#### CURRICULUM VITÆ

FULL NAME: Joseph Daniel Romano, MPhil, MA

CURRENT POSITION: PhD Candidate,

Graduate Research Assistant

**INSTITUTION:** Columbia University

CURRENT EMAIL: jdr2160 /at/ cumc.columbia.edu

ADDRESS: Department of Biomedical Informatics

Columbia University Medical Center

622 W. 168<sup>th</sup> Street

Presbyterian Hospital Bldg. 20<sup>th</sup> Floor

New York, NY 10032

LAST UPDATED: Monday 19<sup>th</sup> March, 2018

### **EDUCATION**

2014–present **Doctor of Philosophy** — Prospective (Biomedical Informatics)

Dissertation advisor: Nicholas Tatonetti, PhD

Columbia University; New York, NY

2014–2017 Master of Philosophy (Biomedical Informatics)

Columbia University; New York, NY

2014–2016 Master of Arts (Biomedical Informatics)

Columbia University; New York, NY

2010–2014 Bachelor of Science (Molecular Genetics)

Research Advisor: Neil Sarkar, PhD, MLIS, FACMI

University of Vermont; Burlington, VT

2011–2012 International Student Exchange Program

C.I.E.L.D. (国際関係学部国際教育センター)

Nihon University; Mishima, Shizuoka prefecture, Japan

# **PUBLICATIONS**

- 1. **Romano JD**, Tharp WG, Sarkar IN. "Exploring Complex Disease Gene Relationships Using Simultaneous Analysis." *UVM Honors College Senior Theses.* 2014.
- 2. **Romano JD**, Tharp WG, Sarkar IN. "Adapting Simultaneous Analysis Phylogenomic Techniques to Study Complex Disease Gene Relationships." *Journal of Biomedical Informatics*; 2014 Apr; (54):10–38.
- 3. **Romano JD**, Tatonetti NP. "VenomKB, a new knowledge base for facilitating the validation of putative venom therapies." *Scientific Data*; 2015 Nov;(2):65.
- 4. Boland MR, Jacunski A, Lorberbaum T, **Romano JD**, Moskovitch R, Tatonetti NP. "Systems Biology Approaches for Identifying Adverse Drug Reactions and Elucidating Their Underlying Biological Mechanisms." WIREs Systems Biology and Medicine; 2016 Mar;8(2):104–22.
- 5. **Romano JD**, Tatonetti NP. "Using a Novel Ontology to Inform the Discovery of Therapeutic Peptides from Animal Venoms." *AMIA Joint Summits on Translational Science Proceedings* (Second place, TBI student paper competition).; 2016 Jul;(2016):209-18.

- 6. Vanguri RS, **Romano JD**, Lorberbaum T, Nwankwo V, Youn C, Tatonetti NP. "nSides: An interactive drug–side effect gateway." *Gateways 2017*.
- 7. Vanguri RS, **Romano JD**, Lorberbaum T, Nwankwo V, Youn C, Tatonetti NP. "A Distributed Infrastructure for Drug Effect Discovery using Spontaneous Reporting Systems." (In review).
- 8. **Romano JD**, Polubriaginof F, Yahi A, Vawdrey D, Tatonetti NP. "Open Data: A review of current methods used for discovery in biomedical research." (In review).
- 9. Romano JD, Nwankwo V, Tatonetti NP. "VenomKB v2.0." (In review).

## CONFERENCE PRESENTATIONS and INVITED TALKS

- VenomSeq—A platform for drug discovery from animal venoms using differential gene expression, AMIA 2018 Informatics Summit, American Medical Informatics Association; San Francisco, CA
- 2018 Computational Resources for Personalized Genomics: High Performance Clusters and Bioinformatics Resources for Analysis and Functional Interpretation of Next-Generation Sequencing Data (tutorial session), American Medical Informatics Association; San Francisco, CA
- 2017 Deep recurrent neural networks identify transgender patients, 2017 AMIA Annual Symposium, American Medical Informatics Association; Washington, DC
- 2017 **High-sensitivity Stroke Phenotyping using Recurrent Neural Networks**, AMIA Joint Summits on Translational Science 2017, American Medical Informatics Association; San Francisco, CA
- Building a Centralized Resource for Computational Venom Research, National Library of Medicine Informatics Training Conference 2016, The Ohio State University College of Medicine; Columbus, OH
- Venom Peptides as Therapeutic Agents: Can we use Phylogenetics to Inform Drug Discovery?, 13th Annual Rocky Mountain Bioinformatics Conference, International Society for Computational Biology; Aspen, CO
- 2015 Data-Driven Clinical Research Generalizability Assessment and Improvement, 8th Annual Mid-Atlantic Healthcare Informatics Symposium, Department of Biomedical and Health Informatics, The Children's Hospital of Philadelphia; Philadelpha, PA
- 2015 Computational Prediction of Therapeutic Uses for Venom Compounds, Research Seminar Series in Biomedical Informatics, Columbia University Department of Biomedical Informatics; New York, NY
- 2015 Observational Health Data Sciences and Informatics (OHDSI), Innovation and Application @ Columbia, Columbia University Data Science Institute; New York, NY

#### FELLOWSHIPS and GRANT SUPPORT

2015-2017 **NLM T15 Training Fellow**, National Library of Medicine

# TEACHING EXPERIENCE

- Spring 2018 BINF G4003 Methods II: Symbolic Methods (Guest Lecture—1 lecture). Columbia University; New York, NY.
- Spring 2017 BINF G4003 Methods II: Symbolic Methods (Teaching Assistant). Columbia University; New York, NY.
- Fall 2016 BINF G4006 Translational Bioinformatics (Guest Lecturer—2 lectures). Columbia University; New York, NY.
- Fall 2016 BINF G4000 Acculturation to Programming and Statistics (Teaching Assistant).
- Columbia University; New York, NY.
- Spring 2014 Linux and UNIX Operating Systems.

  UVM Enterprise Technology Services, UVM TechTeam; Burlington, VT.

Spring 2012 MMG 104 — Introduction to Recombinant DNA Technology (Teaching Assistant).

University of Vermont; Burlington, VT.

Fall 2011 English Language Teaching Assistant.

Nihon University College of International Relations; Mishima, Japan.

#### **HONORS** and **AWARDS**

2010-2014	Honors Scholar, University of Vermont Honors College
2010 – 2014	Vermont Scholar, University of Vermont
2009	AP Scholar with Honors, College Board
2009	National Merit Scholar, National Merit Scholarship Corporation
2009	Eagle Scout, Boy Scouts of America

### **BOARD and COMMITTEE APPOINTMENTS**

2018-present	Chair-Elect, AMIA Student Working Group, American Medical Informatics Association
2015-present	Communications and Social Media Coordinator, AMIA Genomics and Translational
	Bioinformatics Working Group, American Medical Informatics Association
2015	Assistant Editor, Editorial Committee, 15th World Congress on Health and Biomedical Infor-
	matics; São Paulo, Brazil
2014	Committee Member, "Year In Review: Informatics in the Media 2014" Student Review Com-
	mittee American Medical Informatics Association Annual Symposium: Washington D.C.

## REVIEW ACTIVITIES

AMIA Annual Symposium

AMIA Joint Summits on Translational Science

IEEE International Conference on Healthcare Informatics

Nature Scientific Reports

Pacific Symposium on Biocomputing

 $World\ Congress\ on\ Health\ and\ Biomedical\ Informatics$ 

## PROFESSIONAL MEMBERSHIPS and AFFILIATIONS

2017–present	American Society of Clinical Pharmacology and Therapeutics
2016-present	International Society on Toxinology
2014-present	International Society for Computational Biology
2014-present	New York Academy of Sciences
2014-present	American Association for the Advancement of Science
2012-present	American Medical Informatics Association (AMIA)
2011 – 2014	Alpha Zeta academic honors fraternity
2011 – 2014	UVM Biochemistry Society

#### WORK EXPERIENCE

2010–2014 Level IV Lead Technician, University of Vermont Enterprise Technology Services; Burlington, VT.

Student management position at a walk-in software support clinic and telephone computing helpline serving all students, faculty, and staff at the University of Vermont.

- 2007–2014 Front Desk Manager, Cedarbrook at Killington / North Star Lodge; Killington, VT.

  Oversaw all aspects of customer service at a timeshare resort. Includes answering phone calls, checking in/out guests, auditing the resort, and overseeing property maintenance.
- Summer 2010 **Desktop Support Technician**, Central Vermont Public Service; Rutland, VT. Summer internship providing software and hardware support for all CVPS employees, as well as maintaining and troubleshooting the company's enterprise-level network.
- 2009–2010 **IT Technician**, *University of Rochester ITS*; Rochester, NY.

  Provided desktop support to faculty, staff, and students of the University of Rochester. Additionally, managed computing lab facilities available to University affiliates.