

Timothy (Tim) Barry

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EMPLOYMENT

- | | |
|---|----------------------|
| Harvard University
Postdoctoral researcher, Department of Biostatistics
Advisor: Xihong Lin | 2024 – |
| University of Pennsylvania
Postdoctoral researcher, Department of Statistics
Advisor: Eugene Katsevich | July – December 2023 |

EDUCATION

- | | |
|---|-------------|
| Carnegie Mellon University (CMU)
PhD in Statistics
Advisors: Kathryn Roeder (CMU), Eugene Katsevich (University of Pennsylvania) | 2018 – 2023 |
| University of Maryland, College Park
BS in Mathematics with high honors
Minor in Computer Science | 2014 – 2018 |

AWARDS

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|---|------|
| • Harvard Chan Postdoctoral Association Travel Award | 2025 |
| • Howard Hughes Medical Institute Fellowship | 2017 |
| • Maryland Summer Scholars Research Grant | 2016 |
| • Banneker-Key Scholarship, University of Maryland's highest academic scholarship | 2014 |

PAPERS

- | | |
|---|------|
| • T Barry , Z Niu, E Katsevich, X Lin. "The permuted score test for robust differential expression analysis." Preprint. Link . | 2025 |
| • L Fischer, T Barry , A Ramdas. "Multiple testing with anytime-valid Monte-Carlo p-values." Preprint. Link . | 2024 |
| • T Barry , K Mason, E Katsevich, K Roeder. "Robust differential expression testing for single-cell CRISPR screens at low multiplicity of infection." <i>Genome Biology</i> . Link . (Mihaela Serban Memorial Award , American Statistical Association , Pittsburgh chapter) | 2024 |
| • T Barry , K Roeder, E Katsevich. "Exponential family measurement error models for single-cell CRISPR screens." <i>Biostatistics</i> . Link . | 2024 |
| • J Morris, C Caragine, Z Daniloski, J Domingo, T Barry , L Lu, K Davis, M Ziosi, D Glinos, S Hao, E Mimitou, P Smibert, K Roeder, E Katsevich, T Lappalainen, N Sanjana. "Discovery of target genes and pathways at GWAS loci by pooled single-cell CRISPR screens." <i>Science</i> . Link . | 2023 |
| • T Barry , X Wang, J Morris, K Roeder, E Katsevich. "SCEPTRE improves calibration and sensitivity in single-cell CRISPR screen analysis." <i>Genome Biology</i> . Link . (Reviewers' choice , American Society of Human Genetics conference) | 2021 |

- **T Barry***, E Gurarie*, F Cheraghi, I Kajola, W Fagan. “Does dispersal make the heart grow bolder? Avoidance of anthropogenic habitat elements across wolf life history.” *Animal Behaviour* 166. **Joint first authorship*. [Link](#). 2020
- **T Barry**. “Collections in R: Review and Proposal.” *The R Journal* 10.1. [Link](#). 2018

BOOK

T Barry, J Deutsch, E Katsevich. “Hands-on single-cell CRISPR screen analysis.” e-book. [Link](#) 2024

GRANT ACTIVITY

Under review

Name	Agency	Mechanism	Role	Requested funds	Dates
“Statistical advances in CRISPR profiling and screening”	NIH	K99/R00	PI	\$1,015,000	4/01/2026 - 3/30/2031

SOFTWARE PACKAGES

- **sceptre**: statistically rigorous and massively scalable single-cell CRISPR screen analysis. [Link](#).
(First package for single-cell CRISPR screen analysis endorsed by 10x Genomics, main commercial supplier of single-cell experimental kits.)
- **ondisc**: out-of-core and cluster-scale computing on single-cell data. [Link](#).

PROFESSIONAL SERVICE

- *Reviewer*, Annals of Applied Statistics, Biometrika, Frontiers in Genetics, Nature Biotechnology
- *Judge*, NESS student paper competition (2024), ENAR student poster competition (2025)

MENTORING

- Songcheng Dai (Computational Biology Masters student at CMU). Topic: algorithms, data structures, and software for large-scale single-cell data. 2021 - 2022

Updated June 2025.