

Graduate Student Evolutionary genomics jacob.steenwyk@vanderbilt.edu jlsteenwyk.github.io

EDUCATION

Present	Graduate Student Biological Sciences Advisor: Antonis Rokas GPA: 3.97	Vanderbilt University
2016	M.S. Biochemistry and Molecular Biology Advisor: John G. Gibbons GPA: 3.98	Clark University
2015	B.A. Biochemistry and Molecular Biology Advisor: Denis Larochelle Cumulative GPA: 3.84 Science GPA: 3.84	Clark University

AWARDS

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2018	GENETICS Peer Review Training Program, Genetics Society of America
2018	Best poster award, Gordon Research Seminar, Holderness, NH
2018	Best poster award, Gordon Research Conference, Holderness, NH
2018	Best poster award, Department of Biological Sciences, Vanderbilt
	University
2018	T-shirt design contest winner, Department of Biological Sciences,
	Vanderbilt University
2017	Graduate student travel grant, Vanderbilt University
2016	Graduate student council travel awards, Clark University
2015	Summa cum laude, Clark University
2014	Summer research scholar, Bridging the gaps, University of Southern
	California Keck School of Medicine
2013	Global environmental microbiology scholar, Center for dark energy
	biosphere investigations, University of Southern California
2011	Jonas Clark Scholar, Clark University

RESEARCH INTERESTS

- DNA damage and repair
- Gene and genome evolution
- Evolution and function of technologically and medically significant fungi
- Phylogenetics of phylogenomics of fungi and insects

PUBLICATIONS

+ represents equal contributors

Preprint: Mead M.E., S.L. Knowles, H.A. Raja, S. R. Beattie, C.H. Kowalski, **J.L. Steenwyk**, L.P. Silva, J. Chiaratto, L.N.A. Ries, G.G. Goldman, R.A. Cramer, N.H. Oberlies, & A. Rokas (2018). Characterizing the pathogenic, genomic, and chemical traits of *Aspergillus fischeri*, the closest sequenced relative of the major human fungal pathogen *Aspergillus fumigatus*. bioRxiv: doi: 10.1101/430728

Preprint: **Steenwyk, J.L.**, X.-X. Shen, A.L. Lind, G.G. Goldman, & A. Rokas (2018). A robust phylogenomic timetree for biotechnologically and medically important fungi in the genera *Aspergillus* and *Penicillium*. bioRxiv: doi: 10.1101/370429

Preprint: Eidem, H.R., **J.L. Steenwyk**, J. Wisecaver, J.A. Capra, P. Abbot, & A. Rokas (2018). integRATE: a desirability-based data integration framework for the prioritization of candidate genes across heterogeneous 'omics and its application to preterm birth. bioRxiv: doi: 10.1101/302612

Preprint: **Steenwyk, J.**, J. St. Denis, J. Dresch, D. Larochelle, & R. Drewell (2017). Whole genome bisulfite sequencing reveals a sparse, but robust pattern of DNA methylation in the *Dictyostelium discoideum* genome. bioRxiv. doi: 10.1101/166033

- (5) Shen, X.-X.⁺, D.A. Opulente⁺, J. Kominek⁺, X. Zhou⁺, **J.L. Steenwyk**, K.V. Buh, M.A.B. Haase, J.H. Wisecaver, M. Wang, D.T. Doering, J.T. Boudouris, R.M. Schneider, Q.K. Langdon, M. Ohkuma, R. Endoh, M. Takashima, R. Manabe, N. Čadež, D. Libkind, C.A. Rosa, J. DeVirgilio, A.B. Hulfachor, M. Groenewald, C.P. Kurtzman, C.T. Hittinger & A. Rokas (2018). Tempo and mode of genome evolution in the budding yeast subphylum. Cell. doi: 10.1016/j.cell.2018.10.023
- (4) Segal, E.S., V. Gritsenko, A. Levitan, B. Yadav, N. Dror, J.L. Steenwyk, Y. Silberberg, K. Mielich, A. Rokas, N.A.R. Gow, R. Kunze, R. Sharan, & J. Berman (2018). Gene Essentiality Analyzed by In Vivo Transposon Mutagenesis and Machine Learning in a Stable Haploid Isolate of *Candida albicans*. mBio. doi: 10.1128/mBio.02048-18
- (3) **Steenwyk, J.L.** & A. Rokas (2018). Copy number variation in fungi and its implications for wine yeast genetic diversity and adaptation. Frontiers in Microbiology. doi: 10.3389/fmicb.2018.00288
- (2) **Steenwyk, J.** & A. Rokas (2017). Extensive Copy Number Variation in Fermentation-Related Genes among *Saccharomyces cerevisiae* Wine Strains. G3: Genes | Genomes | Genetics. doi: 10.1534/g3.117.040105
- (1) **Steenwyk J.L.**, J.S. Soghigian, J.R. Perfect, & J.G. Gibbons (2016). Copy number variation contributes to cryptic genetic variation in outbreak lineages of *Cryptococcus gattii* from the North American Pacific Northwest. BMC Genomics. doi: 10.1186/s12864-016-3044-0

INVITED TALKS

2019	The Phylogenomics and Evolution Group of North Carolina State
	University, Raleigh, NC (scheduled)
2015	TedXClarkUniversity, Clark University, Worcester, MA

CONTRIBUTED TALKS

2019	Science club at the library, Nashville Public Library, Nashville, TN
	(scheduled)
2018	Nashville Science Club, Jackalope Brewing Co., Nashville, TN
2017	Mycological Society of America, Univ. of Georgia, Athens, GA
2016	Mycological Society of America, Univ. of California Berkeley, Berkeley,
	CA
2016	Graduate Student Multidisciplinary Conference, Clark University,
	Worcester, MA

WORKSHOP TEACHING

Image approach, design and processing, ArtLab, Vanderbilt University, Nashville, TN (scheduled)

POSTER PRESENTATIONS

Cellular and Molecular Fungal Biology, Gordon Research Conference,
Holderness, New Hampshire
Cellular and Molecular Fungal Biology, Gordon Research Seminar,
Holderness, New Hampshire
Department of Biological Sciences Annual Retreat, Vanderbilt University,
Nashville, TN
Exploring the intersection between Art and Science, ArtLab, Vanderbilt
University, Nashville, TN
Bumpus Symposium, Clark University, Worcester, MA
Traina Scholars Presentation, Clark University, Worcester, MA
Summer Research Presentation, Clark University, Worcester, MA
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RESEARCH EXPERIENCE

2016-Pres.	Antonis Rokas Lab at Vanderbilt University, Nashville, TN.	
2015-2016	John Gibbons Lab at Clark University, Worcester, MA. Investigation of	
	copy number variation in the human pathogen, Cryptococcus gattii.	
2015-2016	Robert Drewell Lab at Clark University, Worcester, MA. Investigation of	
	genome-wide methylation patterns in Dictyostelium discoideum.	
2014	Ite A. Laird-Offringa Lab at University of Southern California, Los Angeles,	
	CA. Bridging the Gaps Summer Scholar. Project aim was to map the	
	autoimmune triggering epitope of <i>ELAVL4</i> in small cell lung cancer.	
2013	John Heidelberg and Eric Webb Labs at University of Southern California,	
	Los Angeles, CA. Global Environmental Microbiology Summer Scholar.	
	Investigation of fresh and marine water microbial diversity.	

TEACHING EXPERIENCE

2017-Pres.	Teaching Assistant, Introductory Biology Lab, Vanderbilt University, Nashville, TN
2016	Teaching Assistant, Introduction to Biostatistics, Clark University, Worcester, MA
2014-2015	Teaching Assistant, Cell Biology, Clark University, Worcester, MA
SERVICE	
2018-Pres.	Co-chair, MEGA <i>Microbe</i> , Vanderbilt Institute for Infections, Immunology and Inflammation, Nashville, TN
2018-Pres.	Vice President, Graduate Student Association, Department of Biological Sciences, Vanderbilt University, Nashville, TN
2018-Pres.	Member, American Society of Microbiology Tennessee Chapter
2017-Pres.	Educational outreach booth design and execution, MEGA <i>Microbe</i> , Nashville, TN
2017-Pres.	Communications chair, Inclusivity in Biological Sciences Association, Vanderbilt University, Nashville, TN
2017-Pres.	Member of the Dean of Graduate Student's survey quantitative analysis subgroup, Graduate Diversity and Inclusion Committee, Vanderbilt University, Nashville, TN
2017-Pres.	Judge, Middle Tennessee Science and Engineering Fair, Belmont University, Nashville, TN
2017-2018	Secretary, Graduate Student Association, Department of Biological Sciences, Vanderbilt University, Nashville, TN
2017-2018	Scientific consultant, Little Harpeth Brewing, Nashville, TN
2017	Vanderbilt Student Volunteers for Science, Volunteer Science Teacher, West End Middle School, Nashville, TN
2014-2015	Undergraduate Subcommittee for Department of Chemistry, Biochemistry and Molecular Biology Faculty Search Committee, Clark University, Worcester, MA
2014-2015	Science Education Outreach Blogger, C-DEBI Sci-Curious Blog

MANUSCRIPT REVIEWER

Systematic Biology, Molecular Genetics and Genomics, BMC Genomics, PLoS One, Young Scientists Journal, Scholarly Undergraduate Research Journal