BOB WEEK

Curriculum Vitae

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2022

2019

2018

PhD Bioinformatics & Computational Biology 2020

S.L. Nuismer Lab, IBEST, University of Idaho

Dissertation focused on modeling eco-evolutionary processes and developing statistical methods

2015 **BS Mathematics** Traditional math degree with electives in electrical engineering

PEER-REVIEWED PUBLICATIONS

Quantitative Genetics of Microbiome Mediated Traits 2025

Evolution

University of Idaho

Week, B.; Ralph, P.L.; Tavalire, H.F.; Cresko, W.A.; Bohannan, B.J.M.

doi:tbd

2025 Applying Evolutionary Theory to Understand Host-Microbiome Evolution Week, B.; Russel, S.L.; Schulenburg, H.; Bohannan, B.J.M.; Bruijning, M.

Nature EcoEvo doi:tbd

Host-Parasite Coevolution in Continuous Space 2023

Leads to Variation in Local Adaptation Across Spatial Scales

The American Naturalist doi:10.1086/727470

Week, B.; Bradburd, G.S.

Uncovering Cryptic Coevolution

The American Naturalist

Nuismer, S.L.; Week, B.; Harmon, L.J.

doi:10.1086/717436

A White Noise Approach to Evolutionary Ecology 2021

Week, B.; Nuismer, S.L.; Harmon, L.J.; Krone, S.M.

Journal of Theoretical Biology doi:10.1016/j.jtbi.2021.110660

2021 **Coevolutionary Arms Races**

and the Conditions for the Maintenance of Mutualism

The American Naturalist doi:10.1086/714274

Week, B.; Nuismer, S.L.

A Unified Model of Species Abundance, Genetic Diversity, and Functional Diversity 2021

Reveals the Mechanisms Structuring Ecological Communities

Molecular Ecology Resources

Overcast, I.; Ruffley, M.; Rosindell, J.; Harmon, L.; Borges, P.; Emerson, B.; Etienne, R.S.; Gillespie, R.; Krehenwinkel, H.; Mahler, L.; Massol, F.; Parent, K.; Patiño, J.; Peter, B.; Week, B.; Wagner, C.; Hickerson, doi:10.1111/1755-0998.13514

M.J.; Rominger, A.

Identifying Models of Trait-Mediated Community Assembly

using Random Forests and Approximate Bayesian Computation

Ecology and Evolution

Ruffley, M.; Peterson, K.; Week, B.; Tank, D.; Harmon, L.J.

doi:10.1002/ece3.5773

Approximate Bayesian Estimation of Coevolutionary Arms Races 2019

Nuismer, S.L.: Week, B.

PLOS Computational Biology doi:10.1371/journal.pcbi.1006988

The Measurement of Coevolution in the Wild 2019

Ecology Letters

Week, B.; Nuismer, S.L.

doi:10.1111/ele.13231

Coevolution Slows the Disassembly of Mutualistic Communities

Nuismer, S.L.; Week, B.; Aizen, M.

The American Naturalist doi:10.1086/699218

PREPRINTS

Stochastic Eco-Evolutionary Dynamics of Multivariate Traits 2025

bioRxiv

Week, B.

doi:10.1101/2025.06.07.658444

The Evolution of Microbiome-Mediated Traits

bioRxiv

Week, B.; Morris, A.H.; Bohannan, B.J.M.

doi:10.1101/2024.03.29.587374

AWARDS

2024

2024 - 2027 **EU/DFG Postdoctoral Fellowship** KiTE, Kiel University

The "Kiel Training for Excellence" programme is cofunded by the Marie Sklodowska-Curie Actions from the European Commission's Horizon Europe programme (project number: 101081480) and by Kiel University

Bioinformatics & Computational Biology Fellowship 2018 - 2019

IBEST, University of Idaho

Project aimed to model the duration of coevolutionary associations

2017-2018	Bioinformatics & Computational Biology Fellowship Project aimed to develop a statistical method to measure coevolu	IBEST, University of Idaho ution in continuous space			
2017	Paul Joyce Memorial BCB Fellowship Endowment Nominated by Professor Scott Nuismer because of my "love for mathematics and helping others to appreciate how it can be used to understand biological processes"				
2013-2015	Undergraduate Research in Biology & Mathematics Efforts focused on developing a statistical method to measure co	IBEST, University of Idaho evolution in metapopulations			
PROFESSIONAL	EXPERIENCE				
2024 - 2027	KiTE Postdoctoral Research Fellow Establishing theoretical foundations for the study of microbiome mediated trait dynamics				
2022 - 2024	Postdoctoral Research Fellow B.J.M. Bohannan Lab, University of Orego Extended evolutionary theory for traits jointly determined by host genotype and host microbiome				
2020 - 2022	Postdoctoral Researcher G.S. Bradburd Lab, Michigan State University Developed mathematical and computational approaches to understand coevolution in continuous space				
2018	Visiting Scientist P.J. CaraDonna Lab, Rocky Mountain Biological Laboratory Field ecology training on estimating floral abundance and phenology, recording plant-pollinator interactions and estimating percent cover				
TEACHING EXP	ERIENCE				
2017	Teaching Assistant University of Idaho, Department of Biological Sciences Taught the lab portion of a 300-level ecology and population biology course				
2012 - 2014	Mathematics Tutor Clark Community College, Mathematics Department Part-time work at tutoring center supporting students taking a wide-range of coursework				
PRESENTATION	s —				
2025	Quantitative Genetics of Microbiome Mediated Traits - Talk	ESEB 2025 - Barcelona, Spain			
2025	Quantitative Genetics of Microbiome Mediated Traits - Semina	Theoretical Biology Department - MPI, Plön			
2025	When is Microbial Rescue More or Less Effective than Genetic Rescue? - Poster	Evolutionary Rescue Workshop - MPI, Plön			
2025	Microbiome-Mediated Host Adaptation: A Niche Construction Approach - Poster	Concepts in Evolution Workshop - MPI, Plön			
2024	Host-Parasite Coevolution & Microbiome-Mediated Adaptation - Seminar	TransEvo Core Seminar - CAU, Kiel			
2023	The Evolution of Microbiome-Mediated Traits - Talk	Symbiosis Theory Workshop - Eugene, Oregon			
2023	Modeling Adaptation of Microbiome-Mediated Traits - Talk	EvoWibo - Port Townsend, Washington			
2022	Host-Parasite Coevolution in Continuous Space - Poster	PEQG2022 - Pacific Grove, California			
2021	Coevolutionary Arms Races and The Conditions for The Maintenance of Mutualism - Talk	AmNat2021 - Virtual			
2020	A Bayesian Methodology for Estimating the Distribution of Coevolution within Ecological Communities	- Talk AmNat2020 - Pacific Grove, California			
2018	The Measurement of Coevolution in Nature - Poster	EvoWibo - Port Townsend, Washington			
2017	The Measurement of Coevolution in Mutualisms - Talk	Evolution - Portland, Oregon			
SERVICE & LEAI	DERSHIP				
2022	Code Contributer Developed a nucleotide-based model of coevolution for SLiM. Se	SLIM 4.0 be 619.7 here doi:10.1086/723601			

Developed a nucleotide-based model of coevolution for SLiM. See §19.7 here.

doi:10.1086/723601

2018-2019

Graduate Student Representative
Represented graduate students in the Bioinformatics & Computational Biology program at institutional meetings

Manuscript Reviewer
The American Naturalist, Ecology, Evolution, PCI Evol Biol, Population Ecology, Proceedings of The Royal Society B, Theoretical Population Biology, Molecular Biology & Evolution

SOCIETIES

2021-Present The International Society of Nonbinary Scientists

isnbs.org

2020-Present The American Society of Naturalists

amnat.org

INTERESTS

I am broadly interested in collaborating on any scientific topic where my skills are useful. I am particularly interested in developing and formalizing models to clarify conceptual issues in population biology and community ecology.

SKILLS -

Software: LATEX, Python, R, Linux, Julia, Mathematica,

SLURM, SLIM, C/C++

Statistics: Modeling, Analysis, Inference, Methods

Development

Math: Linear Algebra, Dynamical Systems,

Functional Analysis, Stochastic Processes