## **GILLIAN CHU** Email: gc3045@princeton.edu Homepage: https://gillichu.github.io/

#### **EDUCATION**

PhD	<b>Princeton University</b> Department of Computer Science Advisor: Benjamin J. Raphael	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**

- National Cancer Institute: SSACB (2024). Awarded Best Poster Runner-up. •
- William G. Bowen Merit Fellowship (2023). Internal Princeton per-department fellowship. •
- 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): Five-year fellowship, 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 Advisor: Benjamin J. Raphael

- Designed new statistic to prioritize drug candidates
- Designed new evolutionary model to describe CRISPR-based lineage tracing data and an • efficient algorithm for maximum likelihood phylogeny inference under this model

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

Designed fast multiple sequence alignment method capable of aligning ultra-large datasets •

Aug 2022 - Present

Jan 2021 - Present

• 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

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

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

## Standard Bounties, Consensys

# **Fullstack Software Engineering Intern**

• Built out RESTful API, and smart contract web application

## **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. ٠
- 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. •

*May* – *Aug* 2019

*Sept 2019 – May 2021* 

June 2018 – Aug 2018

## PUBLICATIONS

\* indicates joint first-author

## **Conference Papers**

2. Mai, U.\*, **Chu, G.\***, and Benjamin J. Raphael. "Maximum Likelihood Inference of Time-scaled Cell Lineage Trees with Mixed-type Missing Data." *bioRxiv* (2024) doi: 10.1101/2024.03.05.583638. RECOMB 2024.

1. 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**

4. **Chu, G.**, Warnow, T., "SCAMPP+FastTree: Improving Scalability for Likelihood-Based Phylogenetic Placement." *Bioinformatics Advances*, Volume 3, Issue 1, 2023, vbad008, <u>https://doi.org/10.1093/bioadv/vbad008</u>.

3. Park M, Ivanovic S, **Chu G**, Shen C, Warnow T. UPP2: Fast and Accurate Alignment of Datasets with Fragmentary Sequences, *Bioinformatics*, Volume 39, Issue 1, January 2023, btad007, <u>https://doi.org/10.1093/bioinformatics/btad007</u>.

2. 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, no. 10 (2022): e1010614. <u>https://doi.org/10.1371/journal.pcbi.1010614</u>

1. 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*, no. 6 (2022), 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**, "LAML: Lineage Analysis via Maximum Likelihood." National Cancer Institute (NCI): Spring School on Algorithmic Cancer Biology (SSACB), Apr 2024.

**Poster Presentation**, "SCAMPP+FastTree: Improving Scalability for Likelihood-Based Phylogenetic Placement." International Society for Computational Biology (ISCB) – Latin America, Oct 2022.

<u>Conference Presentation</u>, "User-guided local and global copy-number segmentation for tumor sequencing data." Research in Computational Molecular Biology (RECOMB) – Computational Cancer Biology (CCB), May 2022.

<u>Conference Presentation</u>, "MGDrive: Mosquito Gene Drive Explorer: Landscape Clustering," National Conference on Undergraduate Research, March 2020.

**<u>Retreat Presentation</u>**, "MGDrive: The Original Trilogy," UC Berkeley Computational Biology Retreat, October 2018.

<u>Conference Presentation</u>, "A Technical Overview of Blockchain Development," TiE Inflect Silicon Valley, April 2018.

#### **COMMUNITY SERVICE**

Computer Science Graduate Student Committee Member, Princeton, Fall 2022 - Present

<u>Shield the Bay</u> Co-Founder/Finance, Berkeley, March 2020 – June 2021

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. Ben Raphael,** Professor Department of Computer Science Princeton University Email: braphael@princeton.edu

**Dr. Tandy Warnow,** Professor Department of Computer Science

### Gillian Chu

University of Illinois, Urbana-Champaign Email: <u>warnow@illinois.edu</u>

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

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

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

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

**Dr. Jaspal Sandhu**, Professor of Practice School of Public Health University of California, Berkley Email: <u>jaspal@berkeley.edu</u>