

Marc G Chevrette

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🎓 Education

University of Wisconsin-Madison

Doctor of Philosophy (PhD) – Genetics

Madison, WI

04/2019

Master of Science (MSc) – Genetics

10/2017

• **Advisor:** Cameron R Currie, PhD

• **Research Focus:** Evolution of Microbial Metabolic Diversity, Chemically-mediated Microbiome Interactions, & Antibiotic Discovery

• **Thesis:** Evolution of Antibiotic Biosynthesis in Actinobacteria: A Framework for Drug Discovery

Institut Pasteur

Annecy, France

Certificate – International Course on Antibiotics and Resistance

11/2017

Harvard University Extension

Cambridge, MA

Master of Liberal Arts (ALM) – Biotechnology (Bioengineering & Nanotechnology)

03/2015

• **Advisor:** Tomás Maira-Litrán, PharmD, PhD

• **Research Focus:** Genome-wide Experimental & Computational Characterization of *In Vivo* Fitness Factors in Bacterial Infections

• **Thesis:** Transposon-Directed Insertion Site Sequencing for Determination of Fitness Factors in Pulmonary Infection by *A. baumannii*.

Rensselaer Polytechnic Institute

Troy, NY

Bachelor of Science (BSc) – Molecular Biology & Bioinformatics

12/2010

🏢 Experience

Wisconsin Institute for Discovery & University of Wisconsin-Madison

Madison, WI

Postdoctoral Associate - Department of Plant Pathology - Advisor: Jo Handelsman, PhD

06/2019–present

WiSolve Consulting

Madison, WI

Co-Founder & Senior Consultant

03/2016–present

Director of Technology

01/2018–09/2019

University of Wisconsin-Madison

Madison, WI

Postdoctoral Associate - Department of Bacteriology - Advisor: Cameron R Currie, PhD

04/2019–05/2019

PhD Candidate - Department of Genetics - Advisor: Cameron R Currie, PhD

08/2015–04/2019

Harvard & Georgetown Universities

Cambridge, MA

Lead Computational Biologist - Departments of Biology and Planetary Science - Advisor: Sarah S Johnson, PhD

10/2013–10/2015

Warp Drive Bio

Cambridge, MA

Head of Experimental Genomics

04/2013–08/2015

Brigham & Women's Hospital

Boston, MA

Research Assistant, Microbiology & Computational Biology - Advisor: Tomás Maira-Litrán, PharmD, PhD

03/2013–08/2015

Broad Institute of MIT & Harvard

Cambridge, MA

Research Associate II, Molecular Biology Process Development

01/2011–03/2013

Rensselaer Polytechnic Institute

Troy, NY

Research Associate, Molecular Genetics - Advisor: Eric Rutledge, PhD

05/2010–12/2010

BCR Biotech

Jamestown, RI

Research Assistant, Microbiology

09/2009–12/2009

📄 Publications, Talks, & Abstracts

Peer-reviewed Publications

*contributed equally

2019 EJ Caldera*, **MG Chevrette***, BR McDonald, CR Currie. "Local adaptation of bacterial symbionts within a geographic mosaic of antibiotic coevolution." *Applied & Environmental Microbiology*, In press.

2019 **MG Chevrette**, C Carlos-Shanley, KB Louie, BP Bowen, TR Northen, CR Currie. "Taxonomic and metabolic incongruence in the ancient genus *Streptomyces*." *Frontiers in Microbiology*. [10.3389/fmicb.2019.02170](https://doi.org/10.3389/fmicb.2019.02170)

- 2019 K Throckmorton*, V Vinnik*, R Chowdhury, TB Cook, **MG Chevette**, CD Maranas, BF Pflieger, MG Thomas. "Directed evolution of an adenylation domain specificity code." *ACS Chemical Biology*. [10.1021/acscchembio.9b00532](https://doi.org/10.1021/acscchembio.9b00532)
- 2019 J Yan, **MG Chevette**, D Braun, MK Harper, CR Currie, TS Bugni. "Madurastatin D1 and D2, oxazoline containing siderophores isolated from an *Actinomadura* sp." *Organic Letters*. [10.1021/acs.orglett.9b02159](https://doi.org/10.1021/acs.orglett.9b02159)
- 2019 **MG Chevette***, JR Bratburd*, CR Currie, RM Stubbendieck. "Experimental microbiomes: models not to scale." *mSystems*. [10.1128/mSystems.00175-19](https://doi.org/10.1128/mSystems.00175-19)
- 2019 **MG Chevette**, CM Carlson, HE Ortega, C Thomas, GE Ananiev, KJ Barns, AJ Book, J Cagnazzo, C Carlos, W Flanigan, KJ Grubbs, HA Horn, FM Hoffmann, JL Klassen, JJ Knack, GR Lewin, BR McDonald, L Muller, WGP Melo, AA Pinto-Tomás, A Schmitz, E Wendt-Pienkowski, S Wildman, M Zhao, F Zhang, TS Bugni, DR Andes, MT Pupo, CR Currie. "The antimicrobial potential of *Streptomyces* from insect microbiomes." *Nature Communications*. [10.1038/s41467-019-08438-0](https://doi.org/10.1038/s41467-019-08438-0)
[Highlighted by NPR]
- 2019 RM Stubbendieck, DS May, **MG Chevette**, MI Temkin, E Wendt-Pienkowski, J Cagnazzo, CM Carlson, JE Gern, CR Currie. "Competition among nasal bacteria suggests a role for siderophore-mediated interactions in shaping the human nasal microbiota." *Applied & Environmental Microbiology*. [10.1128/AEM.02406-18](https://doi.org/10.1128/AEM.02406-18)
- 2019 N Liu,* H Li,* **MG Chevette**, L Zhang, L Cao, H Zhou, X Zhou, Z Zhou, PB Pope, CR Currie, Y Huang, Q Wang. "Functional metagenomics reveals polysaccharide-degrading gene clusters and cellobiose utilization pathways in gut microbiota of a wood-feeding termite." *ISME Journal*. [10.1038/s41396-018-0255-1](https://doi.org/10.1038/s41396-018-0255-1)
- 2019 **MG Chevette**, CR Currie. "Emerging evolutionary paradigms in antibiotic discovery." *Journal of Industrial Microbiology & Biotechnology*. [10.1007/s10295-018-2085-6](https://doi.org/10.1007/s10295-018-2085-6)
- 2017 N Adnani, **MG Chevette**, SN Adibhatla, F Zhang, Q Yu, D Braun, J Nelson, SW Simpkins, BR McDonald, CL Myers, J Piotrowski, C Thompson, CR Currie, L Li, SR Rajsiki, TS Bugni. "Co-culture of marine invertebrate-associated bacteria and interdisciplinary technologies enable biosynthesis and discovery of a new antibiotic, keyicin." *ACS Chemical Biology*. [10.1021/acscchembio.7b00688](https://doi.org/10.1021/acscchembio.7b00688)
[Highlighted by Nature]
- 2017 AF Sanchez-Larrayoz, NM Elshamy, **MG Chevette**, Y Fu, P Giunta, RG Spallanzani, K Ravi, GB Pier, S Lory, T Maira-Litrán. "Complexity of complement-resistance factors expressed by *Acinetobacter baumannii* needed for survival in human serum." *Journal of Immunology*. [10.4049/jimmunol.1700877](https://doi.org/10.4049/jimmunol.1700877)
- 2017 **MG Chevette**, F Aicheler, O Kohlbacher, CR Currie, MH Medema. "SANDPUMA: Ensemble predictions of nonribosomal peptide chemistry reveals biosynthetic diversity across Actinobacteria." *Bioinformatics*. [10.1093/bioinformatics/btx400](https://doi.org/10.1093/bioinformatics/btx400)
- 2017 IJ Miller, **MG Chevette**, JC Kwan. (2017). "Interpreting microbial biosynthesis in the genomic age: Biological and practical considerations." *Marine Drugs*. [10.3390/md15060165](https://doi.org/10.3390/md15060165)
[Cover Image for Issue 6, Volume 15 in June 2017]
- 2017 K Blin, T Wolf, **MG Chevette**, X Lu, CJ Schwalen, SA Kautsar, HG Suarez Duran, ELC de los Santos, HUK Kim, M Nave, JS Dickschat, DA Mitchell, E Shelest, R Breitling, E Takano, SY Lee, T Weber, MH Medema. "antiSMASH 4.0 - Improvements in chemistry prediction and gene cluster boundary identification." *Nucleic Acids Research*. [10.1093/nar/gkx319](https://doi.org/10.1093/nar/gkx319)
- 2016 GR Lewin, C Carlos, **MG Chevette**, HA Horn, BR McDonald, RJ Stankey, BG Fox, CR Currie. "Ecology and evolution of Actinobacteria and their bioenergy applications." *Annual Review of Microbiology*. [10.1146/annurev-micro-102215-095748](https://doi.org/10.1146/annurev-micro-102215-095748)
- 2015 SS Johnson, **MG Chevette**, BL Ehlmann, KC Benison. "Insights from the metagenome of an acid salt lake: The role of biology in an extreme depositional environment." *PLOS ONE*. [10.1371/journal.pone.0122869](https://doi.org/10.1371/journal.pone.0122869)

Book Chapters

- 2019 **MG Chevette**, PA Hoskisson, F Barona-Gómez. "Enzyme evolution in secondary metabolism." *Comprehensive Natural Products III: Chemistry and Biology*. In press.

Preprints, Submitted, and Under Review

- BR McDonald, **MG Chevette**, JL Klassen, HA Horn, EJ Caldera, E Wendt-Pienkowski, MJ Cafaro, AC Ruzzini, EB Van Arnam, GM Weinstock, NM Gerardo, M Poulsen, G Suen, J Clardy, CR Currie. "Biogeography and microscale diversity shape the biosynthetic potential of fungus-growing ant-associated *Pseudonocardia*." [Preprint available](#)

Invited Talks

- 2019 "Drugs from bugs of bugs: a novel source for antimicrobials." American Society for Microbiology Microbe. San Francisco, CA. Jun 21, 2019.
- 2019 "Mining microbiomes for antimicrobials." Synthetic Biology for Natural Products Conference. Puerto Vallarta, Mexico. Jun 02, 2019.
- 2018 "Host-associated microbes as a source of new antimicrobials." Natural Product Discovery & Development in the Genomic Era, Society for Industrial Microbiology & Biotechnology. Clearwater Beach, FL. Jan 22, 2018.

- 2017 “Natural natural products: Leveraging chemical ecology in the search for new drugs.” Evolution Seminar Series, JF Crow Institute for the Study of Evolution. Madison, WI. Oct 26, 2017.
- 2017 “Computational insights into the diverse nonribosomal peptide chemistry of Actinobacteria.” Synthetic Biology for Natural Products Conference. Cancun, Mexico. Mar 6, 2017.
[Highlighted in [ACS Synthetic Biology](#)]
- 2016 “Darwinian drug discovery: Chemical ecology at fine and coarse evolutionary scales.” International Chemical Biology Society Annual Conference. Madison, WI. Oct 24, 2016.
[Highlighted in [ACS Chemical Biology](#)]

Patents

- 2017 DC Gray, E Li, BR Bowman, GL Verdine, K Robison, **MG Chevette**, D Udway, PS Wang, A Li, JP Morgenstern. Compositions and methods for the production of compounds. Patent Application PCT/US2017/058800, filed Oct 27, 2017.
Publication Number [WO/2018/081590](#)

Publications, Editorial Review Only

- 2018 **MG Chevette**. “Natural products reawakened: New trends in discovery and development.” *SIMB News Magazine*, Society for Industrial Microbiology and Biotechnology.
- 2018 DR Braun, **MG Chevette**, D Acharya, CR Currie, SR Rajski, TS Bugni. “Draft genome of *Micromonospora* sp. WMMA1996, a marine sponge-associated bacterium.” *Genome Announcements*. [10.1128/genomeA.00077-18](#)
- 2018 DR Braun, **MG Chevette**, D Acharya, CR Currie, SR Rajski, K Ritchie, TS Bugni. “Complete genome of *Dietzia* sp. WMMA184, a marine coral-associated bacterium.” *Genome Announcements*. [10.1128/genomeA.01582-17](#)
- 2017 N Adnani, DR Braun, BR McDonald, **MG Chevette**, CR Currie, TS Bugni. “Draft genome of *Micromonospora* sp. WMMB-235, a marine ascidian-associated bacterium.” *Genome Announcements*. [10.1128/genomeA.01369-16](#)
- 2016 N Adnani, DR Braun, BR McDonald, **MG Chevette**, CR Currie, TS Bugni. “Complete genome sequence of *Rhodococcus* sp. strain WMMA185, a marine sponge-associated bacterium.” *Genome Announcements*. [10.1128/genomeA.01406-16](#)

Internal Seminars

- 2018 “Drugs & bugs of bugs: Insect microbiomes as a source of new antibiotics.” Genetics Colloquium, UW-Madison. Aug 8, 2018.
- 2017 “Host-microbe interactions as a source of new antimicrobials.” Highlights at the Chemistry-Biology Interface Colloquium, UW-Madison. Dec 12, 2017.
- 2017 “Genome-based natural product discovery, modular biosynthesis, & applications.” Highlights at the Chemistry-Biology Interface Colloquium, UW-Madison. Feb 2, 2017.
- 2016 “Genome assembly: Tools & analysis.” Computational Biology, Ecology, & Evolution (ComBEE), UW-Madison. Apr 27, 2016.

Abstracts

- 2019 **MG Chevette**, D Acharya, A Hurley, M Beebe, M Garavito, S Miller, J Handelsman. “Tiny Earth Chemistry Hub: From soil to antibiotics.” Presented at: Tiny Earth Symposium; Madison, WI; Jul 10, 2019.
- 2019 **K Throckmorton**, V Vinnik, TB Cook, R Chowdhury, **MG Chevette**, CD Maranas, BF Pfleger, MG Thomas. “Directed evolution of an adenylation domain specificity code.” Presented at: Synthetic Biology for Natural Products Conference; Puerto Vallarta, Mexico; Jun 2, 2019.
- 2019 **CL Hansen**, **MG Chevette**, M Selvaraj, A Vasquez Echeverri, D Maldonado Perez, C Eno, J Hernandez-Ortiz, F Pelegri. “Helical supramolecular assembly of a germline specific membraneless organelle.” Presented at: Phase Separation in Biology & Disease; New York, NY; Feb 20, 2019.
- 2018 **MG Chevette**, CM Carlson, H Ortega, F Zhang, KJ Grubbs, MT Pupo, TS Bugni, DR Andes, CR Currie. “Insect-associated *Streptomyces* are a rich source of new antimicrobials.” Presented at: Beneficial Microbes; Madison, WI; Jul 9, 2018.
- 2018 **HA Horn**, E Gemperline, K Delaney, **MG Chevette**, L Li, CR Currie. “Host specificity influences chemical response in *in vivo* symbiotic interactions.” Presented at: Beneficial Microbes; Madison, WI; Jul 9, 2018.
- 2018 **BR McDonald**, **MG Chevette**, J Klassen, HA Horn, EJ Caldera, E Wendt-Pienkowski, MJ Cafaro, AC Ruzzini, EB Van Arnam, GM Weinstock, NM Gerardo, MG Poulsen, G Suen, J Clardy, CR Currie. “Biogeography and microscale diversity shapes the biosynthetic potential of fungus-growing ant associated *Pseudonocardia*.” Presented at: Beneficial Microbes; Madison, WI; Jul 9, 2018.
- 2018 **MG Chevette**, CM Carlson, H Ortega, F Zhang, KJ Grubbs, MT Pupo, TS Bugni, DR Andes, CR Currie. “Insect-associated *Streptomyces* are a rich source of new antimicrobials with activity against resistant human pathogens.” Presented at: Perlman Antibiotic Discovery and Development Symposium; Madison, WI; Apr 27, 2018.
- 2018 **DD Acharya**, IJ Miller, Y Cui, DR Braun, **MG Chevette**, M Berres, L Li, J Kwan, CR Currie, TS Bugni. “Chemical cross-talk in bacterial co-cultures affects differential gene expression and antibiotic production.” Presented at: Perlman Antibiotic Discovery and Development Symposium; Madison, WI; Apr 27, 2018.

- 2017 R Zarnowski, **MG Chevrete**, E Dominguez, DR Andes. "Modeling high-throughput proteomics into predictive metabolomics - A novel tool for studies of medical device-associated *Candida spp.* biofilm infections." Presented at: Metabolomics Circle 2017 - Bioanalytical & Omics Science, Wrocław, Poland; Nov 18, 2017.
- 2017 D Acharya, N Adnani, D Braun, IJ Miller, Q Yu, **MG Chevrete**, M Berres, CR Currie, L Li, JC Kwan, TS Bugni. "Chemical cross-talk in bacterial co-cultures affects differential gene expression and antibiotic production." Presented at: American Society for Pharmacognosy Annual Meeting, Portland, OR; Jul 30, 2017.
- 2017 AF Sanchez-Larrayoz, NM Elhosseiny, **MG Chevrete**, Y Fu, P Giunta, G Spallanzani, GB Pier, S Lory, T Maira-Litrán. "The membrane lipid asymmetry transport system plays a key role in protecting *Acinetobacter baumannii* against killing by human complement via the alternative pathway." Presented at: American Society for Microbiology Microbe, New Orleans, LA; Jun 2, 2017.
- 2017 **MG Chevrete**, CM Carlson, C Thomas, TS Bugni, DR Andes, CR Currie. "Evolutionary trends in secondary metabolism reveal insect-associated *Streptomyces* as an underexploited antibiotic resource." Presented at: Perlman Antibiotic Discovery and Development Symposium; Madison, WI; Mar 31, 2017.
- 2017 EJ Caldera, **MG Chevrete**, CR Currie. "The geographic mosaic of antibiotic coevolution in a bacterial symbiont of the fungus-farming ant *Apterostigma dentigerum*." Presented at: Perlman Antibiotic Discovery and Development Symposium; Madison, WI; Mar 31, 2017.
- 2017 J Bratburd, C Keller, E Vivas, **MG Chevrete**, F Rey, L Li, CR Currie. "The human gut microbiota metabolomic response to infection." Presented at: Perlman Antibiotic Discovery and Development Symposium; Madison, WI; Mar 31, 2017.
- 2016 **MG Chevrete**, CR Currie, MH Medema. "prediCAT: An accurate predictive method for substrate specificity of nonribosomal peptide synthetase adenylation domains." Presented at: Kenneth B. Raper Symposium on Microbial Research; Madison, WI; Sep 2, 2016.
- 2016 J Bratburd, BR McDonald, **MG Chevrete**, JL Klassen, HA Horn, CR Currie. "Comparative genomics of fungus-growing ant-associated *Pseudonocardia*." Presented at: Kenneth B. Raper Symposium on Microbial Research; Madison, WI; Sep 2, 2016.
- 2016 HA Horn, E Gemperline, **MG Chevrete**, BR McDonald, J Bratburd, E Mevers, J Clardy, L Li, CR Currie. "Mass spectrometry imaging reveals differential chemical response to pathogens in an ancient ant-microbe symbiosis." Presented at: ISME International Symposium on Microbial Ecology; Montreal, QC, Canada; Aug 21-26, 2016.
- 2016 **MG Chevrete**, CR Currie, MH Medema. "Computational predictions of substrate specificity in nonribosomal peptide synthetases through comparative adenylation domain trees." Presented at: American Society for Microbiology Microbe; Boston, MA; Jun 16-20, 2016.
- 2016 SS Johnson, ML Soni, DJ Collins, KC Benison, MR Mormile, **MG Chevrete**, BL Ehlmann. "Biosignatures in mars analog acid salt lakes." Presented at: USRA Biosignature, Preservation and Detection in Mars Analog Environments; Lake Tahoe, Nevada; May 16-19, 2016.
- 2016 **MG Chevrete**, C Carlson, C Thomas, TS Bugni, CR Currie. "Multifaceted antibiotic profiling across Actinomycete chemical ecology." Presented at: Perlman Antibiotic Discovery and Development Symposium; Madison, WI; Apr 29, 2016.
- 2016 N Adnani, S Adibhatla, E Vazquez-Rivera, GA Ellis, D Braun, **MG Chevrete**, BR McDonald, C Thompson, JS Piotrowski, Q Yu, L Li, CR Currie, TS Bugni. "Driving production of novel natural products through marine microbial interspecies interactions." Presented at: Gordon Marine Natural Products; Ventura, CA; Mar 6-11, 2016.
- 2015 **MG Chevrete**, DW Udway, CR Currie, SS Johnson. "Functional classification and secondary metabolism of an extreme metagenome." Presented at: Kenneth B. Raper Symposium on Microbial Research; Madison, WI; Sep 1, 2015.
- 2015 **MG Chevrete**, BL Ehlmann, KC Benison, SS Johnson. "Microbial diversity and biosynthetic potential of an extreme sediment metagenome." Presented at: Gordon Applied and Environmental Microbiology; South Hadley, MA; Jul 12-17, 2015.
- 2015 **MG Chevrete**, M Vinacur, T Maira-Litrán. "Transposon-directed insertion site sequencing reveals *in vivo* fitness factors in *Acinetobacter baumannii* lung infections." Presented at: Boston Bacterial Meeting; Cambridge, MA; Jun 18-19, 2015.
- 2014 DW Udway, K Robison, **MG Chevrete**, GL Verdine. "Lessons from long read assembly of 100+ Actinomycete genomes." Presented at: Gordon Marine Natural Products; Ventura, CA; Mar 2-7, 2014.
- 2014 K Robison, DW Udway, **MG Chevrete**, GL Verdine. "Long read assembly of >100 Actinomycete genomes." Presented at: Advances in Genome Biology & Technology; Marco Island, FL; Feb 12-15, 2014.
- 2013 S Young, S Steelman, R Daza, **MG Chevrete**, R Lintner, S Gnerre, A Berlin, B Walker, C Nusbaum, R Nicol. "Generation of high-quality draft assemblies with a single sequencing library." Presented at: Sequencing, Finishing, Analysis in the Future; Santa Fe, NM; May 29-31, 2013.
- 2012 S Steelman, R Daza, **MG Chevrete**, P Kompella, P Trang, T Surabian, R Lintner, CZ Zhang, J Jung, M Meyerson, C Nusbaum, R Nicol. "Automated low input mate-pair library construction for Illumina sequencing." Presented at: Advances in Genome Biology & Technology; Marco Island, FL; Feb 15-18, 2012.
- 2011 S Steelman, R Daza, **MG Chevrete**, P Kompella, P Trang, T Surabian, R Lintner, R Nicol. "Microbial mate-pair library construction for de novo detection of structural rearrangements." Presented at: Broad Institute Symposium; Boston, MA; Nov 7-8, 2011.

Honors & Awards

Wisconsin Scientific Teaching Design Institute Fellow	UW-Madison	2019–present
Schlimgen Award for Outstanding Scholarship in Doctoral Studies in Genetics	UW-Madison	2019
Ira L Baldwin Distinguished Predoc. Fellowship for Excellence in Research	Bacteriology, UW-Madison	2018–2019
Chemistry-Biology Interface Predoctoral Fellowship	National Institutes of Health, NIGMS	2016–2018
Passed with Distinction	Preliminary Examination A - Genetics, UW-Madison	07/2017
Issue Cover	Marine Drugs: Connecting Marine Microbial Natural Products to Biosynthetic Pathways	06/2017
Bacteriology Departmental Travel Grant	University of Wisconsin-Madison	2016
Vilas Travel Grant	University of Wisconsin-Madison	2016
Dean's Academic Achievement Award	Harvard University Extension	03/2015
Finalist, Core Value Award: "Courageous: Uncompromising Science"	Warp Drive Bio	2014
Finalist, Core Value Award: "Unbounded: Reimagining the Possible"	Warp Drive Bio	2014
Featured Scientific Researcher – "Who is Broad?"	Broad Institute of MIT & Harvard	01/2012
Rensselaer Alumni Scholarship	Rensselaer Polytechnic Institute	2004–2008
Sal H. Alfiero Scholarship	Rensselaer Polytechnic Institute	2004–2008
Rhode Island State Scholarship	Rensselaer Polytechnic Institute	2004–2008

Service & Outreach

Ad hoc Reviewer	Critical Reviews in Microbiology, World Journal of Microbiology & Biotechnology, Microbial Genetics, Frontiers in Microbiology, FEMS Microbiology Letters, Microbial Cell Factories	
On-air Guest	Perpetual Notion Machine, WORT FM, Link	02/2019
Evolution Coordinating Committee	JF Crow Institute for the Study of Evolution, UW-Madison	01/2017–present
Mentor	Google Summer of Code – antiSMASH, Open Bioinformatics Foundation	03/2016–09/2017
Co-chair	Computational Biology, Ecology, & Evolution (ComBEE), UW-Madison	01/2016–08/2018
Co-organizer	Discovery Niche, Wisconsin Institutes for Discovery	10/2015–11/2015
Volunteer	Wisconsin Science Festival	10/2015
Open Genomics Adviser	Revive & Restore, Long Now Foundation	04/2014–10/2015
Environmental, Health, and Safety Representative	Broad Institute of MIT & Harvard	01/2011–03/2013

Teaching Experience

Microbiology 450: Diversity, Ecology, & Evolution of Microorganisms, UW-Madison	Lecturer	Fa 2018
Programming in R, ComBEE, UW-Madison	Group Leader, Lecturer	Sp 2016
Genetics 468: General Genetics II, UW-Madison	Lecturer, TA	Sp 2016
Microbiology 450: Diversity, Ecology, & Evolution of Microorganisms, UW-Madison	Lecturer, TA	Fa 2016

Professional Societies & Groups

International Chemical Biology Society	2016–present
Natural Products Discovery and Bioengineering Network	2016–present
American Society for Microbiology	2015–present
Computational Biology, Ecology, & Evolution (ComBEE) – UW-Madison	2015–present
JF Crow Institute for the Study of Evolution	2015–present
Society for Industrial Microbiology and Biotechnology	2014–present
Laboratory Robotics Interest Group – New England Chapter	2011–2015