaction AC.
- getByInteractor
 Retrieves interactions by using a participant identifier.
- getByInteractorList
 Retrieves interactions by using a list of participant identifiers.
- getByQuery
 Retrieves interactions by using a Molecular Interaction Query Language (MIQL)
 query (full text searches)



Web Service Methods

Other metadata methods:

- getVersion
 Returns the version of the web service implementation.
- getSupportedDbAcs
 Returns the supported database identifiers
- getSupportedReturnTypes
 Returns the list of available format types for the results.
- A limited number of interactions can be fetched. It is possible to retrieve large datasets using **pagination**. Most methods have two additional parameters:
 - First result. Index for the first result to retrieve.
 - Max results: Number of interactions returned per query.



Take a shower before going to sleep?

SOAP AND REST (PROTOCOLS)



How can I access PSICQUIC?

As PSICQUIC is a Web Service, you can access the data:

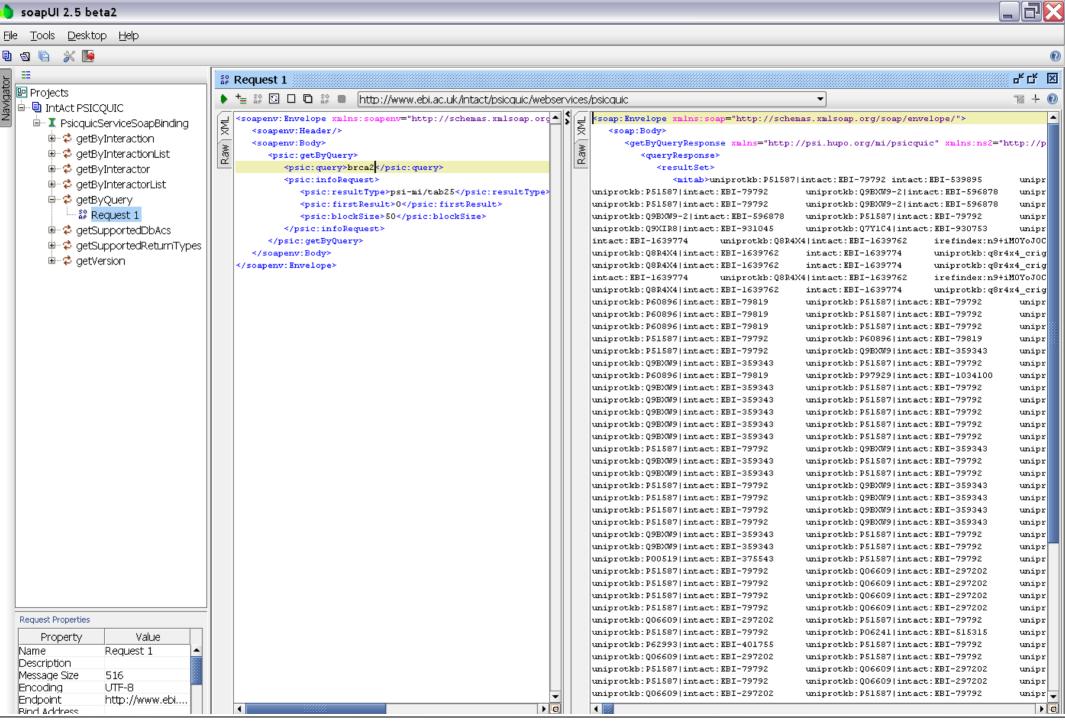
Via SOAP

- A WSDL file exists, and it is the same for all the databases.
- IntAct has developed a Java client, but any other languages can be used.
- You can use it to get interactions in two standard formats: PSI-MI XML and PSI-MI TAB.

Via REST

- Retrieving data directly by using a URL
- Easy to access and data can be obtained just using an internet browser.
- Effective for scripting.

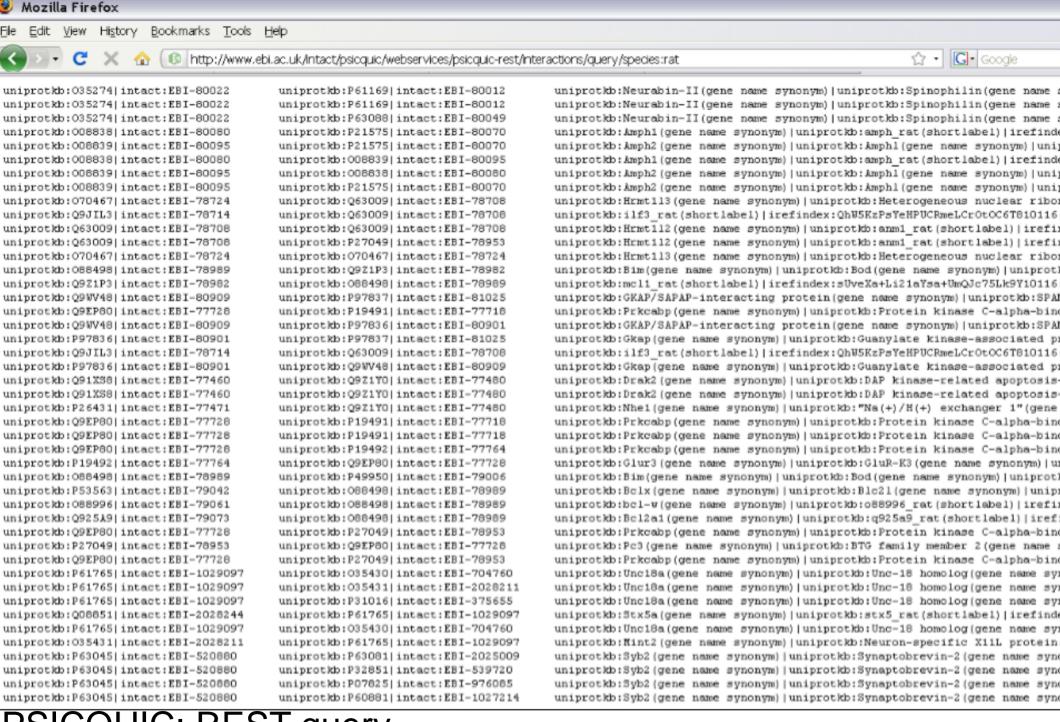






a soapUl 2.5 beta2





PSICQUIC: REST query

http://www.ebi.ac.uk/Tools/webservices/psicquic/intact/webservices/current/search/query/species:rat

Standards and more

FORMATS



Default formats

- The default formats are:
 - PSI-MI XML 2.5.4 (psi-mi/xml25)
 - PSI MITAB 2.5 (psi-mi/tab25)
 - Compressed MITAB (<u>tab25-bin</u>) Only REST
 - Count (count) Only REST
- New formats will be included in the future (work in progress):
 - BioPAX (<u>biopax</u>)
 - And other RDF formats (<u>rdf-xml</u> / <u>rdf-n3</u> / <u>rdf-n3-triple</u> / <u>rdf-turtle</u>)
 - (so it will be possible to use PSICQUIC in the semantic web!)



```
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uniprotkb:P51587|intact:EBI-79792
                                          intact: EBI-539895
                                          uniprotkb:Q9BXW9-2|intact:EBI-596878
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                                                                                                                                                     vnonvm)
                                                                                    uniprotkb:FANCD1(gene name synonym
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uniprotkb:Q9XIR8|intact:EBI-931045
                                          uniprotkb:Q7Y1C4|intact:EBI-930753
                                                                                    uniprotkb: At1g64750 (locus name) | uniprotkb: F13011.6 (orf name) | uniprotk
intact:EBI-1639774
                         uniprotkb:08R4X4|intact:EBI-1639762
                                                                   irefindex:n9+iMOYoJOC9AgBNURd2G9PXjAo10029(rogid)
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uniprotkb:Q9BXW9|intact:EBI-359343
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uniprotkb:Q9BXW9|intact:EBI-359343
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uniprotkb:Q9BXW9|intact:EBI-359343
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biopax

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   xmlns:owlmi="http://purl.org/obo/owl/MI#"
   xmlns:owl="http://www.w3.org/2002/07/owl#"
   xmlns:psimi="http://www.ebi.ac.uk/~intact/psimi.owl#"
   xmlns:xsd="http://www.w3.org/2001/XMLSchema#"
   xmlns="http://www.ebi.ac.uk/intact/"
   xmlns:rdfs="http://www.w3.org/2000/01/rdf-schema#"
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   </br>
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RDF/XML
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    xmlns:owl="http://www.w3.org/2002/07/owl#"
    xmlns:psimi="http://www.ebi.ac.uk/~intact/psimi.owl#"
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    xmlns="http://www.ebi.ac.uk/intact/"
   xmlns:rdfs="http://www.w3.org/2000/01/rdf-schema#"
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    <rdfs:label rdf:datatype="http://www.w3.org/2001/XMLSchema#string">P94102-081303-1</rdfs:label>
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@prefix bp:
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@prefix rdf:
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```

Where are the services?

THE REGISTRY



The PSICQUIC Registry

- It contains a list of the PSICQUIC services from different providers.
- It is a web service itself, and it can be accessed remotely using REST.
- Information can be found about the services, such as the URLs to use, number of interactions provided, versioning, etc.
- The Registry can be found at: http://www.ebi.ac.uk/Tools/webservices/psicquic/registry/registry/action=STATUS



PSICQUIC Registry

Name	Active In	nteractions	Version	SOAP URL	REST URL	REST Example	Restricted	Tags	Comme	
APID	YES	416,124	1.1.5	http://cicblade.dep.usal.es	http://cicblade.dep.usal.es	<u>Example</u>	NO	protein-protein imported spoke clustered		
ChEMBL	YES	581,858	1.1.0	http://www.ebi.ac.uk/Tools	http://www.ebi.ac.uk/Tools	<u>Example</u>	NO	smallmolecule-protein internally curated mimix curation spoke evidence		
<u>InnateDB</u>	YES	9,909	1.1.5	http://imex.innatedb.com/	http://imex.innatedb.com/	<u>Example</u>	NO	protein-protein internally curated spoke rapid curation evidence		
DIP	YES	20,769	1.1.6-SNAPSHOT	http://imex.mbi.ucla.edu/p	http://imex.mbi.ucla.edu/p	<u>Example</u>	NO	protein-protein internally curated imex curation mimix curation spoke evidence		
BioGrid	NO	337,957	1.1.6-SNAPSHOT	http://tyerslab.bio.ed.ac.ul	http://tyerslab.bio.ed.ac.ul	Example	NO	protein-protein internally curated rapid curation spoke evidence		
MINT	YES	124,473	1.1.5	http://mint.bio.uniroma2.it/	http://mint.bio.uniroma2.it/	<u>Example</u>	NO	protein-protein internally curated imex curation mimix curation spoke evidence		
IntAct	YES	228,262	1.1.6-SNAPSHOT	http://www.ebi.ac.uk/Tools	http://www.ebi.ac.uk/Tools	<u>Example</u>	NO	protein-protein smallmolecule-protein nucleicacid-protein		
	More than 14,000,000 binary interactions available from 13 different sources									

http://biotin.uio.no:8080/ps http://biotin.uio.no:8080/ps Example

NO

protein-protein

PSICQUIC: Registry

404,453 1.1.5

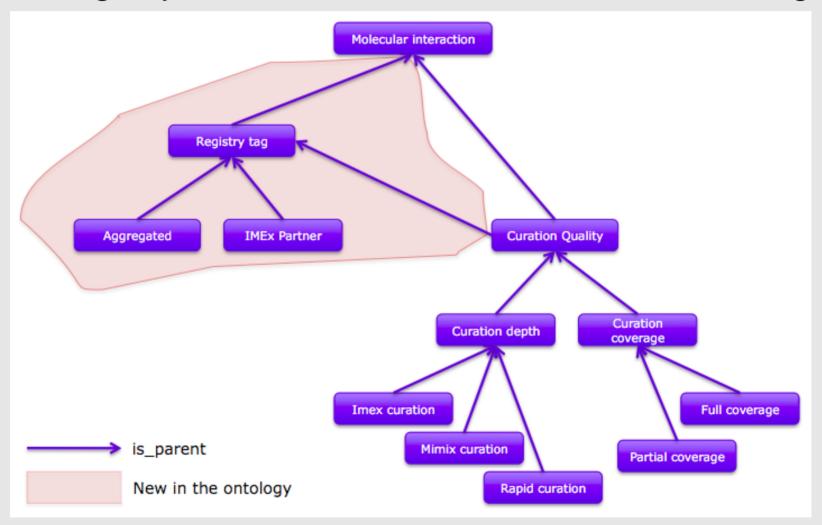
YES

iRefIndex

http://www.ebi.ac.uk/Tools/webservices/psicquic/registry/registry?action=STATUS

Registry Tagging system

The registry classifies the different services with tags.





Querying the registry

- The registry can be accessed with the browser or programmatically (it is a web service).
- Instructions on how to use it can be found here:
 - http://code.google.com/p/psicquic/wiki/Registry
- Querying by tags is work in progress at the moment (it will be explained tomorrow).



Examples

REAL APPLICATIONS



PSICQUIC Applications

- It is clear the value of PSICQUIC to application developers, so indirectly the end-user is benefited too.
- Reduces the time to implement an application that uses data from the different provides, as all of them are accessed the same way.
- Some of the applications:
 - Cytoscape 2.7.x
 - PSICQUIC View
 - Envision2
 - PSICQUIC Client for Android
 - o GMOD client?

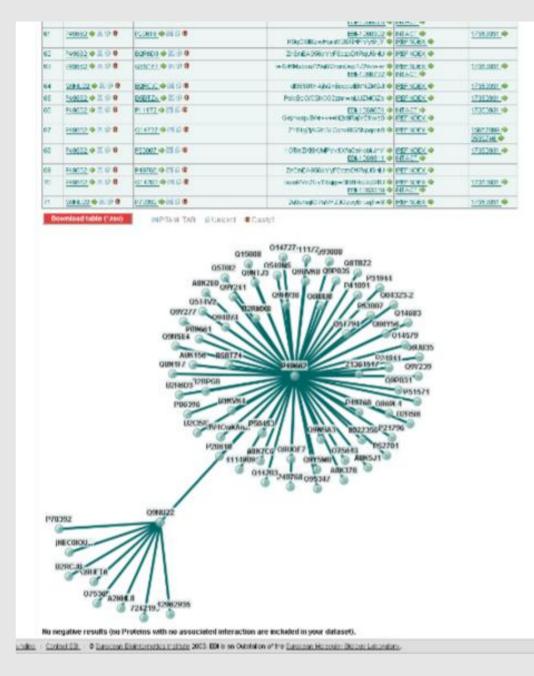


PSICQUIC View

Search: Search Clear Fields »												
<u>C</u>	nEMBL (0)	<u>DIP (0)</u>	IntAct (89)	MINT (22) MPIDB (0) MatrixDB (0) Reactor	me (0) Reactome-Functional-Interactions (29) <u>iRef</u>	findex (31)				
Ext	Export: MITAB 2.5 PSI-XML 2.5.4											
	Name Links Name Links Alt. identifiers molecule A				Alt. identifiers molecule A					Specie		
	molecule A	molecule A	molecule B	molecule B			molecule A	molecule B				
1	<u>P51587;</u> <u>EBI-79792</u>	Uni Prot	EBI-539895		FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human		BRCA2		Human (<u>9606</u>)	-1		
2	<u>P51587;</u> <u>EBI-79792</u>	UniProt	Q9BXW9-2; EBI-596878	UniProt	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human	Q9BXW9-2	BRCA2		Human (9606)	Human (
3	<u>P51587;</u> <u>EBI-79792</u>	Uni Prot	Q9BXW9-2; EBI-596878	Uni Prot	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human	Q9BXW9-2	BRCA2		Human (9606)	Human (
4	<u>P51587;</u> <u>EBI-79792</u>	UniProt	Q9BXW9-2; EBI-596878	Uni Prot	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human	Q9BXW9-2	BRCA2		Human (9606)	Human (
5	CHEBI:15422; EBI-1108845	Sache BI	P51587; EBI-79792	UniProt	atp	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human		BRCA2		Human (
6	CHEBI:15422; EBI-1108845	S Chesi	P51587; EBI-79792	UniProt	atp	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human		BRCA2		Human (
7	CHEBI:15422; EBI-1108845	Chesi	P51587; EBI-79792	Uni Prot	atp	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human		BRCA2		Human (
8	CHEBI:15422; EBI-1108845	Chesi	P51587; EBI-79792	Uni Prot	atp	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human		BRCA2		Human (
9	CHEBI:15422; EBI-1108845	Chesi	P51587; EBI-79792	UniProt	atp	FANCD1; FACD; Fanconi anemia group D1 protein; brca2 human		BRCA2		Human (
10	<u>Q9FL96;</u> <u>EBI-931034</u>	UniProt	Q7Y1C4; EBI-930753	UniProt	At5q45010; K21C13.20; sem12 arath	At5q01630; q7y1c4 arath		brca2b	Mouse-ear cress (3702)	Mouse-e		
11	<u>Q9XIR8;</u> <u>EBI-931045</u>	UniProt	Q7Y1C4; EBI-930753	UniProt	At1q64750; F13O11.6; sem11 arath	At5q01630; g7y1c4 arath		brca2b	Mouse-ear cress (3702)	Mouse-e		
12	Q9XIR8; EBI-931045	Uni Prot	Q7Y1C4; EBI-930753	Uni Prot	At1q64750; F13O11.6; sem11 arath	At5q01630; g7y1c4 arath		brca2b	Mouse-ear cress (3702)	Mouse-6		
13	<u>Q9FL96;</u> <u>EBI-931034</u>	UniProt	Q7Y1C4; EBI-930753	Uni Prot	At5q45010; K21C13.20; sem12 arath	At5q01630; g7y1c4 arath		brca2b	Mouse-ear cress (3702)	Mouse-6		
	nttp://	wwv	v.ebi.	.ac.ı	uk/Tools/webservi	ces/psicquic/view	<u>/</u>	BRCA2	Human (9606)	Human (

Split hand/foot deleted protein 1: dss1_human





Envision2

As an example of PSICQUIC integration

Powerful queries with the Molecular Interaction Query Language

MIQL



Common Query Language

- The Molecular Interactions Query Language (MIQL) allows more powerful and flexible queries.
- It is the default query syntax for PSIQCUIC.
- Designed for fast and effective searches on PSI-MI TAB files.
- All fields (columns) can be searched with specific queries.
- MIQL is a consensus between the different databases, so you should be able to use the same query across different repositories.



MIQL syntax reference

- The MIQL syntax is based on the Lucene syntax[1]. A
 query is broken into terms and operators:
 - Terms: single words or phrases (group of words surrounded by quotes). E.g. brca2 AND "pull down"
 - Fields: search in specific columns. E.g. brca2 AND species:human
 - Term modifiers: wildcard searches, fuzzy searches, proximity and range searches. E.g. brc*
 - Operands: OR (or space), AND, NOT, +, -. E.g. brca2 AND rpa1 / brca2 NOT mouse / +brca2 -mouse -expansion:spoke
 - Grouping and field grouping: brca2 AND (mouse "in vitro")

[1] http://lucene.apache.org/java/docs/queryparsersyntax.html



DIY

HOW TO CREATE YOUR SERVICE



Simplest recipe to implement PSICQUIC



Ingredients:

- PSI-MITAB compliant file.
- Subversion: to get the source code.
- Maven: to run the scripts and start the service.

Steps:

- Generate the MITAB compliant file.
- Get the Reference Implementation (RI):
 - http://code.google.com/p/psicquic/
- Run the script to index the file.
- Start the service with the script provided .



Lots of possibilities

CURRENT AND FUTURE WORK



Future developments

- Smart PSICQUICs: Identification and removal of redundancy
 - Merger and Cluster PSICQUIC services
- PSICQUIC 2.0
 - Overcome the current limitations and many fancy features:
 - Queries using CV terms not possible in the reference implementation (it is possible in IntAct).
 - PSI-MI XML is created from the MITAB, so no n-nary interactions.
 - New features:
 - Redundancy detection mechanism. ROG/RIG ids by default.
 - Built from PSI-MI XML, so complex data available.
- A GMOD component?





Samuel KERRIEN
Jyoti KHADAKE
Sandra ORCHARD
Erick PFEIFFENBERG
Margaret DUESBURY
Marine DUMOUSSEAU
Bruno ARANDA
Henning HERMJAKOB
& former colleagues

Rafael JIMÉNEZ



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The Proteomics Services Team Hinxton Sequence Forum

PSICQUIC

































