

Vincent Emonet

Knowledge graph developer

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Experiences

- Since July 2018 **Data science developer**,
Institute of Data Science, Maastricht University, the Netherlands.
- Developed a cross-platform toolkit to sign and publish RDF Nanopublications using Rust, with bindings to Python, and JavaScript (compiled to WebAssembly). vemonet.github.io/nanopub-rs
 - Developed a python library and documentation to configure and deploy self-hosted open source Large Language Models offline with API and UI. vemonet.github.io/libre-chat
 - Developed a python library to easily deploy SPARQL endpoints using RDFLib. Helped to integrate it to bioregistry.io. github.com/vemonet/rdfliib-endpoint
 - Developed a system with a web service to index standard knowledge representations, such as OWL ontologies and SHACL shapes, from public git repositories, alongside with generating metadata for active SPARQL endpoints (Python, React/TypeScript). Presented as a demo at the Web Conference 2021. index.semanticscience.org
 - Developed a web service and a python library to define, publish, and execute extensible FAIR evaluations of resources online (Python, React/TypeScript). Presented at the International Conference on FAIR Digital Objects in 2022. fair-enough.semanticscience.org and maastrichtu-ids.github.io/fair-test
 - Developed a web service with an interface to annotate biomedical text assisted by NLP, using standard identifiers, and the BioLink model developed for the NIH NCATS Biomedical Data Translator project. Annotated statements are published as RDF Nanopublications. collaboratory.semanticscience.org
 - Developed an API, to serve drug-disease predictions models, and expose the predictions using standards developed for the NIH NCATS Biomedical Data Translator project (Python). openpredict.semanticscience.org
 - Developed a python library and documentation to assist with building knowledge graphs, by mapping structured data to RDF (Python, RML mappings). Published at the SWAT4LS conference. d2s.semanticscience.org
 - Developed the Data Science Research Infrastructure at Maastricht University, an OpenShift cluster for academic researchers and students to deploy Data Science services and workspaces. Managed the cluster with its documentation, and defined the deployment of workspaces for data science on CPU and GPU, such as JupyterHub, RStudio, VSCode server (Kubernetes, Docker). dsri.maastrichtuniversity.nl
 - Deployed and maintained numerous services in production, such as bio2rdf.org and w3id.org/AmIFAIR (Linux).
 - Teaching assistant for the relational database, and semantic web course, at Bachelor level, and for the building and mining knowledge graph course at Master level in the Department of Knowledge and Engineering. Helped students in class, wrote and presented some parts of the courses and practical activities.
- October 2017 **Conversational design engineer**,
to *Match Group*, France.
- June 2018
- Involved in the development and tests of a Natural Language Understanding (NLU) tool to build chatbots.
 - Designed conversational patterns using Google DialogFlow tool.
 - Analyzed conversations flow to improve user engagement using tools such as Google Analytics.
 - Managed the multilingual aspect and translations in 6 languages of the NLU tool developed.
 - Won the French 2018 Digital Innovation Award with the chatbot developed.
- January 2015 **Computer science design engineer - Ontologies & Semantic Web**,
to *CNRS - Laboratory of Informatics, Robotics and Microelectronics of Montpellier (LIRMM)*, France.
- July 2017
- Administration and new features development on the bioportal.lirmm.fr and agroportal.lirmm.fr platforms, used for ontology storage, exploration and text annotation (Ruby on Rails, Java). github.com/sifrproject
 - Contributed to, and maintained compatibility with the NCBO BioPortal developed at Stanford University.
 - Built a web platform to integrate and use Natural Language Processing programs developed in the ADVANSE team (Java, AngularJS). advanse.lirmm.fr
 - 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
 - Set up a Slurm Workload Manager on a supercomputer for research at the LIRMM.
- March 2014 **Bioinformatics design engineer - Biomedical data processing**,
to September 2014 *Université de Laval*, Québec, Canada.
- 2014
- Development of web services to access biomedical data in the RDF format for the bio2rdf.org project. github.com/bio2rdf/bio2rdf-rest
- & June 2013
- Developped components for the Talend ETL for the Semantic Web (Java). github.com/vemonet/talend4sw
- to September 2013
- Converting biomedical databases, such as PubMed and neXtProt, to RDF format using the Talend ETL.
 - Set up a Hortonworks Hadoop platform for Big Data.

- June 2012 **Intern in bioinformatics in an ecology research laboratory,**
to *University of California, Berkeley, USA.*
- August 2012
 - Developed DNA sequence analysis programs to build phylogenetic trees for *Xylella fastidiosa*, using the neighbor-joining method in Python 2.7.
 - Set up an Amazon Web Service machine image for bioinformatics research, 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 biochemistry analysis.

Publications

- Platforms*
 - Google Scholar: <https://scholar.google.fr/citations?user=G59f3woAAAAJ>
 - HAL: https://hal.archives-ouvertes.fr/search/index/q/*/authFullName_s/Vincent+Emonet
 - dblp: <http://dblp.uni-trier.de/pers/hd/e/Emonet:Vincent>
- IDS*
 - V. Emonet, A. Malic, A. Zaveri, A. Grigoriu and M. Dumontier, **Data2Services: enabling automated conversion of data to services**, 2018.
- LIRMM*
 - C. Jonquet et al., **AgroPortal: a vocabulary and ontology repository for agronomy**, *Computers and Electronics in Agriculture*, vol. 144, p. 126-143, 2018.
- Bio2RDF*
 - M. Dumontier, A. Callahan, J. Cruz-Toledo, P. Ansell, V. Emonet, F. Belleau, A. Droit, **Bio2RDF Release 3: A larger, more connected network of Linked Data for the Life Sciences**. *International Semantic Web Conference*, 2014.

Skills

- Languages* Python, TypeScript (React, ExpressJS, NextJS), Rust, Java, Ruby on Rails, Bash, SQL, LaTeX.
- Web* SPARQL, RDF, ontologies (OWL, SKOS, OBO, UMLS), RESTful API, GraphQL API, JSON, XML, HTML, CSS.
- Tools* Git, Linux, Docker, Kubernetes, Talend ETL, Dialogflow, and Agile methodology.
- Biology* Notions in biotechnology, genetics, biochemistry and physiology.

Languages

- French* Mother tongue.
- English* TOEIC Score 2014: 935. Lived 1 year in North America, and 4 years in the Netherlands. Active participation in discussion, reading and writing documents.
- Spanish* Basic notions for communication.

Interests

- Sports* Mountain biking, cyclotourism, hiking.
- Open source enthusiast* I enjoy discovering and using open source softwares, which includes reporting issues, and improving the documentation or implementation when possible.

References

Maastricht University
Michel Dumontier,
Professor of Data Science
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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)