Bren Case

Infectious disease modeling • Bayesian statistics • Experimental design

PhD candidate in Computer Science combining mechanistic modeling and Bayesian statistics for the study of epidemiology and biology. I am especially interested in using these tools for informing how data should be collected, and for assessing control options from a decision-theoretic perspective.

Education

2018- Ph.D. Computer Science, University of Vermont, Burlington, Vermont.

- o Co-advisors: Laurent Hébert-Dufresne and Jean-Gabriel Young
- Anticipated completion: August 2023

2017–2019 MRes. Computer Science, University of Birmingham, Birmingham, UK.

- Thesis: Self-adaptation in non-elitist evolutionary algorithms: a rigorous analysis on discrete problems with unknown structure
- o Committee: Per Kristian Lehre (advisor), Thomas Jansen, Ata Kaban

2013–2017 B.A. Mathematics, Oberlin College, Oberlin, Ohio.

Minor: Computer Science

Publications

Microbial dysbiosis precedes signs of sea star wasting disease in wild populations of the Pycnopodia helianthoides

Andrew R. McCracken, Blair M. Christensen, Daniel Munteanu, **B. K. M. Case**, Melanie Lloyd, Kyle P. Herbert, Melissa H. Pespeni.

Frontiers in Marine Science. 2023. HTML PDF

Spatial epidemiology and adaptive targeted sampling to manage the Chagas disease vector Triatoma dimidiata

B. K. M. Case, Jean-Gabriel Young, Daniel Penados, Carlota Monroy, Laurent Hébert-Dufresne, Lori Stevens.

PLoS Neglected Tropical Diseases. 2022. HTML PDF

Flowers as dirty doorknobs: Deformed wing virus transmitted between Apis mellifera and Bombus impatiens through shared flowers

Phillip Alexander Burnham, Samantha Alger, **Brendan Case**, Humberto Boncristiani, Laurent Hébert-Dufresne, Alison Brody.

Journal of Applied Ecology. 2021. HTML

Self-adaptation in nonelitist evolutionary algorithms on discrete problems with unknown structure

Brendan Case and Per Kristian Lehre.

IEEE Transactions on Evolutionary Computation. 2020. HTML PDF

Submitted or under review

Accurately summarizing an outbreak using epidemiological models takes time

B. K. M. Case, Jean-Gabriel Young, Laurent Hébert-Dufresne.

Under review. Preprint: HTML PDF

The unintended consequences of inconsistent pandemic control policies

Benjamin Althouse, Brendan Wallace, **Brendan Case**, Samuel Scarpino, Antoine Allard, Andrew Berdahl, Easton White, Laurent Hébert-Dufresne.

Under review. Preprint: HTML PDF

Presentations

Conference talks

- Feb 2023 Adapting Survey Designs for Vector Surveillance Using Bayesian Decision Theory: An Application to an Ongoing Tick Monitoring Program in the Southeastern United States¹, National Big Data Health Science Conference, Columbia, South Carolina.
- Jun 2019 **Hidden geometry of infestation in Chagas disease vectors: an approach from epidemiological network theory**, Laboratorio de Entomología Aplicada y Parasitología Research Symposium, Guatemala City, Guatemala.
- May 2019 **Modeling disease spillover using multipartite networks**, *NetSci 2019*, Burlington, VT.
- Apr 2019 **Modeling disease spillover in bees: exploring dilution effects**, *UVM Student Research Conference*, Burlington, VT.

Conference posters

- Mar 2022 Parameter inference in epidemiological modeling: a perspective from Bayesian experimental design², NERCCS 2022: Fifth Northeast Regional Conference on Complex Systems, Buffalo, NY.
- Sep 2019 **QuEST timeline: highlights from the first year**, *NSF National Research Trainee-ship annual meeting*, Evanston, IL.

Teaching

Teaching Assistant

- Spring 2020 **Computability and Complexity**, *University of Vermont*.
 - Fall 2019 Modeling Complex Systems, University of Vermont.
- Spring 2018 **Software Workshop I**, *University of Birmingham*.
 - Fall 2017 Data Structures and Algorithms, University of Birmingham.
- Spring 2017 Foundations of Analysis, Oberlin College.
- Spring 2017 Algorithms, Oberlin College.
 - Fall 2016 Discrete Mathematics, Oberlin College.

Invited Workshops & Lectures

¹best presentation award

²best poster award

- 9/14 2022 Introduction to epidemiological models and disease forecasting, *EPID 394:* Infectious Disease Epidemiology, University of South Carolina.

 Slides
- 8/2 2022 Spatial epidemiology and adaptive targeted sampling to manage domestic Triatomine infestations in Guatemala, *UPenn-Tulane-UPCH Zoonotic Disease Research Lab*, University of Pennsylvania.

 Slides
- 6/23 2022 Introduction to tidy data and network science in R, Big Data Health Science Center T35 trainees, University of South Carolina.

 Notes
- 10/25 2021 **Bayesian Geostatistics and Adaptive Sampling**, *TGIR Adventures in Modeling*, University of Vermont.
- 8/16–8/23 **QuEST Coding Workshop for Incoming Trainees**, University of Vermont. 2021 Notes
- 11/15 2019 **The Rest of the Tidyverse**, *BIOL 381: Foundations of Quantitative Reasoning*, University of Vermont.

Professional Service and Leadership

- April 2023 Reviewer, Physical Review E.
- April 2023 Reviewer, Geospatial Statistics.
- Jan 2023 Reviewer, PLoS Computational Biology.
- Oct 2022 Reviewer, Frontiers in Ecology and Evolution.
- Apr 2022 Reviewer, Swarm and Evolutionary Computation.
- Apr 2022 Judge, Vermont Science Olympiad. Event: Experimental Design.
- Feb 2022 Reviewer, Physical Review E.
- Aug 2021 Reviewer, Swarm and Evolutionary Computation.

Advanced Schools & Workshops

- 4/10–4/14 **Multi-scale modeling of malaria**, *American Institute of Mathematics*, San Jose, 2023 California.
- 12/15–12/20 **Complex Networks Winter Workshop**, *Université Laval*, Quebec City, Canada. 2019
- 6/3–6/5 2019 **VectorBase Workshop**, *Universidad del Valle de Guatemala*, Guatemala City, Guatemala.

Scholarships

- 2022 **T35 Research Traineeship**, National Institute for Allergy and Infectious Diseases & University of South Carolina Big Data Health Science Center, award 5T35Al165252-02.
- 2018-2023 **QuEST National Research Traineeship**, *National Science Foundation & University of Vermont Graduate College*, award DGE-1735316.

2013-2014 Conservatory Dean's Scholarship, Oberlin College.

Selected Software

- Marginal Divergence.jl: a psuedo-Bayesian method for practical identifiability of differential equation models. (Julia)
- Conditional Sampling.jl: sampling from joint distributions conditional on variable transformations (Julia)
- Adaptive targeted sampling using R-INLA (R, RMarkdown)

Skills & Expertise

 $Programming \ \ R \ (tidyverse, \ tidygraph, \ sf/raster, \ caret), \ Julia \ (Differential Equations), \ Python$

languages (graph-tool)

Statistical Stan, R-INLA, nimble, Turing.jl

programming

Visualization ggplot2, ggraph, Inkscape

Community Service

2020-2022 Food Not Bombs, volunteer cook, Burlington, VT.

2019 ANEW Place, meal provider, Burlington, VT.

2017-2018 GoodGym, general member, Birmingham, UK.

2014-2015 Boys and Girls Club, tutor, Oberlin, OH.