Jacob Steenwyk

Graduate Student Evolutionary genomics Dept. Biological Sciences, Vanderbilt University

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Education

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

Awards

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esearch Seminar, Holderness, NH
esearch Conference, Holderness, NH
at of Biological Sciences, Vanderbilt
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Department of Biological Sciences,
Vanderbilt University
el awards, Clark University
versity
dging the gaps, University of Southern
edicine
iology scholar, Center for dark energy
versity of Southern California
niversity

Publications

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

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

In review: Shen X.-X. and 24 others including **Steenwyk J.L.** Tempo and mode of genome evolution in the budding yeast subphylum

- (3) **Steenwyk J.L.**, Rokas A. (2018). Copy number variation in fungi and its implications for wine yeast genetic diversity and adaptation. Frontiers in Microbiology. doi: https://doi.org/10.3389/fmicb.2018.00288
- (2) **Steenwyk J.**, Rokas A. (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.**, 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	Cellular and Molecular Fungal Biology, Gordon Research Conference,
	Holderness, New Hampshire
2018	Cellular and Molecular Fungal Biology, Gordon Research Seminar,
	Holderness, New Hampshire
2018	Department of Biological Sciences Annual Retreat, Vanderbilt University,
	Nashville, TN
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 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

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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	

- **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
- Vanderbilt Student Volunteers for Science, Volunteer Science Teacher, West End Middle School, Nashville, TN
- **2017** Educational outreach booth design and execution, MEGA*Microbe*, 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

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