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Brainomics - A management system for exploring and merging heterogeneous brain mapping data

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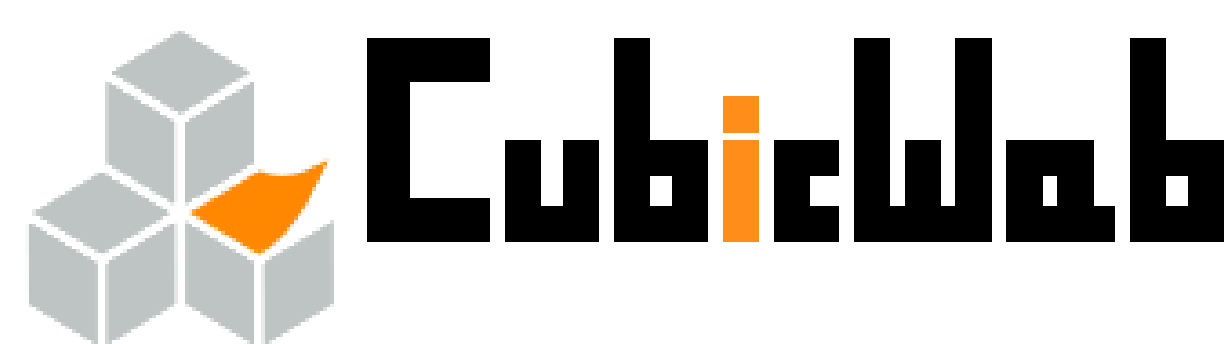
Introduction

- Number of large datasets for brain mapping have been released [1, 2].
- Neuroimaging datasets more routinely include clinical data or genetics data.
- Exploitation requires
 - An efficient organization to integrate all the measures
 - An easy access to the relevant information.
- Neuroimaging [3] and genomics[4] databases are dedicated to their own field of research.

<http://www.brainomics.net/demo/>

- Brings together **brain imaging and genetics data**.
- Relies on a **high-level query language (RQL)**.
- Solution based on **CubicWeb**, a semantic framework.
- Visualizing / exporting data** in several formats.

question	text	score	date
ados_module	Autism Diagnostic Observa		2010-03-19
ados_social	Social Total Subscore of the Classic ADOS	9.0	2010-03-19
ados_stereo_behav	Stereotyped Behaviors and Restricted Interests Total Subscore of the Classic ADOS	4.0	2010-03-19
ados_rsrch_reliable	Was ADOS scored and administered by research reliable personnel?	0.0	2010-03-19
ados_gotham_soc_affect	Social Affect Total Subscore for Gotham Algorithm of the ADOS	4.0	2010-03-19
ados_gotham_rrb	Restrictive and Repetitive Behaviors Total Subscore for Gotham Algorithm of the ADOS	7.0	2010-03-19
ados_gotham_total	Social Affect Total + Restrictive and Repetitive Behaviors Total	16.0	2010-03-19
ados_gotham_severity	Individually Calibrated Severity Score for Gotham Algorithm of the ADOS	6.0	2010-03-19



<http://www.cubicweb.org/>

- ✓ **Data management framework**, 10 years of industrial uses (e.g. [5]).
- ✓ Well established core technologies: **SQL, Python, HTML5, Javascript**.
- ✓ Licensed under **LGPL** since 2008.
- ✓ Used in production environments since 2005.
- ✓ Fine-grained **security system** coupled to the data model definition.
- ✓ **Migration mechanisms** controls model version / ensures data integrity.

Data model

- ✓ Described in Python, using reusable modules called "cubes".
- ✓ Modelisation of Scans, Questionnaires, Genomics results, Behavioral results, Subjects and Studies information.
- ✓ Data model optimized for large volumes (> 2000 subjects).
- ✓ Tested with several publicly available datasets [1, 2].

Query using RQL

- ✓ Similar to the W3C's SPARQL [6].
- ✓ Supports the basic operations (selection, insertion, etc.).
- ✓ Subquerying, ordering, counting, ...

Query all the female subjects of the database, with an age greater than 30
Any S WHERE S is Subject, S age > 30, S gender "female"

Query all the Cmap scans of subjects with an age greater than 25, and that have a score greater than 4.0 for the "algebre" question
Any SA WHERE S is Subject, S age > 25, X is QuestionnaireRun, X concerns S, A is Answer, A questionnaire_run X, A question Q, Q text "algebre", A value > 4, SA is Scan, SA concerns S, SA type "c map"

Views

- ✓ Each query result can be seen using different views.
- ✓ HTML pages, ZIP files, spreadsheets, XCEDE XML, ...
- ✓ May include processing (stat. maps computed on the fly).

Conclusion

- ✓ **Open source solution to manage brain imaging datasets and associated meta data.**
- ✓ **Powerful querying and reporting tool, customized for emerging imaging-genetics field.**

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[1] <http://openfmri.org/data-sets>

[2] fcon_1000.projects.nitrc.org/indi/abide/

[3] Olsen, M.D. (2007). The extensible neuroimaging archive toolkit. Neuroinformatics 5, 11-33

[4] Vallon-Christersson, J. (2009). 'BASE - 2nd generation software for microarray data management and analysis'

[5] Publishing bibliographic records on the Web of data: opportunities for the BnF (French national Library). ESWC 2013

[6] <http://www.w3.org/TR/rdf-sparql-query/>