
Michael Sellers Cuoco

PhD Student, Bioinformatics and Systems Biology

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Research interests Retrotransposon activity in the developing, aging, and diseased human brain.

Education

University of California, San Diego	La Jolla, California
PhD in Bioinformatics and Systems Biology	In Progress
Advised by Rusty Gage and Eran Mukamel	
Trinity College	Hartford, Connecticut
BS in Molecular and Cellular Biology	May 2016
Minor in Models and Data	

Honors and Awards

NSF Graduate Research Fellowship	2022
<i>National Science Foundation (NSF)</i>	
Spot Award	2017
<i>Broad Institute</i>	
Beta Beta Beta National Biology Honors Society	2014
<i>Trinity College</i>	
NESCAC Winter All-Academic Team	2014
<i>Trinity College</i>	

Research experience

PhD Student	2020 – Present
<i>UC San Diego, Salk Institute</i>	La Jolla, California
Mentors: Rusty Gage and Eran Mukamel	
Research Associate	2016 – 2020
<i>Broad Institute</i>	Cambridge, Massachusetts
Mentors: Aviv Regev, Benjamin Izar, Pratiksha Thakore, Yaara Oren	
Undergraduate Researcher	2014 – 2016
<i>Dana-Farber Cancer Institute</i>	Boston, Massachusetts
Mentors: Matthew Meyerson and Alison Taylor	
Undergraduate Researcher	2013
<i>Trinity College</i>	Hartford, Connecticut
HHMI Science Education Alliance-Phage Hunters Advancing Genomics and Evolutionary Science program. (SEA-PHAGES: seaphages.org)	

Research: Published

- Otto, J. E., Ursu, O., Wu, A. P., Winter, E. B., **Cuoco, M. S.**, Ma, S., Qian, K., Michel, B. C., Buenrostro, J. D., Berger, B., Regev, A., Kadoch, C., “Structural and functional properties of mSWI/SNF chromatin remodeling complexes revealed through single-cell perturbation screens.” In: *Molecular cell* (Apr. 2023). DOI: 10.1016/j.molcel.2023.03.013.
- Eraslan, G., Drokhyansky, E., Anand, S., Fiskin, E., Subramanian, A., Slyper, M., Wang, J., Wittenberghe, N. V., Rouhana, J. M., Waldman, J., Ashenberg, O., Lek, M., Dionne, D., Win, T. S., **Cuoco, M. S.**, Kuksenko, O., Tsankov, A. M., Branton, P. A., Marshall, J. L., Greka, A., Getz, G., Segrè, A. V., Aguet, F., Rozenblatt-Rosen, O., Ardlie, K. G., Regev, A., “Single-nucleus cross-tissue molecular reference maps toward understanding disease gene function.” In: *Science (New York, N.Y.)* 376 (6594 May 14, 2022). DOI: 10.1126/science.abl4290.
- Hwang, W. L., Jagadeesh, K. A., Guo, J. A., Hoffman, H. I., Yadollahpour, P., Reeves, J. W., Mohan, R., Drokhyansky, E., Wittenberghe, N. V., Ashenberg, O., Farhi, S. L., Schapiro, D., Divakar, P., Miller, E., Zollinger, D. R., Eng, G., Schenkel, J. M., Su, J., Shiau, C., Yu, P., Freed-Pastor, W. A., Abbondanza, D., Mehta, A., Gould, J., Lambden, C., Porter, C. B. M., Tsankov, A., Dionne, D., Waldman, J., **Cuoco, M. S.**, Nguyen, L., Delorey, T., Phillips, D., Barth, J. L., Kem, M., Rodrigues, C., Ciprani, D., Roldan, J., Zelga, P., Jorgji, V., Chen, J. H., Ely, Z., Zhao, D., Fuhrman, K., Fropf, R., Beechem, J. M., Loeffler, J. S., Ryan, D. P., Weekes, C. D., Ferrone, C. R., Qadan, M., Aryee, M. J., Jain, R. K., Neuberg, D. S., Wo, J. Y., Hong, T. S., Xavier, R., Aguirre, A. J., Rozenblatt-Rosen, O., Mino-Kenudson, M., Castillo, C. F.-D., Liss, A. S., Ting, D. T., Jacks, T., Regev, A., “Single-nucleus and spatial transcriptome profiling of pancreatic cancer identifies multicellular dynamics associated with neoadjuvant treatment.” In: *Nature genetics* 54 (8 July 29, 2022). DOI: 10.1038/s41588-022-01134-8.
- Li, J., Pinto-Duarte, A., Zander, M., **Cuoco, M. S.**, Lai, C.-Y., Osteen, J., Fang, L., Luo, C., Lucero, J. D., Gomez-Castanon, R., Nery, J. R., Silva-Garcia, I., Pang, Y., Sejnowski, T. J., Powell, S. B., Ecker, J. R., Mukamel, E. A., Behrens, M. M., “Dnmt3a knockout in excitatory neurons impairs postnatal synapse maturation and increases the repressive histone modification H3K27me3.” In: *eLife* 11 (May 24, 2022). DOI: 10.7554/eLife.66909.
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- Frangieh, C. J., Melms, J. C., Thakore, P. I., Geiger-Schuller, K. R., Ho, P., Luoma, A. M., Cleary, B., Jerby-Arnon, L., Malu, S., **Cuoco, M. S.**, Zhao, M., Ager, C. R., Rogava, M., Hovey, L., Rotem, A., Bernatchez, C., Wucherpfennig, K. W., Johnson, B. E., Rozenblatt-Rosen, O., Schadendorf, D., Regev, A., Izar, B., “Multimodal pooled Perturb-CITE-seq screens in patient models define mechanisms of cancer immune evasion.” In: *Nature genetics* 53 (3 Mar. 2021). DOI: 10.1038/s41588-021-00779-1.
- He, M. X., **Cuoco, M. S.**, Crowdis, J., Bosma-Moody, A., Zhang, Z., Bi, K., Kanodia, A., Su, M.-J., Ku, S.-Y., Garcia, M. M., Sweet, A. R., Rodman, C., DelloStritto, L., Silver, R., Steinharter, J., Shah, P., Izar, B., Walk, N. C., Burke, K. P., Bakouny, Z., Tewari, A. K., Liu, D., Camp, S. Y., Vokes, N. I., Salari, K., Park, J., Vigneau, S., Fong, L., Russo, J. W., Yuan, X., Balk, S. P., Beltran, H., Rozenblatt-Rosen, O., Regev, A., Rotem, A., Taplin, M.-E., Allen, E. M. V., “Transcriptional mediators of treatment resistance in lethal prostate cancer.” In: *Nature medicine* 27 (3 Mar. 2021). DOI: 10.1038/s41591-021-01244-6.
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- Izar, B., Tirosh, I., Stover, E. H., Wakiro, I., **Cuoco, M. S.**, Alter, I., Rodman, C., Leeson, R., Su, M.-J., Shah, P., Iwanicki, M., Walker, S. R., Kanodia, A., Melms, J. C., Mei, S., Lin, J.-R., Porter, C. B. M.,

- Slyper, M., Waldman, J., Jerby-Arnon, L., Ashenberg, O., Brinker, T. J., Mills, C., Rogava, M., Vigneau, S., Sorger, P. K., Garraway, L. A., Konstantinopoulos, P. A., Liu, J. F., Matulonis, U., Johnson, B. E., Rozenblatt-Rosen, O., Rotem, A., Regev, A., “A single-cell landscape of high-grade serous ovarian cancer.” In: *Nature medicine* 26 (8 June 24, 2020). DOI: 10.1038/s41591-020-0926-0.
- Kinker, G. S., Greenwald, A. C., Tal, R., Orlova, Z., **Cuoco, M. S.**, McFarland, J. M., Warren, A., Rodman, C., Roth, J. A., Bender, S. A., Kumar, B., Rocco, J. W., Fernandes, P. A. C. M., Mader, C. C., Keren-Shaul, H., Plotnikov, A., Barr, H., Tsherniak, A., Rozenblatt-Rosen, O., Krizhanovsky, V., Puram, S. V., Regev, A., Tirosh, I., “Pan-cancer single-cell RNA-seq identifies recurring programs of cellular heterogeneity.” In: *Nature genetics* 52 (11 Nov. 2020). DOI: 10.1038/s41588-020-00726-6.
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- Li, A., Herbst, R. H., Canner, D., Schenkel, J. M., Smith, O. C., Kim, J. Y., Hillman, M., Bhutkar, A., **Cuoco, M. S.**, Rappazzo, C. G., Rogers, P., Dang, C., Jerby-Arnon, L., Rozenblatt-Rosen, O., Cong, L., Birnbaum, M., Regev, A., Jacks, T., “IL-33 Signaling Alters Regulatory T Cell Diversity in Support of Tumor Development.” In: *Cell reports* 29 (10 Dec. 2019). DOI: 10.1016/j.celrep.2019.10.120.
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Research: Preprint

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Teaching / Mentorship	Undergraduate mentor	2021 – Present
	<i>UCSD Biology Undergraduate and Master’s</i>	La Jolla, California
	Bootcamp instructor	Fall 2021, Fall 2022
	<i>Bioinformatics and Systems Biology, UCSD</i>	La Jolla, California
	Teaching assistant	Spring 2015
Service / Outreach	<i>Department of Biology, Trinity College</i>	Hartford, Connecticut
	BIOL 224: Genetics	
	Tutor	2014 – 2016
	<i>Department of Biology, Trinity College</i>	Hartford, Connecticut
	BIOL 182: Evolution of Life	
	BIOL 183: Cellular Basis of Life	
	BIOL 224: Genetics	
	Committee Member	2021 – Present
	<i>Advisory Committee on Diversity</i>	La Jolla, California
	<i>Salk Institute for Biological Studies</i>	
	Director of Onboarding	2021 – Present
	Symposium Organizer	2022
	<i>Graduate Bioinformatics Council</i>	La Jolla, California
	<i>UCSD Bioinformatics and Systems Biology</i>	
	Committee Member	2020 – Present
	<i>Diversity Equity and Inclusion Committee</i>	La Jolla, California
	<i>UCSD Bioinformatics and Systems Biology</i>	
	Seminar Organizer	2021
	Symposium Organizer	Fall 2021
	<i>Diversity and Science Lecture Series</i>	La Jolla, California
	<i>UCSD</i>	
	Volunteer - High Tech High Mesa	Fall 2021
	Volunteer - La Jolla High School	Fall 2021
	<i>SciChats@Salk Education Outreach</i>	La Jolla, California
	<i>Salk Institute for Biological Studies</i>	

Programming Languages

R, Python, Bash

Data Analysis

Single-cell genomics: Seurat, scanpy, pegasus

Pipeline development: Workflow development language (WDL), Snake-make

Job managers: Cromwell, Sun Grid Engine (SGE), Slurm, PBS-Torque

Cloud computing: Google Cloud Platform (GCP), Terra

Visualization: ggplot, matplotlib

Programmatic Reporting

Notebooks / Slides: Quarto, Rmarkdown, Jupyter Notebooks

Websites: Jekyll, Bookdown, Blogdown, Jupyter Book

Software Development

Git, GitHub, GitHub Actions CI/CD