Qile Yang | Curriculum Vitae qile [dot] yang [at] berkeley [dot] edu

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 projects)
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 $(1/5 \text{ admitted})$
	nationwide)
2022	Top school participant, Swedish Biology Olympiad (Fully Swedish)
2023	PostHS scholarship recipient, Equitable College Counselling
2022-23	Swedish National Debate Team member (5 national debating awards)

^{*}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 2023 Finals (Södertälje, Sweden). Apr. 2023. https://qile0317.github.io/KmerGMA.jl/stable/.

Demonstrations & Tutorials

- D1. Yang, Qile & Murrell, B. Discovery of Novel Camelid Germline Immunoglobulin Alleles
 Karolinska Institute's summer research program (SoFo) 2022 (Stockholm, Sweden). Aug.
 2022. https://github.com/Qile0317/SoFoCompBio22/blob/main/FinalPresentation.
 pptx.
- D2. Yang, Qile & et al. Final Presentation: Hallucinating Scaffolds Team iGEM at Berkeley All Hands 3 (Berkeley CA, USA). Dec. 2023.

Selected Media

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]

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

Biology / Chemistry: 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

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

Last updated: July 1, 2025