# Thanasi Bakis

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#### **EDUCATION**

University of California, Irvine, Ph.D., Statistics (Advisor: Volodymyr Minin)

2022-Present

University of California, Irvine, M.S., Statistics

2020-2022

University of California, Irvine, B.S., Data Science (summa cum laude)

2017-2020

Selected coursework:

- Bayesian Statistics (Stan) • Database Design (PostgreSQL)
- Generalized & Longitudinal Linear Models (R)

- Data Structures (C++)
- Deep Generative Models (PyTorch) Machine Learning (Python)

#### **Publications**

Bakis, A. G., & Minin, V. M. (In progress). Predictive variational inference for probabilistic forecasts of hospitalization time

- Developing a model-agnostic inference framework for probabilistic forecasting of time series, inspired by recent advances in variational inference.
- Applying the method to widely-used forecasting models, enabled by the ForwardDiff.jl automatic differentiation library in Julia.

Bakis, A. G., Vora, A. A., Araki, T., Jia, T., Galloway, J. G., Jennings-Shaffer, C., ... & Minin, V. M. (2025). Bayesian inference of antibody evolutionary dynamics using multitype branching processes. arXiv preprint arXiv:2508.09519.

- Implemented a novel branching process model in Julia to jointly analyze genetic sequence data from 3,700+ B cells sampled in an immunlogical experiment—the first application of Bayesian modeling to this large-scale dataset.
- Leveraged state-of-the-art packages Turing, il for Markov chain Monte Carlo-based inference and DifferentialEquations, il for numerical integration.
- Carefully optimized the memory allocation throughout my software to enable parallelization over 80 cores for >1 month of
- · Published all work as an open-source software library & a repository of data analyses to promote reproducibility and extensibility.

Ralph, D. K., Bakis, A. G., Galloway, J., Vora, A. A., Araki, T., Victora, G. D., ... & Matsen IV, F. A. (2025). Inference of germinal center evolutionary dynamics via simulation-based deep learning. arXiv preprint arXiv:2508.09871.

• Developed a deep learning model in TensorFlow for phylogenetic tree data, using the latest research to convert tree data structures to matrix inputs for the neural network without any loss of information.

### WORK EXPERIENCE

U.S. Centers for Disease Control and Prevention (CDC), Center for Forecasting and Outbreak Analytics

2024-2025

- Built a pipeline for the iterative development and evaluation of probabilistic forecasting models for the compositional prevalence of SARS-CoV-2 variants, using Python and Polars.
- Developed a collection of increasingly complex Bayesian multinomial regression models in NumPyro, using the pipeline to argue the merits of different model assumptions in various epidemiologically-relevant scenarios.

## Stats 170A/B, UC Irvine, Teaching Assistant

2021-2022

• Mentored 15 undergraduate data science teams on end-to-end capstone projects, from data acquisition (Python) and database design (PostgreSQL) to statistical modeling and visualization (R).

## iD Tech Camps, Instructor

2018

• Designed and taught an interactive cryptography curriculum in Python for middle and high school students.

### Honors and Achievements

### National Science Foundation, NSF Graduate Research Fellowship

2022

One of 14 Statistics students nationally to secure three years of Ph.D. research funding.

## Department of Statistics, UC Irvine, Newcombe Award

2021

Highest qualifying exam score across the Ph.D./M.S. cohort.

# CyberPatriot, Air & Space Forces Association, 9th place team nationally, of 2200

2017

Cybersecurity competition for Linux server hardening.