

GILLIAN CHUEmail: gchu4@illinois.eduHomepage: <https://gillichu.github.io/>**EDUCATION**

PhD	Princeton University Department of Computer Science	Aug 2022 - May 2027 (Expected)
MS	University of Illinois at Urbana-Champaign Program in Ecology, Evolution and Conservation Thesis: Phylogenetic Placement Advisor: Professor Tandy Warnow	Dec 2020 - August 2022
BS	University of California, Berkeley Department of Computer Science	Aug 2016 - Dec 2020
HSD	Phillips Exeter Academy	Aug 2014 - May 2016

RESEARCH INTERESTS

Computational Biology, Computational/Statistical Genetics, Probabilistic Graphical Models.

My interests are at the intersection of computer science, statistics, and biology. I am interested in building tools that help us understand the evolution of complex and heterogeneous biological systems.

ACADEMIC HONORS AND FELLOWSHIPS

- RECOMB-CCB Scientific Communications (2022) 1st Place (*Awarded \$200*).
- Genetics Society of America: Presidential Membership Initiative (2022). *Awarded 1-year membership to GSA, Early Career Leadership Program and GENETICS Peer Review Training Program.*
- NSF GRFP (5 years), 2021: *Three-year annual stipend of \$34,000.*
- Excellent Graduate Student Instructor, UIUC. Spring 2021. *Introduction to Programming for Engineers and Scientists (CS101).*

INDUSTRY & RESEARCH EXPERIENCE

Research Assistant, Princeton University *Aug 2022 - Present*
Advisor: Ben Raphael

- Leveraging graph motifs to identify drugs with previously unrecognized targets

Research Assistant, University of Illinois at Urbana-Champaign *Jan 2021 - Present*
Advisor: Tandy Warnow

- Designed fast multiple sequence alignment method capable of aligning ultra-large datasets
- Designed fast and scalable phylogenetic placement methods

Research Assistant, University of Illinois at Urbana-Champaign *Jan 2021 – Dec 2021*
Advisor: Mohammed El-Kebir

- Designing interactive visual editor for copy number calls in bulk tumor cell data

Research Assistant, University of California Berkeley*Sept 2019 – May 2021*

Advisor: Priya Moorjani

- Implemented an efficient method of uncovering founder events in modern populations
- Designed an efficient and accurate local ancestry inference method

Research Assistant, University of California Berkeley*Sept 2019 – Dec 2020*

Advisor: Satish Rao

- Designed a distance-based phylogenetic tree inference algorithm

Databricks, San Francisco*May – Aug 2019***Software Engineering Intern**, Observability Team

- Implemented distributed tracing for performance analysis across microservice architecture

Researcher, Sperax*Apr 2018 – Oct 2021*

- Analyzed consensus protocols for distributed systems and implemented a test net
- Designed Decentralized Autonomous Organization (DAO) voting protocol and modeled token economics

TEACHING EXPERIENCE

University of Illinois at Urbana-Champaign

- **Intro to Programming for Engineers and Scientists**. CS101. Graduate Student Instructure, UIUC Department of Computer Science. Spr' 21. **Excellent Graduate Instructor Award**.

University of California, Berkeley

- **Bioinformatics Bootcamp**. Teaching Assistant, Center for Computational Biology. Aug' 20.
- **Operating Systems and System Programming**. CS162. Reader, UC Berkeley EECS. Su'20.
- **Efficient Algorithms and Intractable Problems**. CS170. Undergraduate Student Instructor, UC Berkeley EECS. Fa'18, Spr'19, Fa'19, Spr'20.
- **Discrete Mathematics and Probability**. CS70. Reader, UC Berkeley EECS. Fa'17, Spr'18, Su'18.
- **Building with Blockchain for Web 3.0**. Guest Lecturer, UC Berkeley IEOR. Spr'20.
- **Blockchain Fundamentals**. CS198. Lecturer, UC Berkeley. Spr'18, Fa'18.
- **Blockchain for Enterprise**. Guest Lecturer, UC Berkeley Haas Business. Spr'19. Fa'19.
- **Blockchain for Lawyers**. Guest Lecturer, UC Berkeley Boalt Law. Spr'18.
- **EdX Blockchain Fundamentals**. Course Advisor, UC Berkeley. Spr'18.

PUBLICATIONS

* indicates joint first-author

Journal Papers in Preparation

On fast, accurate local ancestry inference. **Chu, G.**, Nisonoff, H., Moorjani, P. *KernelMix is a modular method to perform local ancestry inference using discriminative classifiers and a conditional random field.*

Journal Papers in Submission

Chu, G., Warnow, T., “SCAMPP+FastTree: Improving Scalability for Likelihood-Based Phylogenetic Placement” bioRxiv doi: 10.1101/2022.05.23.493012.

Conference Papers

Lalani, Z.*, **Chu, G.***, Zaccaria, S., El-Kebir, M., “User-guided local and global copy-number segmentation for tumor sequencing data.” bioRxiv doi: 10.1101/2022.01.15.476457v1. RECOMB-CCB 2022.

Journal Papers

Park M, Ivanovic S, **Chu G**, Shen C, Warnow T. UPP2: Fast and Accurate Alignment of Datasets with Fragmentary Sequences, *Bioinformatics*, 2023; btad007.

Lalani Z*, **Chu G***, Hsu S, Kagawa S, Xiang M, et al. (2022) CNAViz: An interactive webtool for user-guided segmentation of tumor DNA sequencing data. *PLOS Computational Biology* 18(10): e1010614.

Tournebize, R., **Chu, G.**, & Moorjani, P. (2022). Reconstructing the history of founder events using genome-wide patterns of allele sharing across individuals. *PLoS Genetics*, 18(6), e1010243.

Workshop Papers

Y. Wang, Sun J., Wang, X., Wei, Y., Wu, H., **Chu, G.**, Yu, Z., “Sperax: An Approach to Defeat Long Range Attacks in Blockchains,” IEEE INFOCOM 2020 – IEEE Conference on Computer Communications Workshops (INFOCOM WKSHPS), Toronto, ON, Canada, 2020, pp. 574-579. doi: 10.1109/INFOCOMWKSHPS50562.2020.9163036.

PRESENTATIONS

Poster Presentation, “SCAMPP+FastTree: Improving Scalability for Likelihood-Based Phylogenetic Placement.” ISMB-LA, Oct 2022.

Conference Presentation, “User-guided local and global copy-number segmentation for tumor sequencing data.” RECOMB-CCB, May 2022.

Conference Presentation, “MGDrive: Mosquito Gene Drive Explorer: Landscape Clustering,” National Conference on Undergraduate Research, March 2020.

Retreat Presentation, “MGDrive: The Original Trilogy,” UC Berkeley Computational Biology Retreat, October 2018.

Conference Presentation, “A Technical Overview of Blockchain Development,” TiE Inflect Silicon Valley, April 2018.

COMMUNITY SERVICE

Shield the Bay

Co-Founder/Finance, Berkeley, March 2020 – Present

Berkeley ANova

Events Committee Chair, Berkeley, Sept 2016 – June 2018

SKILLS/LANGUAGES

Programming: Python, Java, C, Javascript, R, React, Redux, Solidity, Go, Jsonnet, Scala

Tools/Framework: HTML, Git, Django, Docker, AWS, Remix, CircleCI, Webpack, Jenkins, Kubernetes, Grafana

Genomics: samtools, bwa, GATK

REFERENCES

Dr. Tandy Warnow, Professor
Department of Computer Science
University of Illinois, Urbana-Champaign
Email: warnow@illinois.edu

Dr. Mohammed El-Kebir, Assistant Professor
Department of Computer Science
University of Illinois, Urbana-Champaign
Email: melkebir@illinois.edu

Dr. Priya Moorjani, Assistant Professor
Center for Computational Biology
University of California, Berkeley
Email: moorjani@berkeley.edu

Dr. Satish Rao, Professor
Electrical Engineering and Computer Science
University of California, Berkeley
Email: satishr@berkeley.edu

Dr. John Marshall, Assistant Professor
School of Public Health
University of California, Berkeley
Email: john.marshall@berkeley.edu

Dr. Jaspal Sandhu, Professor of Practice
School of Public Health
University of California, Berkeley
Email: jaspal@berkeley.edu