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Education

2022 – Ph.D. Computer Science (Computational Biology Track)

COLUMBIA UNIVERSITY, 4.09/4.00

Advisor: Elham Azizi

2019 – 2020 M.S. Electrical Engineering and Computer Science

University of California, Berkeley, 4.00/4.00

Advisors: Yun Song, Kannan Ramchandran

Thesis: A Likelihood-based Deconvolution of Bulk Gene Expression Data Using Single-

cell References [T1].

2015 – 2019 B.A. Computer Science & Molecular and Cellular Biology (emph. Immunology)

University of California, Berkeley, 3.95/4.00

Graduated with Highest Distinction and EECS Department Honors.

Research Experience

Summer 2024 Machine Learning & NGS Intern. SANOFI

Advisor: Vikram Agarwal. Modeled the effect of mRNA sequences on *in vitro* transcription (IVT) efficiency. Took the project from processing raw Next Generation Sequencing (NGS) data to ultimately providing sequence designs for the next iteration of in-house lab experiments.

2021 – 2022 Research Engineer. YOSEF LAB, UC BERKELEY

Advisor: Nir Yosef. Developed open-source software for deep probabilistic modeling of single-cell omics data, scvi-tools [J2]. Contributed to novel research work concerning RNA velocity inference and meta-analysis of large-scale scRNA-seq datasets [W2, J1].

2019 – 2020 Graduate Student Researcher. SONG LAB, UC BERKELEY

Advisor: Yun Song. Developed a method for the deconvolution of bulk RNA-seq samples

using a single-cell RNA-seq reference [J3].

2018 – 2019 Undergraduate Student Researcher. BLISS LAB, UC BERKELEY

Advisor: Kannan Ramchandran. Developed a robust method for the federated learning

regime in the presence of adversaries [W3].

2016 – 2017 Research Assistant. Brem Lab, UC Berkeley

Advisor: Rachel Brem. Trained in experimental wet lab protocols concerning gene trans-

formations and knockouts in yeast.

Honors & Awards

2023	1st Place GSK.ai CausalBench Challenge [W1]
2020	Outstanding Graduate Student Instructor Award, UC Berkeley
2019	Graduation with Highest Distinction, UC Berkeley (equiv. summa cum laude)
2019	EECS Department Honors, UC Berkeley
2018	Jim and Donna Gray Scholarship, UC Berkeley
2017	Upsilon Pi Epsilon, UC Berkeley

Publications

Star symbol (*) denotes equal contributions as a co-first author.

JOURNAL ARTICLES

- [J1] Adam Gayoso, Philipp Weiler, Mohammad Lotfollahi, Dominik Klein, **Justin Hong**, Aaron Streets, Fabian J Theis, and Nir Yosef. "Deep generative modeling of transcriptional dynamics for RNA velocity analysis in single cells". In: *Nature Methods* (2023), pp. 1–10. [URL].
- [J2] Adam Gayoso*, Romain Lopez*, Galen Xing*, Pierre Boyeau, Valeh Valiollah Pour Amiri, **Justin Hong**, Katherine Wu, Michael Jayasuriya, Edouard Mehlman, Maxime Langevin, Yining Liu, Jules Samaran, Gabriel Misrachi, Achille Nazaret, Oscar Clivio, Chenling Xu, Tal Ashuach, Mariano Gabitto, Mohammad Lotfollahi, Valentine Svensson, Eduardo da Veiga Beltrame, Vitalii Kleshchevnikov, Carlos Talavera-López, Lior Pachter, Fabian J. Theis, Aaron Streets, Michael I. Jordan, Jeffrey Regier, and Nir Yosef. "A Python library for probabilistic analysis of single-cell omics data". In: *Nature Biotechnology* 40.2 (2022), pp. 163–166. [URL].
- [J3] Dan D Erdmann-Pham*, Jonathan Fischer*, **Justin Hong**, and Yun S Song. "Likelihood-based deconvolution of bulk gene expression data using single-cell references". In: *Genome Research* 31.10 (2021), pp. 1794–1806. [URL].

ARTICLES IN CONFERENCE PROCEEDINGS

[J1] Achille Nazaret*, **Justin Hong***, Elham Azizi, and David Blei. "Stable Differentiable Causal Discovery". In: *Proceedings of the 41st International Conference on Machine Learning (ICML)* (2024). [URL].

REFEREED WORKSHOP PAPERS

- [W1] Achille Nazaret and **Justin Hong**. "BetterBoost-Inference of Gene Regulatory Networks with Perturbation Data". In: *Machine Learning for Drug Discovery (MLDD) Workshop at ICLR*, *Oral Presentation* (2023). [URL]. 1st place submission of GSK.ai Causal-Bench Challenge.
- [W2] Pierre Boyeau*, **Justin Hong***, Adam Gayoso, Michael Jordan, Elham Azizi, and Nir Yosef. "Deep generative modeling for quantifying sample-level heterogeneity in single-cell omics". In: *Machine Learning in Computational Biology (MLCB)*, *Oral presentation* (2022). [URL].

[W3] Avishek Ghosh*, **Justin Hong***, Dong Yin, and Kannan Ramchandran. "Robust Federated Learning in a Heterogeneous Environment". In: *ICML Workshop on Privacy and Security in ML* (2019). [URL].

PREPRINTS

[P1] Ross M Giglio, Nicholas Hou, Adeya Wyatt, **Justin Hong**, Lingting Shi, Mathini Vaikunthan, Henry Fuchs, Jose Pomarino Nima, Seth W Malinowski, Keith L Ligon, et al. "A heterogeneous pharmaco-transcriptomic landscape induced by targeting a single oncogenic kinase". In: *bioRxiv* (2024). [URL].

[P2] Jiacheng Gu, ... **Justin Hong** ..., and Jellert T Gaublomme. "CRISPRmap: Sequencing-free optical pooled screens mapping multi-omic phenotypes in cells and tissue". In: *bioRxiv* (2023). [URL].

THESIS WORK

[T1] **Justin Hong**, Dan D Erdmann-Pham, Jonathan Fischer, and Yun S Song. "A Likelihood-based Deconvolution of Bulk Gene Expression Data Using Single-cell References". Master's Thesis. University of California, Berkeley, 2021. [URL].

Presentations

CONTRIBUTED TALKS

Jul 2024	Irving Institute for Cancer Dynamics Intensive Workshop
Dec 2023	Irving Institute for Cancer Dynamics Monthly Research Meeting
Nov 2022	Machine Learning in Computational Biology Workshop (MLCB), Oral Presentation
May 2022	CZI Assembling Tissue References Workshop
Dec 2021	UC Berkeley Computational Biology Skills Seminar
	Posters
Sep 2024	Single Cell Genomics
Sep 2024	Machine Learning in Computational Biology Workshop (MLCB)
Jul 2024	International Conference on Machine Learning (ICML)
Oct 2023	Single Cell Genomics
Jul 2023	ICML Workshop on Computational Biology

Software

2023	SDCD – A method for inferring causal graphs from labeled interventional data.
2022-2024	MrVI – A library for deep sample-level meta-analysis of single-cell omics data.
2021 - 2022	scvi-tools – A library for deep probabilistic analysis of single-cell omics data.
2019 – 2020	$\ensuremath{\mathtt{RNA}}\textsc{-Sieve}$ – A package for deconvolution of bulk RNA-seq data with single-cell RNA-seq references.

Industry Experience

2020 – 2021 Software Engineer. NURO

Developed infrastructure software for the evaluation and introspection of the autonomy

software stack. Mentored new hires and a summer intern.

Summer 2019 Software Engineer Intern. NURO

Developed software for remote large-scale bot fleet management. Co-inventor of a

patent relevant to this project.

Summer 2018 Software Engineer Intern. PALANTIR

Developed code editing and execution software within the Foundry platform.

Summer 2017 Software Engineer Intern. Affinity

Developed software for customer relationship management in the venture capital space.

Teaching Experience

2023 Teaching Assistant. COLUMBIA UNIVERSITY

Course Title: Probabilistic Models and Machine Learning (STCS 6701)

Responsibilities: Held office hours and recitations, developed and graded assignments.

2019–2020 Head Graduate Student Instructor. UC BERKELEY

Course Title: Probability and Random Processes (EE 126)

Responsibilities: Developed course content, assignments, and exams. Organized staff of over ten student instructors. Lectured as a substitute.

Coordinated the course transition to fully online during the COVID-19 pandemic.

2018–2019 Undergraduate Student Instructor. UC BERKELEY

Course Title: Probability and Random Processes (EE 126)

Responsibilities: Taught discussions sections, led office hours, created course content.

2016–2017 Undergraduate Student Instructor / Course Tutor. UC BERKELEY

Course Title: Structure and Interpretation of Computer Programs (CS 61A)

Responsibilities: Taught discussion sections, led office hours, graded exam content.

Journal and Conference Reviewing

Jul 2024 Machine Learning in Computational Biology Workshop (MLCB)
Nov 2023 Machine Learning in Computational Biology Workshop (MLCB)

Jul 2023 ICML Workshop on Computational Biology

Mentorship

Sp/Su 2024	Khushi Desai, Master's Student, Columbia University
Summer 2024	Léa Bohbot, IICD Alliance Program Summer Intern, École Polytechnique
Summer 2023	Tu Duyen Nguyen, IICD ALLIANCE PROGRAM SUMMER INTERN, ÉCOLE POLYTECHNIQUE