

# Jacob L. Steenwyk

Berkeley Science Fellow & Howard Hughes Medical Institute Awardee of the Life Sciences Research Foundation. Howard Hughes Medical Institute and University of California, Berkeley. jlsteenwyk@berkeley.edu www.jlsteenwyk.com

## **PROFESSIONAL POSITIONS**

Since 2023 Howard Hughes Medical Institute Awardee, Life Sciences Research Foundation

Since 2022 Postdoctoral Associate, Laboratory of Dr. King, University of California, Berkeley

Since 2022 Berkeley Science Fellow, University of California, Berkeley

# FIVE HIGHLIGHTED PUBLICATIONS (Trainees and I are in **bold** font.)

- 1. **Steenwyk, J.L.**, Y. Li, X. Zhou, X.-X. Shen, & A. Rokas (2023). <u>Incongruence in the phylogenomics era</u>. *Nature Reviews Genetics*. DOI: 10.1038/s41576-023-00620-x
- Steenwyk, J.L., M.A. Phillips, F. Yang, S.S. Date, T. Graham, J. Berman, C.T. Hittinger, & A. Rokas (2022). <u>An orthologous gene coevolution network provides insight into</u> <u>eukaryotic cellular and genomic structure and function</u>. *Science Advances*. DOI: 10.1126/sciadv.abn0105
- Steenwyk, J.L.<sup>^</sup>, T.J. Buida III, A.L. LaBella, Y. Li, X.-X. Shen, & A. Rokas<sup>^</sup> (2021). <u>PhyKIT: a UNIX shell toolkit for processing and analyzing phylogenomic data</u>. <u>^Corresponding authors</u>. *Bioinformatics*. DOI: 10.1093/bioinformatics/btab096
- Steenwyk, J.L.<sup>^</sup>, T.J. Buida III, Y. Li, X.-X. Shen, & A. Rokas<sup>^</sup> (2020). <u>ClipKIT: a multiple sequence alignment-trimming software for accurate phylogenomic inference</u>.
   ^Corresponding authors. *PLOS Biology*. DOI: 10.1371/journal.pbio.3001007
- Steenwyk, J.L.\*, A.L. Lind\*, L.N.A. Ries, T.F. dos Reis, L.P. Silva, F. Almeida, R.W. Bastos, T.F. de Campos Fraga da Silva, V.L.D. Bonato, A.M. Pessoni, F. Rodrigues, H.A. Raja, S.L. Knowles, N.H. Oberlies, K. Lagrou, G.H. Goldman^, A. Rokas^ (2020). Pathogenic allodiploid hybrids of Aspergillus fungi. \*Equal contributors; \*Corresponding authors. Current Biology. DOI: 10.1016/j.cub.2020.04.071

#### AWARDS

| 2024 | Trailblazers of Tomorrow National Postdoctoral Symposium, The University of   |
|------|---|
|      | Texas at Austin   |
| 2024 | Englund Emerging Scholar Award, Department of Biological Chemistry, Johns     |
|      | Hopkins Medicine  |
| 2023 | JXTX + CSHL Genome Informatics Scholarship, Biology of Genomes, Cold Spring   |
|      | Harbor Laboratory   |
| 2023 | Honorable mention, Next Generation Faculty Symposium, Stanford.Berkeley.UCSF  |
| 2023 | Howard Hughes Medical Institute Awardee, Life Sciences Research Foundation    |
| 2022 | Fifty 50 Community Fellow, Fifty Years Industries, LLC                        |
| 2022 | Berkeley Science Fellow, Berkeley Postdoctoral Entrepreneurship Program,      |
|      | University of California, Berkeley  |
| 2022 | Graduate Student Excellence Award Finalist, Society for Molecular Biology and |
|      | Evolution   |
| 2022 | Hanna H. Gray Fellows Finalist, Howard Hughes Medical Institute               |

| 2022 | Edward Ferguson Jr. Graduate Award, Graduate School, Vanderbilt University           |
|------|--|
| 2022 | James F. Crow Early Career Researcher Award Finalist, Genetics Society of            |
|      | America  |
| 2022 | Harold M. Weintraub Graduate Student Award, Fred Hutchinson Cancer Research          |
|      | Center   |
| 2021 | Sandler Fellows Finalist, University of California, San Francisco                    |
| 2021 | Honorable mention, Next Generation Faculty Symposium, Stanford.Berkeley.UCSF         |
| 2021 | Presentation award, Canadian Fungal Research Network and Great Lakes                 |
| 0004 | Mycology Conference  |
| 2021 | Graduate Research Excellence Award in Biological Sciences, Vanderbilt University     |
| 2021 | Smriti Bardhan Scholarship, Vanderbilt University                                    |
| 2021 | Registration award, Science Talk '21   |
| 2020 | Favorite Artist Award, Catalyst: A Virtual Sci-Art Exhibition                        |
| 2020 | Oral presentation award, SACNAS – The National Diversity in STEM Virtual             |
| 2020 | Conference   |
| 2020 | Registration scholarship, SACNAS – The National Diversity in STEM Virtual            |
| 2020 | Conference Best Talk Honorable Mention, Canadian Fungal Research Network Meeting     |
| 2020 | Trainee-of-the-Year, Vanderbilt Institute for Infection, Immunology and Inflammation |
| 2020 | Gilliam Predoctoral Fellowship, Howard Hughes Medical Institute                      |
| 2019 | Ann Bernard Martin Award for Excellence in Graduate Research, Vanderbilt             |
| 2019 | University   |
| 2019 | Ruth L. Kirschstein National Research Service Award, National Institutes of Health   |
| 2019 | Ford Foundation Predoctoral Fellowship, Ford Foundation                              |
| 2019 | Graduate student travel grant, Vanderbilt University                                 |
| 2019 | Curb Center Fellow, ArtLab, Vanderbilt University                                    |
| 2018 | GENETICS Peer Review Training Program, Genetics Society of America                   |
| 2018 | Best poster award, Cellular and Molecular Fungal Biology, Gordon Research            |
| 20.0 | Seminar  |
| 2018 | Best poster award, Cellular and Molecular Fungal Biology, Gordon Research            |
|      | Conference   |
| 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 | Bridging the gaps scholar, 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  |
|      | •  |

# **FUNDING**

| Life Sciences | Principal investigator, 08/23-08/26, Investigating the molecular          |
|---------------|---|
| Research      | underpinnings of complex traits like multicellularity, \$231,000. Funding |
| Foundation    | generously provided by the Howard Hughes Medical Institute.               |

Howard Hughes Medical Institute

Principal investigator, 09/22, The evolution of pathways responsible for genome integrity in early animals and close relatives, Hanna H. Gray

Finalist, \$10,000

**Howard Hughes Medical Institute** 

Principal investigator, 09/19-09/22, Examining the loss of diverse DNA repair genes and long-term hypermutation in a lineage of budding yeasts,

Gilliam Fellowship, Individual Predoctoral Fellowship, \$150,000

National Institutes of Health Principal investigator, 08/19-08/22, Examining the loss of diverse DNA repair genes and long-term hypermutation in a lineage of budding yeasts,

Ruth L. Kirschstein National Research Service Award, Individual

Predoctoral Fellowship (Parent F31), \$88,128 (declined)

Ford Foundation Predoctoral Fellow Principal investigator, 08/19-08/22, The consequences of aberrant cell cycle

and DNA repair processes in budding yeast, Individual Predoctoral

Fellowship, \$72,000 (declined)

Curb Center ArtLab Fellow

Principal investigator, 12/18-04/19, Bridging the gap between artist and

scientist, ArtLab, Vanderbilt University, \$300

#### **SOFTWARE**

1) ClipKIT: a multiple sequence alignment-trimming software for accurate phylogenomic inference. Publication PDF; Documentation; Source code

- 2) PhyKIT: a UNIX shell toolkit for processing and analyzing phylogenomic data. <u>Publication</u> PDF; Documentation; Source code
- 3) BioKIT: a versatile toolkit for processing and analyzing diverse types of sequence data. Publication PDF; Documentation; Source code
- 4) OrthoSNAP: a tree splitting and pruning algorithm for retrieving single-copy orthologs from gene family trees. <u>Publication PDF</u>; <u>Documentation</u>; <u>Source code</u>
- orthofisher: a broadly applicable tool for automated gene identification and retrieval. <u>Publication PDF; Documentation; Source code</u>
- 6) LVBRS: a cloud-based suite of workflows for bulk RNA-seq quality control, analysis, and functional enrichment. <u>Publication PDF</u>; <u>Documentation</u>; <u>Source code</u>
- 7) treehouse: a user-friendly application to obtain subtrees from large phylogenies. Publication PDF; Documentation & source code
- 8) ggpubfigs: an R package for creating colorblind friendly figures with ggplot2. <u>Publication PDF</u>; Documentation & source code

# **INVITED TALKS**

| 2024 | Bioinformatics | Group, | Wageningen | University, | Netherlands |
|------|----------------|--------|------------|-------------|-------------|
|------|----------------|--------|------------|-------------|-------------|

- **2024** Evolutionary Rate covariation and Related Methods Workshop, Oregon State University and Colorado State University
- 2024 Biochemistry, Molecular, and Structural Biology Graduate Program, the University of California, Los Angeles
- 2024 Southern California Evolutionary Genetics and Genomics Meeting, Caltech's Center for Evolutionary Science and the University of California's Center for Ecological and Evolutionary Dynamics, Caltech

| 2024 | Postdoc Research Showcase, Molecular and Cell Biology Department, University of California, Berkeley                                 |
|------|--|
| 2024 | Trailblazers of Tomorrow National Postdoctoral Symposium, The University of  |
| 2024 | Texas at Austin Englund Emerging Scholar Symposium, Department of Biological Chemistry, John   |
| 2024 | Hopkins Medicine 32st Fungal Genetics Conference at Asilomar, Pacific Grove, CA  |
| 2023 | Seminar, Stowers Institute   |
| 2023 | Innovative Genomics Institute and the Department of Electrical Engineering and Computer Sciences, University of California, Berkeley |
| 2023 | CanFunNet, Acadia University   |
| 2023 | Lightening Talk, Southern California Systems Biology Conference, University of Southern California                                   |
| 2023 | Departmental Seminar, Plant Pathology and Environmental Microbiology, The  |
|      | Pennsylvania State University  |
| 2022 | Biology Department, Loras College  |
| 2022 | Yeast Genetics Meeting, Genetics Society of America  |
| 2022 | Hanna H. Gray Fellows Finalists Meeting, Howard Hughes Medical Institute   |
| 2022 | Molecular mycology meeting, Technion - Israel Institute of Technology  |
| 2022 | Evolution, Am. Soc. of Naturalists, Soc. for the Study of Evo., and the Soc. of Sys.   |
| 0000 | Biologists, Cleveland, OH (declined due to scheduling conflict)  |
| 2022 | James F. Crow Early Career Researcher Award Finalist, Genetics Society of America  |
| 2022 | Department of Biological Sciences, George Washington University  |
| 2022 | 31st Fungal Genetics Conference at Asilomar, Pacific Grove, CA   |
| 2021 | Artist-in-Residence program, Vanderbilt Institute for Infection,   |
| 2021 | Immunology and Inflammation CanFunNet and Great Lakes Mycology Conference  |
| 2021 | Sandler Fellows Finalists Seminar, University of California, San Francisco   |
| 2021 | Department of Ecology, Evolution, and Organismal Biology, Iowa State University  |
| 2021 | Medical Mycology Trainee Seminar Series, University of Utah (Link)   |
| 2021 | Mycology Graduate Student Organization, University of Georgia  |
| 2021 | MicroSeminar, International Society for Microbial Ecology (Link)   |
| 2021 | Alliance for Diversity in Science and Engineering, Young Researchers Conference  |
| 2021 | Andrew Murray Lab seminar, Harvard University, Cambridge   |
| 2020 | Institute of Insect Sciences, Zhejiang University  |
| 2020 | Evan Eichler Lab seminar, University of Washington, Seattle  |
| 2020 | Genetics Society of America, Early Career Scientist Seminar Series   |
| 2020 | Nicole King Lab seminar, University of California Berkeley   |
| 2020 | The National Diversity in STEM Conference, SACNAS  |
| 2020 | Canadian Fungal Research Network Meeting   |
| 2020 | Trainee-of-the-year talk, Vanderbilt Institute for Infection, Immunology and Inflammation  |
| 2020 | Day of Wond'ry, Vanderbilt University, Nashville, TN   |
| 2019 | Genetics Society of America, Early Career Scientist Seminar Series   |
| 2019 | Gordon Research Conference, Molecular Mechanisms in Evolution, Easton, MA  |
| 2019 | Gordon Research Seminar, Molecular Mechanisms in Evolution, Easton, MA (declined)  |
| 2019 | Focal Point, ArtLab, Vanderbilt University, Nashville, TN  |
| 2019 | 30th Fungal Genetics Conference at Asilomar, Pacific Grove, CA   |

| 2019<br>2018<br>2015 | Phylogenomics and Evolution Group, North Carolina State University, Raleigh, NC ArtLab Seminar Series, Vanderbilt University, Nashville, TN TedXClarkUniversity, Clark University, Worcester, MA |
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## **CONTRIBUTED TALKS**

| Plant and Microbial Biology, Postdoc Seminar Series, University of California,  |
|---|
| Berkeley, Berkeley, CA  |
| Students' Mycology Colloquium, Mycological Society of America                   |
| Evolution Seminar Series, Vanderbilt University (Link)                          |
| DNA Damage and Response Journal Club, Vanderbilt University, Nashville, TN      |
| Research in Progress Seminar, Vanderbilt University, Nashville, TN              |
| Biological Sciences Annual Retreat, Vanderbilt University, Nashville, TN        |
| Science club at the library, Nashville Public Library, Nashville, TN            |
| Nashville Science Club, Jackalope Brewing Company, Nashville, TN                |
| Mycological Society of America, University of Georgia, Athens, GA               |
| Mycological Society of America, University of California Berkeley, Berkeley, CA |
| Graduate Student Multidisciplinary Conference, Clark University, Worcester, MA  |
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## TRAINEE ADVISING

# **Graduate Students**

**2023-Pres.** Saelin Bjornson, University of Melbourne (co-advised with Heroen Verbruggen)

# **Undergraduates**

| 2024-Pres. | Veronica L | Uy, Universit | y of California, | Berkeley |
|------------|------------|---------------|------------------|----------|
|------------|------------|---------------|------------------|----------|

**2022-2024** Charu Balamurugan, Vanderbilt University (co-advised with Antonis Rokas)

2019-2022 Qianhui (Olivia) Zheng, Vanderbilt University (co-advised with Antonis Rokas)

**2018-2021** Megan A. Phillips, Vanderbilt University (co-advised with Antonis Rokas)

## **COMPANY ADVISING**

| <b>Since 2023</b> | Advisor, ForensisGroup, Inc., Pasadena, CA                   |
|-------------------|--|
| 2022-2023         | Scientific Advisor, WittGen Biotechnologies, CA              |
| 2022-2023         | Scientific Consultant, Latch Al Inc., San Francisco, CA      |
| 2017-2018         | Scientific Consultant, Little Harpeth Brewing, Nashville, TN |

## **WORKSHOP TEACHING**

| 2024 | Instructor, Phylogenomics and Population Genomics: Inference and Applications, University of Barcelona, Barcelona, Spain |
|------|--|
| 2024 | Lead instructor, Workshop on Phylogenomics, Evolution and Genomics, Český<br>Krumlov, Czech Republic                     |
| 2019 | Organizer and instructor, Values-based leadership, Vanderbilt University, Nashville, TN                                  |
| 2019 | Founder and instructor, 'A beginner's guide to making figures in R,' Vanderbilt University, Nashville, TN                |
| 2019 | Instructor, Workshop on Phylogenomics, Evolution and Genomics, Český Krumlov,<br>Czech Republic                          |
| 2019 | Instructor, Workshop on Genomics, Evolution and Genomics, Český Krumlov,<br>Czech Republic                               |

## **TEACHING EXPERIENCE**

| 2020 | Guest lecture, Science Communication Tools and Techniques, Vanderbilt |
|------|---|
|      | University, Nashville, TN   |

2017-2019 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

### POSTER PRESENTATIONS

| 2024 | Life Sciences Research Foundation Fellows Annual Meeting, Chicago, IL  |
|------|--|
| 2023 | Genome Informatics, Cold Spring Harbor Laboratory  |
| 2023 | The International Choanoflagellates & Friends Workshop   |
| 2022 | Decoding the Genome, Howard Hughes Medical Institute   |
| 2021 | Gilliam Fellows Meeting, Howard Hughes Medical Institute   |
| 2021 | Biology of Genomes, Cold Spring Harbor Laboratories  |
| 2021 | Science Talk '21, Science Talk   |
| 2020 | Gilliam Fellows Meeting, Howard Hughes Medical Institute   |
| 2020 | Vanderbilt Institute for Infection, Immunology and Inflammation Annual Symposium,  |
|      | Virtual Conference   |
| 2020 | The Allied Genetics Conference, Virtual Conference   |
| 2019 | Investigators Science Meeting, Howard Hughes Medical Institute, Bethesda, MD   |
| 2019 | Gilliam Fellows Annual Meeting, Howard Hughes Medical Institute, Bethesda, MD  |
| 2019 | Molecular Mechanisms in Evolution, Gordon Research Conference, Easton, MA  |
| 2019 | Molecular Mechanisms in Evolution, Gordon Research Seminar, Easton, MA   |
| 2019 | 30th Fungal Genetics Conference at Asilomar, Pacific Grove, CA   |
| 2019 | Asperfest pre-meeting at 30 <sup>th</sup> Fungal Genetics Conference at Asilomar, Pacific  |
|      | Grove, CA  |
| 2018 | Cellular and Molecular Fungal Biology, Gordon Research Conference, Holderness,   |
|      | NH   |
| 2018 | Cellular and Molecular Fungal Biology, Gordon Research Seminar, Holderness, NH   |
| 2018 | Department of Biological Sciences Annual Retreat, Vanderbilt University, Nashville, TN   |
| 2015 | Bumpus Symposium, Clark University, Worcester, MA  |
| 2015 | Traina Scholars Presentation, Clark University, Worcester, MA  |
| 2015 | Summer Research Presentation, Clark University, Worcester, MA  |
|      | Talling in the second s |

## **EDUCATION**

| Vanderbilt University | Biological Sciences                | Ph.D., GPA: 3.97 | 2016–2022 |
|-----------------------|------------------------------------|------------------|-----------|
| Clark University      | Biochemistry and Molecular Biology | M.S., GPA: 3.98  | 2015–2016 |
| Clark University      | Biochemistry and Molecular Biology | B.A., GPA: 3.84  | 2011–2015 |

#### SERVICE

2024-Pres. Member, Conferences Committee, Genetics Society of America

**2024-Pres.** Co-chair, Postdocs Research Showcase, Molecular and Cell Biology, University of California, Berkeley

**2023-Pres.** Mentor, Alumni Mentorship Program Early Career Mentorship Program, Genetics Society of America

2023-Pres. Project Partner, Data Science Discovery Program, University of California, Berkeley

**2023-Pres.** Chair of Alumni Affairs, The Evolutionary Studies Initiative at Vanderbilt, Vanderbilt University, Nashville, TN

- 2023-Pres. Founder, Genomely Internship in Genome Sciences Program, Genomely
- 2020-Pres. Founder and Chief Officer, SciArt with Purpose, https://jlsteenwyk.com/sciart.html
- **2023-2024** Executive Board Member, Berkeley Postdoctoral Entrepreneurship Program, University of California, Berkeley
- **2023-2024** Chair, TAGC Undergraduate Travel Award Subcommittee, Genetics Society of America
- **2023-2024** Representative for Early Career Scientists, Allied Program Committee, The Allied Genetics Conference 2024, Genetics Society of America
- 2022 Panelist at the Diversity, Equity, and Inclusion Discussion, Yeast Genetics Conference, Genetics Society of America
- 2022 Scientist-Artist: Embracing Duality, ArtLab, Vanderbilt University
- **2019-2022** Member, Steering Committee, Early Career Leadership Program, Genetics Society of America
- **2019-2022** Inclusion Coordinator, The Evolutionary Studies Initiative at Vanderbilt, Vanderbilt University, Nashville, TN
- **2019-2022** Graphic Illustrator, The Evolutionary Studies Initiative at Vanderbilt, Vanderbilt University, Nashville, TN
- 2017-2022 Educational outreach booth design and execution, MEGA Microbe, Nashville, TN
- **2017-2022** Member of the Dean of Graduate Student's survey quantitative analysis subgroup, Graduate Diversity and Inclusion Committee, Vanderbilt University, Nashville, TN
- **2017-2022** Judge, Middle Tennessee Science and Engineering Fair, Belmont University, Nashville, TN
- **2019-2021** Co-chair, Communication and Outreach Subcommittee, Genetics Society of America
- **2018-2021** Volunteer Deputy, American Society of Microbiology Vanderbilt University Chapter, Nashville, TN
- **2017-2021** Communications chair, Inclusivity in Biosciences Association, Vanderbilt University, Nashville, TN
- 2020 Panelist at the Communication and Outreach Workshop, The Allied Genetics Conference, Genetics Society of America
- **2019-2020** President, Inclusivity in Biosciences Association, Vanderbilt University, Nashville, TN
- **2019-2020** Co-chair, MEGA*Microbe*, Vanderbilt Institute for Infections, Immunology and Inflammation, Nashville, TN
- **2018-2019** Vice President, Inclusivity in Biosciences Association, Vanderbilt University, Nashville, TN
- **2013-2019** Administrator and Owner, Molecular Biology and Biochemistry for Researchers and Students Group, LinkedIn
- Peer review workshop leader, 30<sup>th</sup> Fungal Genetics Conference at Asilomar, Pacific Grove, CA
- **2018-2019** Vice President, Graduate Student Association, Department of Biological Sciences, Vanderbilt University, Nashville, TN
- **2018-2019** Vice co-chair, MEGA*Microbe*, Vanderbilt Institute for Infections, Immunology and Inflammation, Nashville, TN
- **2017-2018** Secretary, Graduate Student Association, Department of Biological Sciences, Vanderbilt University, Nashville, TN
- 2017 Vanderbilt Student Volunteers for Science, Volunteer Science Teacher, West End Middle School, Nashville, TN
- **2014-2015** Director, Clark University Emergency Medical Services, Clark University, Worcester, MA

- **2014-2015** 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
- **2013-2014** Secretary, Clark University Emergency Medical Services, Clark University, Worcester, MA

#### **SOCIETIES**

Genetics Society of America, Society of Systematic Biologists, American Society for Microbiology, Mycological Society of America, Society for the Advancement of Chicanos/Hispanics and Native Americans in Science

### MANUSCRIPT REVIEWER

Nature Communications; Proceedings of the National Academy of Sciences, Molecular Biology and Evolution; Systematic Biology; PLOS Pathogens, GigaScience, Methods in Ecology and Evolution; Genome Biology and Evolution; Genetics; G3 Genes|Genomes|Genetics; FEMS Yeast Research; and others

#### **ART SHOWS**

| 2021 | Science Talk '21, Science Talk  |
|------|---|
| 2020 | Catalyst: A Virtual Sci-Art Exhibition, Michigan State University                   |
| 2020 | Day of Wond'ry, Vanderbilt University, Nashville, TN                                |
| 2020 | Fire-Exhibition, Kefi Collective at Vanderbilt University, Nashville, TN            |
| 2019 | Biomedical Sciences Winter Show, Vanderbilt University, Nashville, TN               |
| 2019 | Focal point, ArtLab, Vanderbilt University, Nashville, TN                           |
| 2019 | Connecting the Dots, ArtLab, Vanderbilt University, Nashville, TN                   |
| 2018 | ArtLab opening reception, ArtLab, Vanderbilt University, Nashville, TN              |
| 2018 | The Intersection between Art and Science, ArtLab, Vanderbilt University, Nashville, |
|      | TN  |

#### POPULAR SCIENCE ARTICLES

- 6. **Steenwyk, J.L.** & K. Giffin. The silver lining of bioinformatics. *Genes to Genomes*. September 12, 2022
- 5. Simopoulos, M.A.C., A.F. Cisneros, A.D. Mendoza, C. Bautista, **J.L. Steenwyk**, N. Ahmad. Hurdles and advances to making science gender-neutral, *ecrLife*. November 26, 2020
- 4. Mendoza, A.D., C. Bautista, E.A. Marnik, C.M.A. Simopoulos, & **J.L. Steenwyk**. Navigating fake news as a scientist, *ecrLife*. October 8, 2020
- 3. **Steenwyk, J.L.** & M. Jonika. How to get started in science communication, *ecrLife*. August 21, 2020
- 2. **Steenwyk, J.L.** & A. Rokas. A new hybrid fungus is found in hospitals and linked to lung disease, *The Conversation*. June 4, 2020
- 1. **Steenwyk**, **J.L.** & A. Rokas. An outlaw yeast thrives with genetic chaos and could provide clues for understanding cancer growth, *The Conversation*. May 21, 2019

# **PUBLICATIONS** (Trainees and I are in **bold** font.)

# **Preprints/Submitted**

13. O. Lemke\*, B.M. Heineike\*, S. Viknander, N. Cohen, J.L. Steenwyk, L. Spranger, F. Li, F.

- Agostini, C.T. Lee, S.K. Aulakh, J. Nielsen, A. Rokas, J. Berman, A. Zelezniak, T.I. Gossmann, & M. Ralser (2024). (\*Equal contributors). <u>The Role of Metabolism in Shaping Enzyme Structures Over 400 Million Years of Evolution</u>. *bioRxiv*. DOI: 10.1101/2024.05.27.596037
- 12. Li, N., D.M. Geiser, **J.L. Steenwyk**, C. Tsuchida, S. Koike, S. Slinski, & F.N. Martin (2024). A systematic approach for identifying unique genomic sequences for *Fusarium oxysporum* f. sp. *lactucae* race 1 and development of molecular diagnostic tools. *Submitted*.
- Steenwyk, J.L., G.I. Martínez-Redondo, T.J. Buida III, E. Gluck-Thaler, X.-X. Shen, T. Gabaldón, A. Rokas, & R. Fernández (2024). <a href="Phykit">Phykit: A Multitool for Phylogenomics</a>. <a href="Preprints">Preprints</a>. DOI: 10.20944/preprints202404.1514.
- Fan, Y., M. Du, W. Zhang, W. Deng, E. Yang, S. Wang, L. Yan, L. Zhang, S. Kang, J.L. Steenwyk, Z. An, X. Liu, & M. Xiang (2024). <u>The Genomes of Nematode-Trapping Fungi Provide Insights into the Origin and Diversification of Fungal Carnivorism</u>. *bioRxiv*. DOI: 10.1101/2024.03.21.586190.
- 9. Zhang, W., Y. Fan, W. Deng, Y. Chen, S. Wang, S. Kang, **J.L. Steenwyk**, M. Xiang, & X. Liu (2024). Characterization of Genome-wide Phylogenetic Conflict Uncovers Evolutionary Modes of Carnivorous Fungi. *bioRxiv*. DOI: 10.1101/2024.03.21.586083.
- 8. Turnbull, R.\*, **J.L. Steenwyk**\*, S. Mutch, P. Scholten, V.W. Salazar, J.L. Birch, & H. Verbruggen (2023). \*Equal contributors. <u>Orthoflow: phylogenomic analysis and diagnostics with one command</u>. *Research Square*. DOI: 10.21203/rs.3.rs-3699210/v1
- 7. Goncalves, C., M.-C. Harrison, **J.L. Steenwyk**, D.A. Opulente, A.L. LaBella, J.F. Wolters, X. Zhou, X.-X. Shen, M. Groenewald, C.T. Hittinger, & A. Rokas (2023). <u>Diverse signatures of convergent evolution in cacti-associated yeasts</u>. *bioRxiv*. DOI: 10.1101/2023.09.14.557833.
- 6. **Bjornson, S.**, H. Verbruggen, N. Upham, & **J.L. Steenwyk** (2023). Reticulate Evolution: Detection and Utility in the Phylogenomics Era. *Preprints*. DOI: 10.20944/preprints202309.0905.v2.
- 5. Liu, H.\*, **J.L. Steenwyk**\*, X. Zhou, D.T. Schultz, K.M. Kocot, X.-X. Shen, A. Rokas^, & Y. Li^ (2023). \*Equal contributors; ^Corresponding authors. <u>A genome-scale Opisthokonta tree of life: toward phylogenomic resolution of ancient divergences</u>. **bioRxiv**. DOI: 10.1101/2023.09.20.556338
- 4. G.H. Goldman<sup>^</sup>, Valero, C.\*, Pinzan, C.F.\*, P.A. de Castro, N. van Rhijn, K. Earle, H. Liu, M.A.C. Horta, O. Kniemeyer, T. Krüger, A. Pschibul, A.A. Brakhage, J.L. Steenwyk, M.E. Mead, A. Rokas, S.G. Filler, H. Cabral, E. Deljabe, M.J. Bromley, G. Palmisano, A.S. Ibrahim, S. Gago<sup>^</sup>, & T.F. dos Reis<sup>^</sup> (2023). \*Equal contributors; <sup>^</sup>Corresponding authors. A phylogenetic approach to explore the *Aspergillus fumigatus* conidial surface-associated proteome and its role in pathogenesis. bioRxiv. DOI: 10.1101/2023.08.22.553365v1
- Steenwyk, J.L., S. Knowles, R.W. Bastos, C. Balamurugan, D. Rinker, M.E. Mead, C.D. Roberts, H.A. Raja, Y. Li, A.C. Colabardini, P.A. de Castro, T.F. dos Reis, D. Canóvas, R.L. Sanchez, K. Lagrou, E. Torrado, F. Rodrigues, N.H. Oberlies, X. Zhou, G.H. Goldman<sup>^</sup>, & A. Rokas<sup>^</sup> (2023). Corresponding authors. Evolutionary origin, population diversity, and diagnostics for a cryptic hybrid pathogen. bioRxiv. DOI: 10.1101/2023.07.03.547508.
- 2. Le, H.G.B.H.<sup>^,\*</sup>, **J.L. Steenwyk**\*, N. Manske, M. Smolin, A. Abdulali, A. Kamat, R.

- Kanchana, K. Giffin, A. Andere, & K. Workman<sup>^</sup> (2022). \*Equal contributors; ^Corresponding authors. <u>Latch Verified Bulk-RNA Seq toolkit: a cloud-based suite of workflows for bulk RNA-seq quality control, analysis, and functional enrichment</u>. bioRxiv. DOI: 10.1101/2022.11.10.516016.
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