#### GILLIAN CHU

Email: <u>gc3045@princeton.edu</u> Homepage: <u>https://gillichu.github.io/</u>

#### **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

- 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: 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
- Designing a new probabilistic model to study cell differentiation

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

Jan 2021 - Present

- 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

### **Standard Bounties**, Consensys

June 2018 – Aug 2018

**Fullstack Software Engineering Intern** 

• Built out RESTful API, and smart contract web application

### **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**

\* indicates joint first-author

## **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**

- 4. **Chu, G.**, Warnow, T., "SCAMPP+FastTree: Improving Scalability for Likelihood-Based Phylogenetic Placement." *Bioinformatics Advances*, Volume 3, Issue 1, 2023, vbad008, <a href="https://doi.org/10.1093/bioadv/vbad008">https://doi.org/10.1093/bioadv/vbad008</a>.
- 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, <a href="https://doi.org/10.1093/bioinformatics/btad007">https://doi.org/10.1093/bioinformatics/btad007</a>.
- 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. https://doi.org/10.1371/journal.pcbi.1010614
- 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**

<u>Poster Presentation</u>, "SCAMPP+FastTree: Improving Scalability for Likelihood-Based Phylogenetic Placement." ISCB-LA, Oct 2022.

<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.

**Retreat Presentation**, "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

## **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

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, Berkley Email: jaspal@berkeley.edu