Richard G. Côté

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Profile

15+ years practical experience with large software development projects. Skill and training in life sciences and IT.

Methodical approach to complex problem solving.

Accustomed to work in an Agile environment (Scrum / Kanban).

Professional Experience

R&D Specialist – Software Engineer [Since July 2013]

Nestle Institute of Health Sciences, Lausanne Project management, software development and data architecture for a multi-omics and clinical data warehouse and R interfaces. Member of the cross-project architecture team. Member of the Nestle Big Data task force.

Senior Software Engineer

[May 2005 - May 2013]

European Bioinformatics Institute, Cambridge
Back-end, frond-end and web service development for
several highly accessed projects at the EBI, including the
PRoteomics IDEntification (PRIDE) proteomics data
repository, the Ontology Lookup Service (OLS) and the
Protein Identifier Cross-Reference (PICR) service.

Systems Manager

[October 2004 - March 2005]

McGill University, Montreal | | |

Controlled Vocabularies

Web Services Struts

Hibernate Scrum MuS

Software Engineering

Scripting XML Schema

JAXB WordPress

Data Warehousing
R. Jackson REST Tomcat
GIEINTEILJORM

ava Gimp

MS Office SVN

Photoshop

Business Intelligence and requirement gathering for the McGill University Life Sciences Complex. Coordination with hardware and software

Confluence

Software Developer

[January 2002 – October 2004]

Sequence Bioinformatics, Montreal 🙌

Analysis and bioinformatics consulting to implement or develop custom solutions. Projects included building a web-based LIMS for a breast and ovarian cancer research project (Centre Hospitalier de l'Université de Montréal and Royal Victoria Hospital), integrating external third-party analysis tools and reference databases into a centralize SAAS platform (CGI Life Science Services Centre) and performing various network and software administration tasks.

Research Assistant

[March 2000 - December 2001]

Ste-Justine Hospital, Paediatric Oncology Research Centre, Montreal ■◆■

Port legacy data entry software into web-accessible LIMS. Hardware and software support. Automate common analysis tasks using scripting. Prepare publication-grade figures and graphics.

Skills

Education

D.E.S.S. Applied Computer ScienceUniversité de Montréal

2000 - 2001

B.Sc. Honours Biochemistry McGill University, Montreal

M.Sc. Biochemistry

McGill University, Montreal •••

1997 - 2000

D.E.C. Pure Sciences

CEGEP de l'Outaouais, Hull

1992 - 1994

1994 - 1997

Selected Publications

Côté RG et al., The PRoteomics IDEntification (PRIDE) Converter 2 Framework, Mol. Cell. Proteomics. 2012, PMID: 22949509.

Côté RG et al., The Ontology Lookup Service: bigger and better, Nucleic Acids Res. 2010, PMID: 20460452.

Côté RG et al., The Protein Identifier Cross-Referencing (PICR) service: reconciling protein identifiers across multiple source databases, BMC Bioinformatics 2007, PMID: 17945017.

Full list (26) available upon request.

Restudio Spring Java Script Spring MVC Jetty Git Minimal Data Standards Data Modeling

Ontologies XML MavenOracle

Other information

 $\label{thm:english} \mbox{English and French fluently written and spoken}.$

References are available upon request.

Keywords: Agile Apache Back-End development Clinical Data Confluence Controlled Vocabulary CSS Data Architecture Data Warehouse Database Design Eclipse ElasticSearch Genomics Gimp Git IntelliJ Hibernate Jackson JUnit JAXB JDBC JS JSP Java JavaScript Jetty Jira Kanban LSF Lucene Maven Minimal Data Standards MS Office MySQL nginx ORM Ontologies Oracle PHP Photoshop PostgreSQL Proteomics R REST R Studio SOAP SVN Schema Scrum Software Engineering Struts Spring SpringMVC Web Application Web Services Scripting SVN Tomcat WordPress Web Services XML Schema