RICHARD CÔTÉ

grglcote@gmail.com

+41 (0) 78 653 7213

in ch.linkedin.com/in/rcote/

15+ years practical experience with large software development projects in Life Sciences and IT.

Methodical approach to complex problem solving.

Accustomed to work in an Agile corporate or academic environment.

| 2013 | R&D Specialist - Software Engineer & Technical Project Manager Nestlé Institute of Health Sciences, Lausanne Nestlé Institute of Health Sciences |
|-----------|--|
| 2005 | Senior Software Engineer European Bioinformatics Institute, Cambridge EMBL-EBI |
| 2004 | Systems Manager McGill University, Montréal MCGill UNIVERSITY |
| 2002 | Software Engineer Sequence Bioinformatics, Montréal |
| 2000 | D.E.S.S. Applied Computer Science Université de Montréal, Montréal Université de Montréal |
| 2000 | Software Engineer Hôpital Sainte-Justine, Montréal CHU Sainte-Justine Le centre hospitalier universitaire mêre-enfant |
| 1997 & | M.Sc. Biochemistry McGill University, Montréal Montréal |
| 1994 🔗 | B.Sc. Biochemistry (Honours) McGill University, Montréal WCGill UNIVERSITY |

SKILLS

Languages Java, Python, C#, R, PHP

IDE & Tools IntelliJ, Android Studio,

Visual Studio, R Studio, Eclipse, Git, Subversion, Maven, Confluence, Jira

Database MySQL, PostgreSQL,

Oracle, Data Modelling, Data Warehouse, Master Data Management, ETL,

XML Schema,

Java Hibernate, Spring,

technologies SpringMVC, Struts, JaxB,

JaxWS, Jackson, Swing, Lucene, JSP, Tomcat, Jetty

Mobile Android, Xamarin

Graphics Gimp, Photoshop,

Illustrator

Others Wordpress, MS Office

Perfectly fluent in English and French
Dual nationality ■◆■
B-Permit ••

SELECTED PUBLICATIONS

Côté RG et al., <u>The PRoteomics</u>
<u>IDEntification (PRIDE) Converter 2</u>
<u>Framework</u>, Mol. Cell. Proteomics. 2012.

Côté RG et al., <u>The Ontology Lookup</u> <u>Service: bigger and better</u>, Nucleic Acids Res. 2010.

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

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

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2013



R&D Specialist - Software Engineer & Technical Project Manager, Nestle Institute of Nestlé Institute of Health Sciences, Lausanne

Digital Nutrition & Health: Scrum master, technical project manager, back-end API development and data architecture for connected nutrition intelligence platform.

- Scrum master for development team (4 FTEs)
- Technical project manager for multiple activity streams

Roadmap coordination across project activity streams and internal clients Feature prioritization and sprint planning for the development of multiple mobile applications

Clinical trial protocol design

Mobile application acceptance testing

Coordinate and communicate with internal testers

Monitor testing plan and manage bug reports and new feature requests

Design a data integration and data warehouse schema to integrate and distribute nutritional information

Data modelling and database design

Implement ETL for data load and model validation

Lead architecture for data flow across systems

Collaborate in API definition and refactoring

BioSystems Informatics: Project management, software development and data architecture for a multi-omics and clinical data warehouse with R interfaces. System-to-system integration.

Technical project manager for data integration and distribution project

Write project brief and project plan

Milestone reporting

Design data integration and data warehouse schema to integrate and distribute multiple types of scientific data

Validate, update, refactor and maintain ETL designed from 3rd party contractor Collaborate in the design and implementation of R data access library for data analyst

- Design and implement XML schema for system-to-system message-based integration
- Implement master data management solutions for controlled vocabulary annotations, queries and validation
- Lead project to upgrade production Laboratory Information Management System (LIMS)

First-line support between end-users and vendor

standardize installations between dev/test/prod systems

Implement and secure system-to-system integration

Internal and external compliance testing

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2005

Senior Software Engineer 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.

 Refactoring of the PRIDE proteomics data repository from a small-scale proof of concept to a production-ready system

Data modelling of back-end database

Collaborate in implementing middleware code for data I/O

Collaborate in implementing front-end presentation layer

Design and implementation of the Ontology Lookup Service, a new system for controlled vocabulary gueries and master data management

Data modelling of back-end database

Lead implementation of middleware logic

Lead implementation of front-end presentation layer and web services

Open source strategy for distribution of code and data

Design and implementation of the Protein Identifier Cross-Reference (PICR) service, a new system for protein identifier data integration

Data modelling of back-end database

Lead implementation of middleware logic

Lead implementation of front-end presentation layer and web services

Integration strategies for automated mapping of data submitted to PRIDE

Design and implementation of the PRIDE Converter tool, a rich java application that can convert mass spectrometry file formats into a PRIDE-compatible submission format.

Lead UX design and implement a rich Java client

Integration with previously described web services (OLS, PICR) as part of the submission process to generate well-annotated, syntactically and semantically validated XML files

- Write and publish peer-reviewed articles relating to work on PRIDE, PICR and OLS
- Present PRIDE, PICR and OLS in various scientific conferences, workshops and tutorials
- Supervision of trainee master student (6-month internship).