

# Charles Comiter

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CONTACT              *Email:* ccomiter@mit.edu  
INFORMATION        *Linkedin:* linkedin.com/in/charles-comiter/  
                             *Website:* ccomiter1.github.io

RESEARCH            **Areas:** Machine Learning for Biomedicine, Computational Biology  
INTERESTS           **Topics:** Multimodal Learning, -omics, Medical Imaging, Natural Language  
                             **Applications:** Pathology, Cardiology, Clinical Decision Support

EDUCATION           **Massachusetts Institute of Technology**, Cambridge, Massachusetts, USA  
                             Department of Electrical Engineering and Computer Science  
                             Ph.D., Computer Science, September 2021 – September 2026  
                             Received S.M. in Computer Science in May 2024  
                             Advisors: Collin M. Stultz and Jian Shu  
                             Collaborators: Aviv Regev, Charles M. Rudin, K. Kobayashi, and J. Klughammer  
                             Affiliations: MGH, Broad Institute, Human Cell Atlas, RLE, CSAIL  
                             Thesis Topic: Developing data driven AI models that generate clinically useful information—such as radiology reports or molecular profiles—from more accessible sources, such as histology images and EKGs

**Yale University**, New Haven, Connecticut, USA  
B.S., Computer Science, with Distinction, August 2016 – May 2020  
Significant coursework in Music and Statistics  
Senior Thesis: "Special Norms of the Laplacian Matrix and their Approximation",  
advised by Dan Spielman

PUBLICATIONS        Charles Comiter, Xingjian Chen, Eeshit Dhaval Vaishnav, Koseki J. Kobayashi-Kirschvink, Metamia Ciampricotti, Ke Zhang, Jason Murray, Francesco Monticolo, Jianhuan Qi, Ryota Tanaka, Sonia E. Brodowska, Bo Li, Yiming Yang, Scott J. Rodig, Angeliki Karatza, Alvaro Quintanal Villalonga, Madison Turner, Kathleen L. Pfaff, Judit Jane-Valbuena, Michal Slyper, Julia Waldman, Sebastian Vigneau, Jingyi Wu, Timothy R. Blosser, Asa Segerstolpe, Daniel L. Abravanel, Nikhil Wagle, Shadmehr Demehri, Xiaowei Zhuang, Charles M. Rudin, Johanna Klughammer, Orit Rozenblatt-Rosen, Collin M. Stultz, Jian Shu, Aviv Regev. Inference of single cell profiles from histology stains with the Single Cell omics from Histology Analysis Framework (SCHAF). *In Revision/Submission to Nature*. 2025

Koseki J. Kobayashi-Kirschvink, Charles Comiter, Shreya Gaddam, Taylor Joren, Emanuelle I Grody, Johain R Ounadjela, Ke Zhang, Baoliang Ge, Jeon Woong Kang, Ramnik J Xavier, Peter T.C. So, Tommaso Biancalani, Jian Shu, Aviv Regev. Prediction of single-cell RNA expression profiles in live cells by Raman microscopy with Raman2RNA. *Nature Biotechnology*. 2024

Yun Zhang, Jeremy A Miller, Jeongbin Park, Boudewijn P Lelieveldt, Brian Long, Tamim Abdelaal, Brian D Aevertmann, Tommaso Biancalani, [Charles Comiter](#), Oleh Dzyubachyk, Jeroen Eggermont, Christoffer Mattsson Langseth, Viktor Petukhov, Gabriele Scalia, Eeshit Dhaval Vaishnav, Yilin Zhao, Ed S Lein, Richard H Scheuermann. Reference-based cell type matching of in situ image-based spatial transcriptomics data on primary visual cortex of mouse brain. *Nature Scientific Reports*. 2023

Brian Long, Jeremy Miller, The SpaceTx Consortium (including [Charles Comiter](#)). SpaceTx: a roadmap for benchmarking spatial transcriptomics exploration of the brain. *arXiv*. 2023

PROFESSIONAL  
EXPERIENCE

**Broad Institute of MIT and Harvard**, Cambridge, Massachusetts, USA  
***Research Intern, Regev Lab*** **June 2020 – September 2021**  
Development of deep learning models for modality transfer and domain integration in biological imaging and single-cell omics. Collaborations include the Human Tumor Atlas Pilot Project, Human Cell Atlas, and Memorial Sloan Kettering Cancer Center.

**Facebook**, New York, New York, USA  
***Software Engineer*** **September 2020 – September 2021**  
Member of the Responsible Artificial Intelligence / Fairness in AI team. Wrote statistical and infrastructure software to evaluate the fairness of Facebook’s AI systems.

**Facebook**, New York, New York, USA  
***Software Engineer Intern*** **August – November 2019**  
ML Infrastructure for Messenger team. Designed and implemented a domain-specific language and companion compiler for AI feature engineering. Received highest possible performance review.

**Dropbox**, San Francisco, California, USA  
***Software Engineer Intern*** **May – August 2019**  
Filesystem Infrastructure team. Engineered cache implementations for file access/permissions metadata for major latency improvements.

**Accenture**, Chicago, Illinois, USA  
***Technology Analyst Intern*** **June – August 2018**  
Fullstack development and rapid prototyping of web applications for clients.

PATENTS

**Live-cell label-free prediction of single-cell omics profiles by microscopy**. U.S. Patent Application US20240371184A1; International Publication WO2023091970A1 (PCT/US2022/079989). [Charles Comiter](#), Koseki J. Kobayashi-Kirschvink, Tommaso Biancalani, Peter T.C. So, Jian Shu, Aviv Regev. Filed November 16, 2022. Published May 25, 2023 (WO); November 21, 2024 (US)

HONORS AND  
AWARDS

Siebel Scholar Fellowship, November 2023  
EECS Departmental Fellowship, September 2021

INVITED TALKS	<u>Inference of Single Cell Profiles from Histology Stains with SCHAF</u> , University of Hong Kong Blood Engineering Lab, April 2023
	<u>Inference of Single Cell Profiles from Histology Stains with SCHAF</u> , La Jolla Institute for Immunology, April 2023
	<u>Inference of Single Cell Profiles from Histology Stains with SCHAF</u> , Models, Inference, and Algorithms, Broad Institute of MIT and Harvard, March 2023
	<u>Inference of Single Cell Profiles from Histology Stains with SCHAF</u> , Klarman Cell Observatory Scientific Advisory Board, Broad Institute of MIT and Harvard, June 2021
POSTER PRESENTATIONS	Xingjian Chen, <u>Charles Comiter</u> , Collin M. Stultz, Jian Shu. Generation of cellular images from single-cell gene expression with GenVinci, MIT-MGB AI Cures Conference, Cambridge, MA, April 2024
	<u>Charles Comiter</u> , Koseki J. Kobayashi-Kirschvink, Aviv Regev, Jian Shu. ImageOmicsNet: Linking Images and Genomics with Machine Learning, MIT-MGB AI Cures Conference, Cambridge, MA, April 2023
TEACHING	Graduate Teaching Assistant: <u>AI and Decision Making in Medicine: From Disease to Therapy</u> , MIT, Fall 2025
	Graduate Teaching Assistant: <u>Computational Tools for MD/PhD Students</u> , MIT, January 2025
	Graduate Teaching Assistant: <u>Quantitative Methods for Natural Language Processing</u> , MIT, Fall 2022
	Undergraduate Learning Assistant: <u>Introductions to Systems Programming and Computer Organization</u> , Yale, Spring 2020
	Undergraduate Learning Assistant: <u>Mathematical Tools for Computer Science</u> , Yale, Fall 2018
	Mentorship: Mentoring Undergraduate and Early Graduate Students in the Stultz and Shu Laboratories at MIT
SERVICE	Reviewer: <i>Nature Genetics</i> , 2025 Reviewer: <i>Nature Biotechnology</i> , 2024 Reviewer: <i>Nature Methods</i> , 2023 - 2024
SKILLS AND MISCELLANY	Technical: Python, PyTorch, NumPy, Scanpy, Pandas, C, GoLang, Bash, LaTeX
	Other: Spanish (Working Proficiency), Yiddish (Extremely Elementary), Classical Percussion (Specialty in Four Mallet Marimba, First Prize in the 2014 Great Plains International Marimba and 2015 New World Symphony Side by Side Masterclass Competitions, Percussionist in Yale Symphony Orchestra and adjacent ensembles)
	Miscellany: Creator of <i>Yale Memes for Special Snowflake Teens</i> (Featured on NPR), National Merit Scholarship Semi-Finalist (2016), Dreyfoos Arts High School Salutatorian