

Ronald J. Nowling

5927 W Beloit Rd.
West Allis, WI 53219
rnowling@gmail.com
<http://rnowling.github.io/>
<http://www.github.com/rnowling>
954.496.2314

Education

<u>University of Notre Dame</u> , Notre Dame, IN	August 2010 – July 2016
• Ph.D. in Computer Science & Engineering	July 2016
• M.S. in Computer Science & Engineering	May 2015
• Research areas: Math Modeling, Computational Physics, Machine Learning, Bioinformatics, Distributed Systems	
<u>Emporia State University</u> , Emporia, KS	June 2016 – Present
• Graduate Math Classes	
<u>Eckerd College</u> , St. Petersburg, FL	August 2006 – May 2010
• B.S. Computer Science, Mathematics	May 2010
• Thesis: <i>Nature-inspired Metaheuristics for Combinatorial Optimization Problems</i>	

Skills

- Fields: Software Engineering, Math Modeling, Computational Physics, Distributed Systems, Machine Learning, Bioinformatics, Open Source
- Programming Languages: Python, Java, Scala, SQL, CUDA, C/C++, bash
- Libraries: Scikit-learn, Numpy, LAPACK, Apache Spark MLlib
- Testing: Python unittest, JUnit, ScalaTest, Jenkins
- Version Control: Git, Mercurial, SVN, CVS
- Build Systems: Gradle, sbt, lein, CMake
- Distributed and Parallel Frameworks: Apache Spark, Apache Mesos, Condor, CUDA
- Storage and Databases: Gluster, MySQL, SQLite

Professional Experience

<u>Data Science Engineer</u> , AdRoll, Inc.	August 2016 – Present
• Implementing and devising algorithms and models for real-time bidding and conversions	

Software Engineer, Red Hat, Inc.

June 2014 – August 2016

- Engaged in internal consulting around scalable data processing and applications of machine learning
- Proposed and developed method for building customer profiles around interests in knowledgebase article using topic modeling and page view data. Prototyped in Python with scikit-learn and implemented in Apache Spark for production. Presented results at Spark Summit East 2016 and engaged with internal customers to apply pipeline to business problems.
- Consulted on identification of duplicate articles to support a customer portal search engine. Established clear requirements and consistent guidelines for evaluating methods. Demonstrated simple, first-pass method for identifying duplicate articles and proposed extensions to identify false positives.
- Ported pipeline for transforming, cleaning, and summarizing page view data to Apache Spark, reducing run times from days to 1 hour.
- Introduced usage of generative models for generating large, sophisticated data sets for testing big data pipelines without exposing real, confidential data. Presented use cases at Apache Big Data Europe 2015 and contributed open-source implementations to the Apache Bigtop project. Earned appointments as a committer and project management committee (PMC) member.
- Contributed new features and fixes to Apache Spark including performance improvements and new features to machine learning algorithms, random data generators, and tests.
- RPM packaging, integration testing, and deployment of internal analytics stack based on Apache Spark. Provisioning, configuration, and administration of team's Linux clusters

Committer / PMC Member, Apache Bigtop

March 2015 – Present

- Open-source big data distribution including Hadoop, Spark, HBase, Hive, etc.
- Implemented and maintain synthetic data generators and BigPetStore Spark example application
- Various bug fixes and improvements to build system and RPM packages

Research Assistant, University of Notre Dame

August 2010 – July 2016

- Applied machine learning and stochastic models to bioinformatics problems. I demonstrated that machine learning algorithms can work as well as specialized bioinformatics algorithms, enabling scientists to reduce development and maintenance costs by using existing libraries. I used numerical experiments to identify sources of bias when using Random Forests for variable selection and presented solutions that work with existing implementations.
- Derived, implemented, and validated stochastic differential equations and numerical linear algebra algorithms for computational chemistry. Implemented models using parallel, distributed, and GPU programming.
- Served as TA for 3 semesters of Programming Paradigms. Designed and graded assignments, tutored students, and gave guest lectures. Recipient of the GAANN Fellowship (Fall 2012 – Spring 2014) and Kaneb Center Outstanding Teaching Assistant Award (2012).

Simbios OpenMM Visiting Scholar, Stanford University

July 30 – August 24, 2012

- GPU implementations of numerical linear algebra algorithms for computational chemistry

Undergraduate Research Assistant, University of Connecticut

Summers 2005 – 2010

- Collaborated on development of database-driven, Java applications for conversion and processing of molecular structure data

Ford Scholar and Research Assistant, Eckerd College

Summer 2007 – Spring 2010

- Used Genetic Algorithms to solve combinatorial optimization problems; developed novel encoding scheme
- Replaced proprietary file format with SQLite in DARWIN, software for semi-automated identification and cataloging of dolphins using computer vision techniques

Professional Service and Volunteer Efforts

Advisor, Nevada Institute of Personalized Medicine

August 2015 – Present

Co-organizer, Wisconsin Big Data Conference

May 2016 – August 2016

Co-organizer, Milwaukee Big Data Meetup

March 2016 – Present

Oral Presentations

RJ Nowling. Insights into Customer Behavior from Clickstream Data. Spark Summit East (February 2016), New York City, USA.

RJ Nowling. Synthetic Data Generation for Realistic Analytics Examples and Testing. Apache: Big Data Europe (October 2015), Budapest, Hungary.

RJ Nowling. A domain-driven, generative data model for BigPetStore. The 4th IEEE International Conference on Big Data and Cloud Computing (December 2014), Sydney, Australia.

Publications

* denotes equal contribution

RJ Nowling and J Vyas. “A domain-driven, generative data model for BigPetStore.” *Proceedings of the 4th IEEE International Conference on Big Data and Cloud Computing*, 2014.

JC Sweet, **RJ Nowling**, TM Cickovski, CR Sweet, VS Pande, and JA Izquierre. “Long Timestep Molecular Dynamics on the Graphical Processing Unit.” *J. Chem. Theory Comput.*, 9(8):3267–3281, 2013.

RJ Nowling*, JL Abrudan*, DA Shoue, B Abdul-Wahid, M Wadsworth, G Stayback, FH Collins, MA McDowell, and JA Izquierre. “Identification of Novel Arthropod Vector GPCRs.” *Parasit. Vectors*, 6:150, 2013.

HJC Ellis, G Weatherby, **RJ Nowling**, J Vyas, M Fenwick, and MR Gryk. “A Software Architecture for NMR Spectral Data Translation.” *CISE*, 15(1):76-83, 2013.

RJ Nowling and TM Cickovski. "Prototype to Release: Software Engineering for Scientific Software." *Biomedical Computation Review*, Fall 2012.

RJ Nowling, J Vyas, G Weatherby, MW Fenwick, HJC Ellis, and MR Gryk. "CONNJUR Spectrum Translator: An open-source application for reformatting NMR spectral data." *J Bio NMR*, 50:83-89, 2011.

RJ Nowling and H Mauch. "Priority Encoding Scheme for Solving Permutation and Constraint Problems with Genetic Algorithms and Simulated Annealing." *Proceedings of the 8th International Conference on Information Technology - New Generations*, 2011.

HJC Ellis, J Vyas, **RJ Nowling**, TO Martyn and MR Gryk. "Iterative Development Of An Application To Support Nuclear Magnetic Resonance Data Analysis Of Proteins." *Proceedings of the 8th International Conference on Information Technology - New Generations*, 2011.

J Vyas*, **RJ Nowling***, T Meusberger, D Sargeant, K Kadaveru, MR Gryk, V Kundeti, S Rajasekaran, and MR Schiller. "MimoSA: a system for minimotif annotation." *BMC Bioinformatics*, 11:328, 2010.

J Vyas, **RJ Nowling**, MW Maciejewski, S Rajasekaran, MR Gryk, and MR Schiller. "A proposed syntaxfor Minimotif Semantics, version 1." *BMC Genomics*, 10:360, 2009.

Poster Presentations

RJ Nowling. BigTop Bazaar: Simulating Customer Dynamics Driven By Booth Preferences at a Conference. Poster Presented at the 2015 Annual Conference of the Great Lakes Section of the Society for Industrial and Applied Math (2015), Grand Rapids, MI.

RJ Nowling, M Wadsworth, JL Abrudan, DA Shoue, B Abdul-Wahid, GM Stayback, FH Collins, MA McDowell, and JA Izaguirre. Identifying GPCRs in the Genome of the Sand Fly *P. papatasi* using Ensemble*. Poster Presentation at the 7th Annual Arthropod Genomics Symposium (2013), Notre Dame, IN.

RJ Nowling, CR Sweet, and JA Izaguirre. Extending Long Timestep Molecular Dynamics (LTMD) to Explicit Solvent. Poster Presentation at the Midwest Theoretical Chemistry Conference (2013), Urbana-Champaign, IL.

RJ Nowling, JL Abrudan, DA Shoue, B Abdul-Wahid, M Wadsworth, GM Staybak, FH Collins, MA McDowell, and JA Izaguirre. Evaluation and Development of GPCR Classifiers for Vectors. Poster Presentation at the Second Annual Eck Institute for Global Health Research Retreat (2013), Notre Dame, IN.