# GILLIAN CHU

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## **EDUCATION**

PhD Princeton University Aug 2022 - May 2027 (Expected)

Department of Computer Science

MS University of Illinois at Urbana-Champaign Dec 2021 - June 2022 (Expected)

Program in Ecology, Evolution and Conservation

Thesis: Phylogenetic Placement Advisor: Professor Tandy Warnow

**BS** University of California, Berkeley Aug 2016 - Dec 2020

Department of Computer Science

**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 people interpret genetic data.

#### ACADEMIC HONORS AND FELLOWSHIPS

- RECOMB-CCB Scientific Communications (2022) 1st Place (Awarded \$200).
- Genetics Society of America: Presidential Membership Initiative (2022). Awarded 1year 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, Warnow Lab

Jan 2021 - Present

Advisor: Tandy Warnow

- Designing fast multiple sequence alignment method capable of aligning ultra-large datasets
- Improving protein multiple sequence alignments with new outlier detection method
- Studying the impact of query alignment methods on microbial abundance profiling
- Designing new metagenomics abundance profiling method

## Research Assistant, El-Kebir Lab

Jan 2021 - Dec 2021

Advisor: Mohammed El-Kebir

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

### Research Assistant, Lawrence Berkeley National Lab

Sept 2020 – Jan 2021

Advisor: Jessica Granderson

• Improved prediction of unusual peak energy-use events for smart building technologies

## 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

### Research Assistant, University of California Berkeley

Sept 2018 – Dec 2019

Advisor: John Marshall

• Optimized probabilistic gene drive model by redesigning movement kernel

#### 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

#### Consensys, San Francisco

*June – Aug 2018* 

**Software Engineering Intern**, Standard Bounties

• Built RESTful API, React is library and smart contract webapp using distributed file storage

## Office of Intellectual Property & Industry Research, Berkeley System Administrator

*May* 2016 – *Feb* 2017

Implemented and tested Apex web portal used by hundreds of researchers for patent process

#### TEACHING EXPERIENCE

#### **University of Illinois at Urbana-Champaign**

• <u>Intro to Programming for Engineers and Scientists.</u> 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.
- <u>Efficient Algorithms and Intractable Problems.</u> CS170. Undergraduate Student Instructor, UC Berkeley EECS. Fa'18, Spr'19, Fa'19, Spr'20.

• <u>Discrete Mathematics and Probability.</u> 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**

#### 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

Park, M., Ivanovic, S., Chu, G., Shen, C., Warnow, T., "UPP2: Fast and Accurate Alignment Estimation of Datasets with Fragmentary Sequences." bioRxiv doi: 10.1101/2022.02.26.482099

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.

#### **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**

Tournebize, R., **Chu, G.**, and Moorjani, P., "Inferring the History of Founder Events in Human Populations." bioRxiv doi: 10.1101/2020.09.07.286450. (*to appear in PLoS Genetics*).

#### 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**

<u>Conference Presentation</u>, "User-guided local and global copy-number segmentation for tumor sequencing data." RECOMB-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**

#### Shield the Bay

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

#### Berkelev 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

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### 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, Berkley Email: jaspal@berkeley.edu