Vincent Emonet

Semantic web design engineer

Experiences

January 2015 Computer science design engineer - Ontologies & Semantic Web,

to CNRS - Laboratory of Informatics, Robotics and Microelectronics of Montpellier, France.

July 2017 • Administration and new features development on bioportal.lirmm.fr and agroportal.lirmm.fr platforms, used for ontology storage, exploration and text annotation (Ruby on Rails & Java): github.com/sifrproject

Kept compatibility with main version of the NCBO BioPortal (tests)

- Packaged, integrated and documented Natural Language Processing programs, available at *advanse.lirmm.fr* (Java, Angular JS, Linux administration): *gite.lirmm.fr/advanse*
- o Enhanced and packaged the YAM++ ontologies matching tool (Java): gite.lirmm.fr/opendata
- Developped a web interface for ontologies mappings validation (JEE): yamplusplus.lirmm.fr

03/2014 to Bioinformatics design engineer - Biomedical data processing,

09/2014 *Université de Laval*, Québec, Canada.

& \circ Development of web services to access biomedical data in the RDF format for the bio2rdf.org project: 06/2013 to github.com/bio2rdf/bio2rdf-rest

09/2013 O Developped components for the Talend ETL for the Semantic Web (Java): github.com/vemonet/talend4sw

 $\circ \ \ \mathsf{Converting} \ \ \mathsf{biomedical} \ \ \mathsf{databases} \ \ \mathsf{to} \ \ \mathsf{RDF} \ \ \mathsf{format} \ \ \mathsf{(PubMed, neXtProt)} \ \mathsf{using} \ \ \mathsf{Talend} \ \ \mathsf{ETL} \ \ \mathsf{and} \ \ \mathsf{Java}$

Set up a Hadoop platform for Big Data on CentOS server

June 2012 Intern in bioinformatics in an ecology research laboratory,

to University of California, Berkeley, USA.

August 2012 • Developped DNA sequence analysis programs using Python 2.7

o Set up an Amazon Web Service machine image based on Linux

Education

2012 - 2014 Master degree, Physiology, biotechnology and informatics, University of Poitiers, France, Biotechnology.

2011 – 2012 Bachelor's degree, Bioinformatics, University of Poitiers, France, Biotechnology specialization.

2009 – 2011 **Technical College Diploma, Bioengineering**, *Toulon, France*, Biology and biochimistry analysis.

Publications

Platforms • HAL: https://hal.archives-ouvertes.fr/search/index/q/*/authFullName_s/Vincent+Emonet

odblp: http://dblp.uni-trier.de/pers/hd/e/Emonet:Vincent

LIRMM • Anne Toulet et al., Metadata model in an ontology portal (french), 2016

o C. Jonquet et al., Reusing the NCBO BioPortal technology for agronomy to build AgroPortal, 2016

o A. Annane et al., Multilingual Mapping Reconciliation between English-French biomedical ontologies

Bio2RDF • M. Dumontier et al., Bio2RDF Release 3: A larger, more connected network of Linked Data for the Life Sciences. International Semantic Web Conference, 2014

Skills

Languages Java, Ruby (on Rails), HTML, JavaScript, CSS, SQL, Regex, Python

Web RDF, SPARQL, ontologies (OWL, SKOS, OBO, UMLS), RESTful API, Big Data (Hadoop), JSON, XML

Tools Git, Linux (Shell, Debian, RedHat), Docker, ETL Talend et méthodologie Agile

Biology Notions in biotechnology, genetics, biochemistry and physiology

Languages

English TOEIC Score 2014: 935. 5 months in the USA. Active participation in discussion, reading and writing documents Spanish Basic notions for communication

References LIRMM lab

Computer science researchers

- Clément Jonquet (jonquet@lirmm.fr)
- Zohra Bellahsene (bella@lirmm.fr)
- Jérôme Azé (jerome.aze@lirmm.fr)

Bio2RDF project François Belleau,

bio2rdf.org project co-founder (francois.belleau@gmail.com)

UC Berkeley

Adam Retchless, bioinformatics researcher (adam@retchless.us)