

Education

2023-27	B.A. Computer Science, B.A. Data Science*, Minor Bioengineering, <i>University of</i>
	California, Berkeley
2022	SAT Score: 1510/1600
2020-23	IB Diploma Higher level Math AA, Biology, Chemistry, International English Gym-
	nasium Sodermalm

Research Experience

2025-	ML Researcher, Song Lab, University of California, Berkeley.
2024	Bioinformatics/AI engineer, Generation Lab, Inc.
2023-25	Researcher and Bioinformatician, Conboy Lab, University of California, Berkeley.
2023	Researcher, Hallucinating Scaffolds team, iGEM at Berkeley. (Advisor: Chris Ander-
	son)
2023-	Immunoinformatics/ML Researcher, Borch Lab, Washington University in St. Louis.
	(Advisor: Nicholas Borcherding)
2023	NGS Method developer, Sahlin Group, SciLifeLab / Stockholm University. (Advisor:
	Kristoffer Sahlin)
2022-23	Researcher, Karlsson-Hedestam/Murrell Lab, Karolinska Institute. (Advisor: Ben-
	jamin Murrell)

Awards & Honors

2023	Finalist, National Swedish Research fair $(1/55)$
2023	Top 60, National Swedish Programming Olympiad Site
2022	Top 115, National Swedish Chemistry Olympiad (Fully Swedish)
2022	Double Admit to Karolinska Institute's summer research programs (1/5 admitted
	nationwide)
2022	Top school participant, Swedish Biology Olympiad (Fully Swedish)

^{*}With Domain Emphasis of Computational Methods in Molecular and Genomic Biology

Publications

Journal Articles

- J1. Yang, Qile, Safina, K. R., Nguyen, K. D. Q., Tuong, Z. K. & Borcherding, N. scRepertoire
 2: Enhanced and efficient toolkit for single-cell immune profiling. PLOS Computational
 Biology 21, 1–10. https://doi.org/10.1371/journal.pcbi.1012760 (June 2025).
 Media: ASCA, Chinese Software Developer Network [In Chinese], Jianshu [In Chinese]
 - BioRxiv preprint: https://doi.org/10.1101/2024.12.31.630854.
- J2. Yang Qile. APackOfTheClones: Visualization of clonal expansion with circle packing. Journal of Open Source Software 9, 6868. https://doi.org/10.21105/joss.06868 (2024).
 Media: R views (by Rstudio), R-bloggers, Zhihu [In Chinese], 360doc [In Chinese], Tencent [In Chinese].
- J3. Cruz, J. M., Yeung, H., Alzalzalee, R., Yang, Qile, Kabir, H., McDonough, S., Mei, X., Conboy, M. J. & Conboy, I. M. In old mice, Exercise Induces Inflammation and Fibrosis unless Alk5-inhibitor and Oxytocin are used. *Journal of Cellular Physiology* 240, e70054. https://onlinelibrary.wiley.com/doi/abs/10.1002/jcp.70054 (2025).

Working papers

- W1. Cruz, J. M., Alzalzalee, R., Yeung, H., Mahmood, Z., Yang, Qile, Morshedian, N., Conboy, M. J., Mazahery, A. R., Nevado, J. B. & Conboy, I. M. Plasma dilution after myocardial infarction rescues cardiac repair, heart performance, and promotes recovery of motor function and endurance in old mice Manuscript Under Review. 2025.
- W2. Borcherding, N., Sun, B., DeNardo, D., Mudd, P. A., Ellebedy, A., Yang, Qile & Brestoff, J. R. *Ibex: Deep autoencoders for single-cell BCR sequencing* Manuscript to be submitted. 2025. https://github.com/BorchLab/Ibex.

Presentations

Posters

- P1. Yang, Qile, Sahlin, K. & Murrell, B. KmerGMA: A seed-based approach for fast homology searching Swedish National Science Fair Finals (Södertälje, Sweden). Apr. 2023. https://qile0317.github.io/KmerGMA.jl/stable/.
- P2. Yang, Qile et al. Reimagining BCR Modeling: Small-Scale, Interpretable, Chemistry Informed Veridical Data Science for Biology (Berkeley, CA, USA). July 2025. https://www.eventbrite.com/e/veridical-data-science-for-biology-2025-tickets-1384456339179.
 - Upcoming.

Demonstrations & Tutorials

- D1. Yang, Qile & Murrell, B. Discovery of Novel Camelia Germline Immunoglobulin Alleles
 Karolinska Institute's Summer Research Program in Computational Biology (Stockholm,
 Sweden). Aug. 2022. https://github.com/Qile0317/SoFoCompBio22.
- D2. Yang, Qile et al. Final Presentation: Hallucinating Scaffolds Team <u>iGEM</u> at Berkeley All Hands 3 (Berkeley CA, USA). Dec. 2023.

Selected Media Coverage

2025	Association of Single Cell Analysis (ASCA), scRepertoire 2: Enhanced and effi-
	cient toolkit for single-cell immune profiling [J1]
2024	Chinese Software Developer Network, Immune repertoire analysis - scRepertoire [in
	Chinese [J1]
2023	R-views (by RStudio), May 2023: "Top 40" New CRAN Packages [J2]

Professional Organizations

2025-	Member, Gerontological Society of America (GSA)
2025-	Member, The Antibody Society
2025-	Member, Boston Protein Design and Modelling Club (BPDMC)
2025-	Member, Association of Single Cell Analysis (ASCA)
2025-	Member, Bioinformatics.org

Other Distinctions

2025	Top 20 Global Fellow, undisclosed BioTech/AI entrepreneurship program; eligible for
	mid six-figure conditional funding ($< 0.1\%$ acceptance rate)
2023	PostHS scholarship recipient, Equitable College Counselling
2023	Swedish National Debate Team member, Swedish Schools Debating Organization
2023	World Schools Debating Championship Sponsorship (Withdrew due to external
	constraints), The Scholarship Fund in Memory of Ellinor Carlsson [In Swedish]

Relevant Coursework

Computational Biology: Introduction to Computational Molecular and Cell Biology | Machine Learning, Statistical Models, and Optimization for Molecular Problems

Machine Learning / Statistics: Introduction to Artificial Intelligence | Principle & techniques of Data Science | Human Contexts and Ethics of Data

 $\textbf{Biology / Chemistry:} \ \ \text{Chemical Structure and Reactivity | Organic Chemistry Laboratory | Introduction to Bioengineering | Careers in Biotechnology}$

Mathematics: Linear Algebra and Differential Equations | Multi-variable Calculus | Discrete Mathematics and Probability Theory | Introduction to Probability and Statistics

 $\textbf{Computer Science:} \ \, \textbf{Structure and Interpretation of Computer Programs} \mid \textbf{Data Structures} \mid \textbf{Interpretation to Competitive Programming}$

Last updated: July 6, 2025