#### 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 13<sup>th</sup> August, 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

# PEER-REVIEWED PUBLICATIONS

- 1. **Romano J.D.**, Tharp W.G., & Sarkar I.N. (2014). Exploring Complex Disease Gene Relationships Using Simultaneous Analysis. *UVM Honors College Theses*, 35.
- 2. **Romano J.D.**, Tharp W.G., & Sarkar I.N. Adapting Simultaneous Analysis Phylogenomic Techniques to Study Complex Disease Gene Relationships. (2014). *Journal of Biomedical Informatics*, 54, 10-38.
- 3. **Romano J.D.**, & Tatonetti N.P. (2015). VenomKB, a new knowledge base for facilitating the validation of putative venom therapies. *Scientific Data*, 2, 150065.
- 4. Boland M.R., Jacunski A., Lorberbaum T., **Romano J.D.**, Moskovitch R., & Tatonetti N.P. (2016). Systems Biology Approaches for Identifying Adverse Drug Reactions and Elucidating Their Underlying Biological Mechanisms. *WIREs Systems Biology and Medicine*, 8(2), 104-122.

- 5. **Romano J.D.**, & Tatonetti N.P. (2016). Using a Novel Ontology to Inform the Discovery of Therapeutic Peptides from Animal Venoms. *AMIA Summits on Translational Science Proceedings*, 2016, 209-218. (Second place; TBI Student Paper Competition)
- 6. Vanguri R.S., **Romano J.D.**, Lorberbaum T., Nwankwo V., Youn C., & Tatonetti N.P. (2017). nSides: An interactive drug—side effect gateway. *Gateways 2017*.

# PREPRINTS (NON-REFEREED)

1. **Romano J.D.**, Nwankwo V., & Tatonetti N.P. (2018). VenomKB v2.0: A knowledge repository for computational toxinology. *bioRxiv*, 295204.

### MANUSCRIPTS IN REVIEW

1. **Romano J.D.**, Bernauer M., McGrath S.P., Nagar S.D., & Freimuth R.R. A Decade of Translational Bioinformatics: A Retrospective Analysis of "Year-in-Review" Presentations.

#### MANUSCRIPTS IN PREPARATION

- 1. **Romano J.D.**, Polubriaginof F., Yahi A., Vawdrey D., & Tatonetti N.P. Open Data: A review of methods and impact in biomedical science.
- 2. **Romano J.D.**, Li H., Realubit R., Karan C., & Tatonetti N.P. VenomSeq: Discovery of drug candidates from animal venoms.
- 3. **Romano J.D.**, Saint-Mézard M., & Tatonetti N.P. An ontology-aware semantic API for computational toxinology.
- 4. **Romano J.D.**, & Tatonetti N.P. Data-Driven Natural Product Drug Discovery: A review of current informatics techniques, knowledge gaps, and paths forward.

# CONFERENCE PRESENTATIONS and INVITED TALKS

- 2018 Engineering Innovative Informatics Solutions to Support Venom-Based Drug Discovery, 2018 GRC on Venom Evolution, Function and Biomedical Applications, West Dover, VT.
- VenomSeq—A platform for drug discovery from animal venoms using differential gene expression, AMIA 2018 Informatics Summit, 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), AMIA 2018 Informatics Summit, San Francisco, CA.
- 2017 Deep recurrent neural networks identify transgender patients, 2017 AMIA Annual Symposium, Washington, DC.
- 2017 **High-sensitivity Stroke Phenotyping using Recurrent Neural Networks**, AMIA Joint Summits on Translational Science 2017, San Francisco, CA.
- 2016 Building a Centralized Resource for Computational Venom Research, National Library of Medicine Informatics Training Conference 2016, Columbus, OH.
- Venom Peptides as Therapeutic Agents: Can we use Phylogenetics to Inform Drug Discovery?, 13th Annual Rocky Mountain Bioinformatics Conference, Aspen, CO.
- 2015 Data-Driven Clinical Research Generalizability Assessment and Improvement, 8th Annual Mid-Atlantic Healthcare Informatics Symposium, 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.

### TEACHING EXPERIENCE

Summer 2018	World Scholars $SAT+$	$summer\ program$	(Guest Lecturer).
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Yale Divinity School; New Haven, CT.

Spring 2018 BINF G4003 — Methods II: Symbolic Methods (Guest Lecturer).

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.

# PROFESSIONAL SERVICE

2019 Scientific Program Committee Member, AMIA Informatics Summit.

2018-present Chair-Elect, AMIA Student Working Group, American Medical Informatics Association.

2016, 2018 Committee Member, Translational Bioinformatics Year-in-Review committee, AMIA Trans-

lational Bioinformatics Summit.

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

Informatics; 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.

# GRANT SUPPORT

2015–2017 Training in Biomedical Informatics at Columbia University. T15 LM007079 National

Library of Medicine. (PI: Hripcsak).

2014-present Drug Effect Discovery Through Data Mining and Integrative Chemical Biology.

R01 GM107145 National Institute for General Medical Sciences. (PI: Tatonetti).

## HONORS, AWARDS, and FELLOWSHIPS

2015–2017 Biomedical Informatics Research Fellow, US National Library of Medicine

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

## REVIEW ACTIVITIES (JOURNALS and CONFERENCE PROCEEDINGS)

AMIA Annual Symposium

AMIA Joint Summits on Translational Science

IEEE International Conference on Healthcare Informatics

Nature Communications
Nature Scientific Reports

Pacific Symposium on Biocomputing

Scientific Data

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

#### PRIOR WORK EXPERIENCE

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

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.