
Michael Sellers Cuoco

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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
National Science Foundation (NSF)
Spot Award 2017
Broad Institute
Beta Beta Beta National Biology Honors Society 2014
Trinity College
NESCAC Winter All-Academic Team 2014
Trinity College

Research experience

PhD Student 2020 – Present
UC San Diego, Salk Institute La Jolla, California
Mentors: Rusty Gage and Eran Mukamel

Research Associate 2016 – 2020
Broad Institute Cambridge, Massachusetts
Mentors: Aviv Regev, Benjamin Izar, Pratiksha Thakore, Yaara Oren

Undergraduate Researcher 2014 – 2016
Dana-Farber Cancer Institute Boston, Massachusetts
Mentors: Matthew Meyerson, Alison Taylor

Undergraduate Researcher 2013
Trinity College Hartford, Connecticut
HHMI Science Education Alliance-Phage Hunters Advancing Genomics and Evolutionary Science program. (SEA-PHAGES: seaphages.org)

Research: Published

- Eraslan, G., Drokhylyansky, 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.
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- 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.
- Bi, K., He, M. X., Bakouny, Z., Kanodia, A., Napolitano, S., Wu, J., Grimaldi, G., Braun, D. A., **Cuoco, M. S.**, Mayorga, A., DelloStritto, L., Bouchard, G., Steinharter, J., Tewari, A. K., Vokes, N. I., Shannon, E., Sun, M., Park, J., Chang, S. L., McGregor, B. A., Haq, R., Denize, T., Signoretti, S., Guerriero, J. L., Vigneau, S., Rozenblatt-Rosen, O., Rotem, A., Regev, A., Choueiri, T. K., Allen, E. M. V., “Tumor and immune reprogramming during immunotherapy in advanced renal cell carcinoma.” In: *Cancer cell* 39 (5 Mar. 13, 2021). DOI: 10.1016/j.ccell.2021.02.015.
- 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.
- Jerby-Arnon, L., Neftel, C., Shore, M. E., Weisman, H. R., Mathewson, N. D., McBride, M. J., Haas, B., Izar, B., Volorio, A., Boulay, G., Cironi, L., Richman, A. R., Broye, L. C., Gurski, J. M., Luo, C. C., Mylvaganam, R., Nguyen, L., Mei, S., Melms, J. C., Georgescu, C., Cohen, O., Buendia-Buendia, J. E., Segerstolpe, A., Sud, M., **Cuoco, M. S.**, Labes, D., Gritsch, S., Zollinger, D. R., Ortogero, N., Beechem, J. M., Nielsen, G. P., Chebib, I., Nguyen-Ngoc, T., Montemurro, M., Cote, G. M., Choy, E., Letovanec, I., Cherix, S., Wagle, N., Sorger, P. K., Haynes, A. B., Mullen, J. T., Stamenkovic, I., Rivera, M. N., Kadoch, C., Wucherpfennig, K. W., Rozenblatt-Rosen, O., Suvà, M. L., Riggi, N., Regev, A., “Opposing immune and genetic mechanisms shape oncogenic programs in synovial sarcoma.” In: *Nature medicine* 27 (2 Jan. 27, 2021). DOI: 10.1038/s41591-020-01212-6.

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Research: Preprint

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

Pipeline development: Workflow development language (WDL), Snake-make
HPC Job managers: Sun Grid Engine (SGE), Slurm, PBS-Torque
Cloud computing: Cromwell, 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