

HAMISH M. BLAIR

✉ hmlblair@stanford.edu  hmlblair  hmlblair.github.io  0009-0000-0091-2032

EDUCATION & RESEARCH EXPERIENCE

Stanford University

September 2022 - July 2027 (expected)

Ph.D. in Mathematics; advised by **Rhiju Das** (Biochemistry) and **George Papanicolaou** (Mathematics).

- Part of the *Ribonanza* initiative for training structure-informed RNA foundation models
- Advanced methods for extracting RNA structural information from large-scale experimental datasets.

University of Sydney

March 2018 - December 2021





Bachelor of Science and Bachelor of Advanced Studies, with majors in mathematics and physics. Applied mathematics honours supervised by **Leo Tzou**.

- Graduated with first class honours and the university medal
- Advanced theoretic results in computed tomography as part of the **applied geometric analysis group**
- Awarded the **Joye Medal** for greatest proficiency across the pure and applied mathematics honours programs.

PUBLICATIONS & THESES

1. Shujun He et al. **Ribonanza: deep learning of RNA structure through dual crowdsourcing**. *Under review at Nature Methods*.
2. **Hamish Blair**, Wipapat Kladwang, Rhiju Das. Prediction and reduction of read-depth bias in MaP-seq experiments. *In progress*.
3. **Hamish Blair**, Rhiju Das. Mutational profiling beyond the transcriptome scale with cmuts. *In progress*.
4. **Hamish Blair**, George Papanicolaou. Advances in sparse autocorrelation approaches to Cryo-EM. *In progress*.
5. **Hamish Blair**. Microlocal analysis and the geometry of distributions. *Honours thesis*.

SOFTWARE

-  **rn-filter** : fine-tuned foundation model to aid in the design of RNA mutational profiling experiments.
-  **cmuts** : modern multi-threaded RNA/DNA mutation counting software for use in machine learning pipelines.
-  **fld** : design of RNA mutational profiling libraries at the scale required for training foundation models.
-  **bioeq** : PyTorch-based software for training geometric deep learning models on biomolecular data, with custom CUDA kernels for reduced memory usage and up to a 10× inference speedup.

TEACHING & SERVICE

Stanford University, Course Assistant

- *Calculus I*; Fall 2022 (under Lernik Asserian)
- *Applied Matrix Theory*; Spring 2023 (under Gene Kim) and Fall 2024 (under Emmanuel Candès)
- *Fundamental Concepts of Analysis*; Fall 2023 (under Cole Graham)

Stanford University, Teaching Assistant

- *Linear Algebra, Multivariable Calculus, and Modern Applications*; Winter 2024 (under Mark Lucianovic) and Winter 2025 (under Christine Taylor)

Reviewer, *Nature Communications*

2024

Early Career Reviewer, *Science*

2025

SKILLS

Software Development:

- Proficiency in Python (5 years), MATLAB (3 years), C (2 years), and C++ (2 years).
- Experience with Mathematica (1 year), CUDA (1 year) and Lua (6 months).

Machine Learning & Data Science:

- Extensive experience with computational applied mathematics, especially numerical linear algebra (`numpy`, `scipy.linalg`, `cuBLAS`) and optimization (`cvxpy`, `scipy.optimize`) packages
- Three years' experience using PyTorch to train and fine-tune deep learning models
- Experience with multi-GPU fine-tuning and inference of foundation models via PyTorch Lightning.