BEN TEO

(+1) 608 960 2789 \$\display \text{bteo@wisc.edu} \$\display \text{https://github.com/bstkj}\$
Personal website: https://bstkj.github.io

EDUCATION

University of Wisconsin-Madison

Aug 2018 - Fall 2024

Ph.D. Dissertator in Statistics

Carnegie Mellon University

Aug 2013 - May 2017

B.S. Mathematics (Additional Minor in Computer Science)

RESEARCH INTERESTS

Statistical and computational techniques for scalable inference of trait models on phylogenetic networks.

PUBLICATIONS

Teo, B., Bastide, P., Ané, C. 2024. Leveraging graphical model techniques to study evolution on phylogenetic networks. *arXiv* preprint. https://doi.org/10.48550/arXiv.2405.09327.

Teo, B., Rose, J. P., Bastide, P., and Ané, C. 2023. Accounting for within-species variation in continuous trait evolution on a phylogenetic network. *Bulletin of the Society of Systematic Biologists*. https://doi.org/10.18061/bssb.v2i3.8977.

Tang, S., Lee, T. S., Li, M., Zhang, Y., Xu, Y., Liu, F., **Teo, B.**, and Jiang, H. 2018. Complex pattern selectivity in macaque primary visual cortex revealed by large-scale two-photon imaging. Current Biology, 28(1):38–48. https://doi.org/10.1016/j.cub.2017.11.039.

MANUSCRIPTS

Teo, B., Bacharach, M., Kolb, A., Ané, C., Roch, S., Brandt, C. Capabilities and limitations of explicit network methods for recombinant detection: a case study using Bovine Alphaherpesvirus 1. *In preparation*.

TALKS

Leveraging graphical model techniques to study evolution on phylogenetic networks. 3rd Joint Congress on Evolutionary Biology. Montreal, Canada. July 2024

Belief propagation for continuous trait evolution on phylogenetic networks. UW-Madison SGSA Student Seminar. Madison, WI. Nov 2022

Accounting for within-species variation in continuous trait evolution on a phylogenetic network. Evolution conference. Cleveland, OH. June 2022

SOFTWARE

Contributor to: PhyloGaussianBeliefProp.jl, PhyloNetworks.jl

HONORS AND AWARDS

SRGC Conference Presentation Award, UW-Madison Graduate School

Mayr Symposium, Finalist. 3rd Joint Congress on Evolutionary Biology

Statistics Poster Competition, 2nd place. CMU Undergraduate Research Symposium

May 2017

UW ACADEMIC EXPERIENCE

Teaching Assistant

Fall 2018 - Fall 2022

Stat 301/324/371: Introductory Applied Statistics for Social Sciences/Life Sciences/Engineers

Stat 349: Time Series, Stat 424: Experimental Design, Stat 479: Bayesian Analysis

Stat 850: Theory/Application of Regression and Analysis of Variance II

Graduate Student Mentor

Stat 699: Directed Study Fall 2021

Research Assistant

Statistics Department Spring 2023 – Spring 2024

Institute for Foundations of Data Science

Summer 2021

PAST PROJECTS

Sun Prairie Community Schools Data DashboardFall 2021Phylogenetic analysis of Algonquian using SNAPPSpring 2021Phylogenetic analysis of Algonquian using BEAST2Summer 2020

CMU ACADEMIC EXPERIENCE

Project Assistant: Modeling/clustering neuron response to parametrized visual stimuli

Center for Neural Basis of Cognition, Lee Lab

Summer 2017

Undergraduate Research: Hidden Markov Models to determine neuron population structure

Statistics Department

Summer 2016 – Spring 2017

WORK EXPERIENCE

Research Programmer

CMU, Computational Biology Department, Murphy Lab

Oct 2017 - July 2018

- Developer and Tester for CellOrganizer Project, a software package for learning generative models of cell/organelle shape/distribution from 2D/3D fluorescence micrographs.
- Worked with the OME-XML file format for reading and writing micrograph pixel/meta-data.
- Worked variously on segmentation related tasks for basal bodies and white blood cells.
- Compiled CellOrganizer source code for deployment, and built Docker container to export the compiled executables to other Docker-supported platforms.
- Created Bash tool scripts and XML tool definition files to enable CellOrganizer tools to be accessed through Galaxy Project's GUI.
- Linked CellOrganizer Project's GitLab repositories with Jenkins CI to enable automated testing.

TECHNICAL SKILLS

Programming Languages Julia, R, Python, MATLAB, C, Bash

Markup Languages Markdown, XML, LaTeX

Software & Tools Git, Docker, Jenkins CI, Galaxy Project, BEAST2