Jacob Steenwyk

Graduate Student
Evolutionary genomics

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Dept. Biological Sciences, Vanderbilt University

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Education		
Present	Graduate Student Biological Sciences Advisor: Antonis Rokas Current 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		
2017	Graduate student travel grant, Vanderbilt Un	niversity
2016	Graduate student council travel awards, Cla	rk 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	
2011	biosphere investigations, University of Sout Jonas Clark Scholar, Clark University	nern California
Publication	s *co-first authors	
In prep:	Lind A.L.*, Steenwyk J. L.* , Ries L., Gold report, evolution, and function of clinically	
Preprint:	Steenwyk J. , St. Denis J., Dresch J., Laroch Whole genome bisulfite sequencing reveals DNA methylation in the <i>Dictyostelium disce</i> 10.1101/166033	a sparse, but robust pattern of
Preprint:	Eidem H.R., Steenwyk J. L. , Wisecaver J., A. (2018). integRATE: a desirability-based the prioritization of candidate genes across application to preterm birth	data integration framework for
Submitted:	Shen XX. and 24 others including Steenw genome evolution in the budding yeast subp	

Steenwyk J. L., Rokas A. (2018). Copy number variation in fungi and its

implications for wine yeast genetic diversity and adaptation. Frontiers in

Steenwyk J., Rokas A. (2017). Extensive Copy Number Variation in

Microbiology. doi: https://doi.org/10.3389/fmicb.2018.00288

Fermentation-Related Genes among *Saccharomyces cerevisiae* Wine Strains. G3: Genes|Genomes|Genetics. doi: 10.1534/g3.117.040105

(1) **Steenwyk J.L.**, Soghigian J.S., Perfect J.R. and Gibbons J.G. (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

Contributed Talks

2017	Mycological Society of America, Univ. of Georgia, Athens, GA
	Extensive Copy Number Variation in Fermentation-Related Genes Among
	Saccharomyces cerevisiae Wine Strains
2016	Mycological Society of America, Univ. of California Berkeley, Berkeley,
	CA
	Population structure and copy number variation in the fungal pathogen
	Cryptococcus gattii
2016	Graduate Student Multidisciplinary Conference, Clark University,
	Worcester, MA
	Population structure and copy number variation in the fungal pathogen
	Cryptococcus gattii
2015	TedXClarkUniversity, Clark University, Worcester, MA
	Instinct and Intelligence

Poster Presentations

2018	Exploring the intersection between Art and Science, ArtLab, Vanderbilt
	University, Nashville, TN
2015	Bumpus Symposium, Clark University, Worcester, MA
	Population structure and copy number variation in the fungal pathogen
	Cryptococcus gattii
2015	Traina Scholars Presentation, Clark University, Worcester, MA
	Mapping the Auto-Immune Triggering Epitope of ELAVL4
2015	Summer Research Presentation, Clark University, Worcester, MA
	Mapping the Auto-Immune Triggering Epitope of ELAVL4

Research Experience

2016-Pres.	Antonis Rokas Lab at Vanderbilt University, Nashville, TN.	
2015-16	John Gibbons Lab at Clark University, Worcester, MA. Investigation of	
	copy number variation in the human pathogen, Cryptococcus gattii.	
2015-16	Robert Drewell Lab at Clark University, Worcester, MA. Investigation of	
	genome-wide methylation patterns in <i>Dictyostelium discoideum</i> .	
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,
2016	Nashville, TN
2016	Teaching Assistant, Introduction to Biostatistics, Clark University,
20112015	Worcester, MA
2014-2015	Teaching Assistant, Cell Biology, Clark University, Worcester, MA
Service	
2018-Pres.	Co-chair, MEGAMicrobe, Vanderbilt Institute for Infections, Immunology
	and Inflammation, Nashville, TN
2017-Pres.	Communications chair, Inequalities in Biological Sciences Association,
	Vanderbilt University, Nashville, TN
2017-Pres.	Secretary, Graduate Student Association, Department of Biological
	Sciences, 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-Pres.	Scientific consultant, Little Harpeth Brewing, Nashville, TN
2017	Vanderbilt Student Volunteers for Science, Volunteer Science Teacher,
	West End Middle School, Nashville, TN
2017	Educational outreach booth design and execution, MEGAMicrobe,
	Nashville, TN
2014-15	Undergraduate Subcommittee for Department of Chemistry, Biochemistry
	and Molecular Biology Faculty Search Committee, Clark University,
	Worcester, MA
2014-15	Science Education Outreach Blogger, C-DEBI Sci-Curious Blog

Manuscript reviewer

BMC Genomics, Public Library of Sciences (PLoS) One, Young Scientists Journal, Scholarly Undergraduate Research Journal