JL Weissman

pronouns: they/them/theirs or she/her/hers **website:** https://microbialgamut.com/

Positions

Assistant Professor of Computational Biology, Schmid College of Science and Technology, Chapman University, Orange, CA, 2023 -

Adjunct Assistant Professor of Biological Sciences, Dana and David Dornsife College of Letters, Arts and Sciences, University of Southern California, Los Angeles, CA, 2023 -

Postdoctoral Fellow, Department of Marine and Environmental Biology, University of Southern California. Los Angeles. CA. 2020 - 2022

Advisor: Jed A. Fuhrman

Education

Ph.D., Behavior, Ecology, Evolution, and Systematics (BEES; Fall 2019), University of Maryland College Park, College Park, MD

Investigating the Distribution of CRISPR Adaptive Immune Systems Among Prokaryotes Advisors: Philip L.F. Johnson & William F. Fagan

Graduate Certificate, Computation and Mathematics for Biological Networks (COMBINE; Fall 2019), University of Maryland College Park, College Park, MD

Network-Based Prediction of Novel CRISPR-Associated Genes in Metagenomes

B.A., Mathematics and Biology (Spring 2015), Bard College, Annandale-on-Hudson, NY Modeling Stochastic Traps

Advisors: Csilla Szabo & Bruce Robertson

Summer Schools:

EMergent Ecosystem Responses to ChanGE Summer Program, EMERGE, NH, 2022 Microbial Diversity Course, Marine Biological Laboratory, Woods Hole, MA, 2018 The Search for Selection, NIMBioS, Knoxville, TN, 2018 Complex Systems Summer School, Santa Fe Institute, Santa Fe, NM, 2017 Quantitative Laws II, Lake Como School of Advanced Studies, Italy, 2016

External Funding

Simons Foundation Postdoctoral Fellowship in Marine Microbial Ecology (\$246, 100), Simons Foundation, 2020-2023

Drafting an Atlas of Prokaryotic Immune Strategy in the Global Oceans

Publications

Preprints/Submitted:

- 1. Dimbo E.R.O., J.A. Fuhrman, <u>J.L. Weissman</u>. No Relationship between Minimum Doubling Time and Genome Length in Microbial Eukaryotes. submitted
- 2. Weissman J.L., E.R.O. Dimbo, A.I. Krinos, C. Neely, Y. Yagües, D. Nolin, S. Hou, S. Laperriere, D.A. Caron, B. Tully, H. Alexander, J.A. Fuhrman. Estimating the maximal growth rates of eukaryotic microbes from cultures and metagenomes via codon usage patterns. bioRxiv
- 3. Fletcher-Hoppe C., Y.C. Yeh, Y. Raut, <u>J.L. Weissman</u>, J.A. Fuhrman. **Symbiotic diazotrophic** UCYN-A strains co-occurred with El Niño, relaxed upwelling, and varied eukaryotes over 10 years off Southern California Bight. *bioRxiv*
- 4. Ignacio-Espinoza J.C., S. Laperriere, Y. Yeh, <u>J.L. Weissman</u>, S. Hou, M. Long, J.A. Fuhrman. Ribosome-linked mRNA-rRNA chimeras reveal active novel virus host associations. bioRxiv

Peer Reviewed:

- 1. Weissman J.L., M. Peras, T.P. Barnum, J.A. Fuhrman. 2022. Benchmarking community-wide estimates of growth potential from metagenomes using codon usage statistics. mSystems 7(5), e00745-22.
- 2. Tao J., W. Wang, <u>J.L. Weissman</u>, Y. Zhang, S. Chen, Y. Zhu, C. Zhang, and S. Hou. 2022. **Size-fractionated microbiome observed during an eight-month long sampling in Jiaozhou Bay and the Yellow Sea**. *Scientific Data* 9(1), 1-8.
- 3. Gleich S.J, J.A. Cram, <u>J.L. Weissman</u>, D.A. Caron. 2022. **NetGAM: Using generalized additive models to improve the predictive power of .ecological network analyses constructed using time-series data**. *ISME Communications* 2(1), 23
- 4. Weissman J.L., S. Hou, J.A. Fuhrman. 2021. Estimating maximal microbial growth rates from cultures, metagenomes, and single cells via codon usage patterns. Proceedings of the National Academy of Sciences 118(12), e2016810118.
- Weissman J.L., E.O. Alseth, S. Meaden, E.R. Westra, J.A. Fuhrman. 2021. Immune Lag Is a Major Cost of Prokaryotic Adaptive Immunity During Viral Outbreaks. Proceedings of the Royal Society B 288, 20211555.
- 6. Weissman J.L., ..., P.L.F. Johnson, D. Karig, W.F. Fagan, S. Bewick. 2021. Exploring the functional composition of the human microbiome using a hand-curated microbial trait database. *BMC Bioinformatics* 22(1), 1471-2105.
- 7. Tully B.J, J. Buongiorno, ..., L.E. Valentin-Alvarado, <u>J.L. Weissman</u>, BVCN Instructor Consortium. 2021. The Bioinformatics Virtual Coordination Network: an open-source and interactive learning environment. *Frontiers in Education* 6, 394.
- 8. Weissman J.L., A. Stoltzfus, E.R. Westra, P.L.F. Johnson. 2020. Avoidance of Self During CRISPR Immunization. Trends in Microbiology 28(7), 543-553.
- Weissman J.L. & P.L.F. Johnson. 2020. Network-Based Prediction of Novel CRISPR-Associated Genes in Metagenomes. mSystems 5(1), e00752-19.
- Martí-Carreras J., ..., <u>J.L. Weissman</u>, V. Zalunin, A. Efremov, B. Busby. 2020. NCBI's Virus Discovery Codeathon: Building the "FIVE" – Federated Index of Viral Experiments API index. Viruses 12(12), 1424.
- 11. Weissman J.L., W.F. Fagan, P.L.F. Johnson. 2019. Linking high GC content to the repair of double strand breaks in prokaryotic genomes. *PLOS Genetics* 15(11), e1008493.
- 12. Weissman J.L., R. Laljani, W.F. Fagan, P.L.F. Johnson. 2019. Visualization and prediction of CRISPR incidence in microbial trait-space to identify drivers of antiviral immune strategy. The ISME Journal 13(10), 2589-2602.
- 13. Bewick S., E. Gurarie, <u>J.L. Weissman</u>, J. Beattie, C. Davati, R. Flint, P. Thielen, F. Breitwieser, D. Karig, W.F. Fagan. 2019. **Trait-Based Analysis of the Human Skin Microbiome**. *Microbiome* 7(1), 101.
- 14. Weissman J.L., W.F. Fagan, P.L.F. Johnson. 2018. Selective maintenance of multiple CRISPR arrays across prokaryotes. The CRISPR Journal 1(6), 405-413.
- Weissman J.L., R. Holmes, R. Barrangou, S. Moineau, W.F. Fagan, B. Levin, P.L.F. Johnson. 2018. Immune Loss as a Driver of Coexistence During Host-Phage Coevolution. The ISME Journal 12(2), 585-597.

Outreach Publications:

- 1. Weissman J. L., S. Hou, J.A. Fuhrman. Using DNA to Predict How Fast Bacteria Can Grow. Frontiers Young Minds.
- 2. Weissman J. L., H.H. Yiu, P.L.F. Johnson. 2019. What Bacteria Do When They Get Sick. Frontiers Young Minds.

Fellowships and Awards

COMBINE Network Science Fellowship (\$35, 859, UMD/NSF DGE-1632976, 2018)
Devra Kleiman Memorial Scholarship (\$2,500, UMD, 2018)
GAANN Mathematical Biology Fellowship (\$74,598, UMD/U.S. Department of Education, 2015)
The Flagship Fellowship (\$50,000, UMD, 2015)

The Dean's Fellowship (\$10,000, UMD, 2015)

The Harry J. Carman Scholarship (Bard College, 2014)

The John Bard Scholarship (Division of Science, Math and Computing, Bard College, 2013)

George I. Alden Scholar (George I. Alden Trust, 2012)

The Excellence and Equal Cost Scholarship (Bard College, 2011)

The Bishop Scholarship (The Bishop Scholarship Foundation, 2011)

Invited Talks

- 1. Physics of Life Sciences, Massachusetts Institute of Technology, MA, 2022
- 2. Department of Biology, California State University Los Angeles, CA, 2022
- 3. Department of Biology, University of Minnesota Duluth, MN, 2022
- 4. Simons Collab. on Comp. Biogeochem. Modeling of Marine Ecosystems (CBIOMES), 2021
- 5. Center for Dark Energy Biosphere Investigations (C-DEBI), 2021
- 6. Department of Biology, University of Pittsburgh, PA, 2021
- 7. Center for Advanced Biotechnology and Medicine, Rutgers University, NJ, 2021
- 8. Department of Biology, Chapman University, CA, 2021
- 9. Department of Biology, Westchester University, PA, 2021
- 10. Department of Biology, Lake Forest College, IL, 2021
- 11. Department of Biology, San Diego State University, CA, 2021
- 12. Computation and Mathematics for Biological Networks (COMBINE), UMD, MD, 2020
- 13. Department of Marine and Environmental Biology, USC, CA, 2020
- 14. Environment and Sustainability Institute, University of Exeter, Penryn, UK, 2019
- 15. Clemson University, Clemson, SC, Dec 2018

Teaching

Instructor, Qlife Quantitative Biology Winter School: Quantitative Viral Dynamics Across Scales, Département de Biologie, École normale supérieure, Paris, France, March 2022

Facilitator, Marine and Environmental Biology Statistics Reading Group, USC, 2020-2022

Instructor, Bioinformatics Virtual Coordination Network, 2020-2021

https://biovcnet.github.io/

Guest Lecturer, University of Maryland College Park

- BIOL709F: Statistics and Modeling for Biologists, Spring 2018
- BSCI464: Microbial Ecology, Spring 2018
 BSCI405: Population and Evolutionary Genetics, Spring 2017

Teaching Assistant, Calculus for the Life Sciences, Department of Biology, University of Maryland College Park, Fall 2015

Teaching Assistant, Bridge to Enter Advanced Mathematics (BEAM), The Art of Problem Solving Foundation, Cambridge, MA, Summer 2012 and 2015

Mentoring

Panelist, Path of Professorship Workshop, MIT, 2022

Panelist, Effective Time Management for New Graduate Students (Orientation), USC, 2022 Workshop Organizer, Mentoring Undergraduate Researchers: A Practical Guide for Graduate Student Mentors, University of Maryland College Park, 2019

Undergraduate Researchers Mentored (*coauthor, †presented at symposium or conference) University of Southern California:

Edward-Robert Dimbo* (Winter 2021 - Summer 2022; GGURE)

James Rosas[†] (Summer 2021; National Summer Undergraduate Research Project, nsurp.org) Yuniba Yagües*† (Summer 2020; National Summer Undergraduate Research Project, nsurp.org) Oscar Escobedo[†] (Summer 2020; National Summer Undergraduate Research Project, nsurp.org)

University of Maryland College Park:

Rohan Laljani *† (Fall 2017 - Summer 2019)

Vinay Veluvolu † (Summer 2018 - Summer 2019)

Julia Gall[†] (Fall 2018 - Spring 2019; College Park Scholars)

Cori Butkiewicz (Summer 2017 - Spring 2018)

Service

DEI Taskforce Member, Schmid College, Chapman University, 2022-

Facilitator, DEI Journal Club on Ableism in Evolutionary Biology, USC, 2022

Reviewer, CAREER Awards, NSF, 2022

Session Co-Chair, Ocean Sciences Meeting, 2022

DEI Committee Member, Dept. of Marine and Environmental Biology (MEB), USC, 2021-2022 **DEI Subcommittee on Reporting Member,** MEB, USC, 2021-2022

Judge, ASM Early Career Flash Talks, October 2021

Conference Organizing Committee, Holistic Bioinformatics Approaches Used in Microbiome Research, Bioinformatics Virtual Coordination Network, Summer 2020-Summer 2021

Facilitator, Pride Month Workshop on Queering Professionalism, USC, June 2021

Co-Organizer, Pride Month Programming for MEB, USC, June 2021

Poster Judge, CRISPR 2021 Meeting, June 2021

Secretary-Treasurer, BEES Student Taskforce (BEESst), UMD-CP, Fall 2018-Spring 2019

Co-President, BEES Student Taskforce (BEESst), UMD-CP, Fall 2017-Spring 2018

Graduate Mentor, Biological Sciences Graduate Program, UMD-CP, Fall 2016, 2017

Journal Reviewer: ISME J, PNAS, Current Biology, Nature Communications, mSystems, Microbiome, Proceedings B, PLoS Comp Bio, Marine Genomics, npj Biofilms and Microbiomes, BMC Bioinformatics, J Theor Biol

Outreach

Co-Organizer, Postdoc Outreach Project with LA Public Library, USC, 2020-2022 Project Mentor, Terps in Space, 2017, 2018, 2022
Tutor & Mentor, Joint Educational Project, USC, Spring 2020
Lead Organizer, Frontiers Young Minds Writing Group, USC, Winter-Spring 2020
Panel Reviewer, Student Spaceflight Experiments Program (SSEP) 2016-2019
Maryland Day Organizer, Biological Sciences, UMD-CP, 2016-2018
Project Leader, Bard Math Circle, Annandale-on-Hudson, NY, 2013-2015
Public Talk, GradTerps Exchange, University of Maryland, Spring 2019
Public Talk, Skype a Scientist, Summer 2020

Conference Presentations (*poster, †talk)

Predicting maximal microbial growth rates from genomes and metagenomes for prokaryotes and eukaryotes

- • Marine Microbes, Gordon Research Conference and Seminar, Switzerland, May $2022^{\dagger\star}$
- Fifth Workshop On Trait-Based Approaches to Ocean Life, Knoxville, TN, January 2022*

Estimating maximal microbial growth rates from cultures, metagenomes, and single cells via codon usage patterns

- Microbial Ecology & Evolution Virtual (MEEVirtual), online, August 2020*
- CBIOMES Annual Meeting, online, June 2020*

Linking selection for high GC content to repair of double strand breaks in prokaryotic genomes

• Microbial Population Biology, Gordon Research Conference and Seminar, NH, July 2019*

Ecology Shapes Microbial Immune Strategy: Temperature and Oxygen as Determinants of the Incidence of CRISPR Adaptive Immunity

- CRISPR Ecology and Evolution, the Royal Society, London, UK, February 2019*
- Microbial Eco-Evolutionary Dynamics, Instituto Gulbenkian De Ciência, Oeiras, Portugal, October 2018*

Is Having more than one CRISPR array adaptive?

- \bullet Microbial Population Biology, Gordon Research Conference and Seminar, NH, July 2017^\star Immune Loss as a Driver of Coexistence During Host-Phage Coevolution

 - Molecular Coevolution Workshop, Princeton Center for Theoretical Science, April 2016^{\star}