Dr. Stephen J. Beckett

😐 🗠 Computational Ecology and Quantitative Viral Dynamics 🌬 🥐



About me

Computational ecologist with 10+ years of experience in scientific research spanning infectious disease dynamics, microbial ecology, and public health communication. I like interdisciplinary challenges, providing training and mentorship, and open-source coding practices. Current interests lie in investigating the ecology of aquatic microbial communities; and advancing the integration of infectious disease dynamics and human behaviour.

Areas of specialization

- · mathematical modeling and simulation · network analysis
- · interactive data dashboards
 - timeseries analysis · Bayesian inference

Research Interests

- · marine microbial ecology · virus-microbe ecology and
 - evolution biophysics epidemic dynamics
- network structure · dynamics of complex systems

Editorial Boards

- · PLOS Computational Biology.
- · Mathematics in Medical and Life Sciences.
- beckett@umd.edu
- sjbeckett.github.io
- 0000-0002-4410-2960
- p_YVeZ4AAAAJ
- sjbeckett
- in stephen-beckett
- Phage on toast
- Download CV.pdf

Short Resumé

2023-**Associate Research Scientist**

DEPARTMENT OF BIOLOGY & INSTITUTE FOR HEALTH COMPUTING · University of Maryland, College Park

Personalized COVID-19 vaccine recommendation website; cofounded the Chesapeake Aquatic Viral Ecology network, member of UMD Pandemic Readiness Initiative leading a wastewater surveillance and communication project.

2019-2023 Research Scientist I and II

School of Biology · Georgia Tech., Atlanta 🥊

Deadership for multiple COVID-19 response projects - including modeling transmission, risk, and prevalence: covid19risk website (>60M users); lead multitrophic ecological model-data integration and data analysis (20 coauthors); multiple art-science collaborations inc. showings at Atlanta Science Festival.

2015-2019 Postdoctoral Fellow

School of Biology · Georgia Tech., Atlanta 🥊

■ Marine microbial ecology - with a special interest in viral-host interactions; as well as the interplay of bottom-up (primarily resource driven) and top-down controls (viruses, grazers) on marine microbial communities. Organized GT's 2017 Postdoctoral Research Symposium.



DEGREES

2015 **Biological Sciences**

PнD · University of Exeter <u>ш</u>

Mathematics in the Living Environment 2011 MRES · University of York 🏛

2010 **Geography and Mathematics** BSc (Hons) · University of Leeds 🏛

UNIVERSITY OF LEEDS

Programming

julia MATLAB

python 🏶

FUNDING (TOTAL: >\$400,000)

2024 University of Maryland Grand Challenges

2022

2021 Rockefeller Foundation

SELECT PUBLICATIONS (>24 PUBS, >2500 CITATIONS, FULL LIST)

Disentangling top-down drivers of mortality underlying diel population dynamics of Prochlorococcus in the North Pacific Subtropical Gyre. Nat. Comms.

Real-time, interactive website for US-county-level 2020 COVID-19 event risk assessment. Nat. Hum. Behav.

2017 Lysis, lysogeny, and virus-microbe ratios. Nature.

Improved community detection in weighted bipartite 2016 networks. Roy. Soc. Op. Sci.

SELECT CURRICULUM DEVELOPMENT (MENTORED 4 UNDERGRADUATE AND 8 PHD STUDENT RESEARCHERS.)

Introduction to Python programming for life sciences

LEAD · University of Maryland, College Park

Fundamentals of coding using Python with a focus on applications in the life sciences. Students will learn fundamental coding and apply their knowledge through projects.

2025 Infectious disease dynamics: a systems approach

CO-LEAD · University of Maryland, College Park

Introduced systems thinking to understand complex disease interactions, and used interactive notebooks to run and analyze simulations of infectious disease models. 8 students.

Last updated: October 15, 2025