Michael Sellers Cuoco

Bioinformatics and Systems Biology PhD Student

Curriculum Vitae

June 2020

• University of California, San Diego

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Education

May 2016 - May 2018 Postbaccalaureate coursework

Harvard Extension School

Cambridge, Massachusetts

Sep 2012- May 2016 BS in Cellular and Molecular Biology

Trinity College

Hartford, Connecticut

➤ Minor: Models and Data

➤ Major GPA: 3.62

Honors and Awards

2017 Spot Award

Broad Institute

Cambridge, Massachusetts

Awarded to nominees to acknowledge and demonstrate appreciation and

recognition for their exceptional contributions.

2014 TriBeta National Biology Honors Society

Trinity College

Hartford, Connecticut

➤ Dedicated to improving the understanding and appreciation of biological study and extending boundaries of human knowledge through scientific research. Members must meet the national membership criteria.

2014 NESCAC Winter All-Academic Team

Trinity College

Hartford, Connecticut

Student-athletes must have reached sophomore academic standing, and be in good standing in their sport with a cumulative grade point average of at least 3 50

Experience

Jun 2011- Jul 2011

Aug 2016 - Aug 2020 Research Associate

Aviv Regev Lab, Broad Institute, MIT & Harvard

Cambridge, Massachuseets

➤ Collaborated with teams of physicians and scientists to characterize signatures of cancer drug resistance by single-cell and bulk transcriptome and chromatin profiling of patient, mouse, and cell line samples. Conducted functional validation primarily by in vitro by genetic perturbation experiments.

May 2015 - Aug 2015 Undergraduate Student

Matthew Meyerson Lab, Dana-Farber Cancer Institute, Harvard Medical School

Boston, Massachusetts

➤ Undergraduate thesis: Engineered an in vitro model of chromosome arm 8p loss by CRISPR/Cas9 editing and artificial telomere recombination to investigate the functional consequences of the common cancerous alteration.

May 2014 - Aug 2014 Undergraduate Student

Matthew Meyerson Lab, Dana-Farber Cancer Institute, Harvard Medical School

Boston, Massachusetts

Sep 2012- Dec 2013 Undergraduate Student

SEA-PHAGES, Genomics Research Program, Trinity College

Hartford, Connecticut

Highschool Student

Alan D'Andrea Lab, Dana-Farber Cancer Institute, Harvard Medical School

Boston, Massachusetts

Professional Training

2018 Harvard Biotech Incubator

Harvard Biotech Club

Boston, Massachusetts

➤ Worked directly with company founding members and key opinion leaders, performing due diligence and market research to identify clinical indications for therapeutic technology.

2018 Patent Law Short Course

Harvard Biotech Club

Boston, Massachusetts

Reviewed basic concepts of patent law through weekly case readings and workshops at a local firm.

2017 Heatlhcare Innovation & Commercialization Short Course

Harvard Biotech Club

Boston, Massachusetts

➤ Weekly modules addressed various aspects of the commercialization process of biomedical technology including intellectual property, market sizing, clinical paths, and FDA regulation.

Teaching

2015-2016 **Tutor**

Trinity College

Hartford, Connecticut

➤ Tutored struggling students one-on-one by request

2015-2016 Teaching Assistant

Trinity College

Hartford, Connecticut

➤ Hosted study sessions on topics in genetics

2018 CodeRATS

Broad Institute

Cambridge, Massachusetts

➤ Managed the leadership team for a series of institute-wide introduction to programming workshops.

Publications

Published

Dec 5, 2019 IL-33 Signaling Alters Regulatory T Cell Diversity in Support of Tumor Development.

A Li, RH Herbst, D Canner, JM Schenkel, OC Smith, JY Kim, M Hillman, A Bhutkar, MS Cuoco, CG Rappazzo, P Rogers, C Dang, L Jerby-Arnon, O Rozenblatt-Rosen, L Cong, M Birnbaum, A Regev, T Jacks

Cell reports doi.org/10.1016/j.celrep.2019.10.120

Dec 12, 2018 Acquired HER2 mutations in ER+ metastatic breast cancer confer resistance to estrogen receptordirected therapies.

U Nayar, O Cohen, C Kapstad, <u>MS Cuoco</u>, AG Waks, SA Wander, C Painter, S Freeman, NS Persky, L Marini, K Helvie, N Oliver, O Rozenblatt-Rosen, CX Ma, A Regev, EP Winer, NU Lin, N Wagle

Nature genetics doi.org/10.1038/s41588-018-0287-5

Nov 6, 2018 A Cancer Cell Program Promotes T Cell Exclusion and Resistance to Checkpoint Blockade.

L Jerby-Arnon, P Shah, MS Cuoco, C Rodman, MJ Su, JC Melms, R Leeson, A Kanodia, S Mei, JR Lin, S Wang, B Rabasha, D Liu, G Zhang, C Margolais, O Ashenberg, PA Ott, El Buchbinder, R Haq, FS Hodi, GM Boland, RJ Sullivan, DT Frederick, B Miao, T Moll, KT Flaherty, M Herlyn, RW Jenkins, R Thummalapalli, MS Kowalczyk, I Cañadas, B Schilling, ANR Cartwright, AM Luoma, S Malu, P Hwu, C Bernatchez, MA Forget, DA Barbie, AK Shalek, I Tirosh, PK Sorger, K Wucherpfennig, EM Van Allen, D Schadendorf, BE Johnson, A Rotem, O Rozenblatt-Rosen, LA Garraway, CH Yoon, B Izar, A Regev

Cell doi.org/10.1016/j.cell.2018.09.006

Sep 14, 2017 The neuropeptide NMU amplifies ILC2-driven allergic lung inflammation.

A Wallrapp, SJ Riesenfeld, PR Burkett, RE Abdulnour, J Nyman, D Dionne, M Hofree, MS Cuoco, C Rodman, D Farouq, BJ Haas, TL Tickle, JJ Trombetta, P Baral, CSN Klose, T Mahlakõiv, D Artis, O Rozenblatt-Rosen, IM Chiu, BD Levy, MS Kowalczyk, A Regev, VK Kuchroo

Nature doi.org/10.1038/nature24029

Apr 29, 2015 Whole genome comparison of a large collection of mycobacteriophages reveals a continuum of phage genetic diversity.

WH Pope, CA Bowman, DA Russell, D Jacobs-Sera, DJ Asai, SG Cresawn, WR Jacobs, RW Hendrix, JG Lawrence, GF Hatfull *eLife* doi.org/10.7554/eLife.06416

Preprints

Jun 5, 2020 Cycling cancer persister cells arise from lineages with distinct transcriptional and metabolic programs

Y Oren, M Tsabar, HF Cabanos, <u>MS Cuoco</u>, E Zaganjor, PI Thakore, M Tabaka, CP Fulco, SA Hurvitz, DJ Slamon, G Lahav, A Hata, JS Brugge, A Regev

BioRxiv doi.org/10.1101/2020.06.05.136358

Apr 20, 2020 Integrated analyses of single-cell atlases reveal age, gender, and smoking status associations with cell type-specific expression of mediators of SARS-CoV-2 viral entry and highlights inflammatory programs in putative target cells

C Muus, MD Luecken, G Eraslan, A Waghray, G Heimberg, L Sikkema, Y Kobayashi, ED Vaishnav, A Subramanian, C Smillie, K Jagadeesh, ET Duong, E Fiskin, E Torlai Triglia, C Becavin, M Ansari, P Cai, B Lin, J Buchanan, S Chen, J Shu, AL Haber, H Chung, DT Montoro, T Adams, H Aliee, SJ Allon, Z Andrusivova, I Angelidis, O Ashenberg, K Bassler, C Becavin, I Benhar, J Bergenstrahle, L Bergenstrahle, L Bolt, E Braun, LT Bui, M Chaffin, E Chichelnitskiy, J Chiou, TM Conlon, MS Cuoco, M Deprez, NA Fisc BioRxiv doi.org/10.1101/2020.04.19.049254

Mar 20, 2020 Transcriptional mediators of treatment resistance in lethal prostate cancer

MX He, MS Cuoco, J Crowdis, A Bosma-Moody, Z Zhang, K Bi, A Kanodia, M-J Su, C Rodman, L DelloStritto, P Shah, KP Burke, B Izar, Z Bakouny, AK Tewari, D Liu, SY Camp, NI Vokes, J Park, S Vigneau, L Fong, O Rozenblatt-Rosen, A Regev, A Rotem, M-E Taplin, EM Van Allen

BioRxiv doi.org/10.1101/2020.03.19.998450

Oct 21, 2019 Pan-cancer single cell RNA-seq uncovers recurring programs of cellular heterogeneity

GS Kinker, AC Greenwald, R Tal, Z Orlova, MS Cuoco, JM McFarland, A Warren, C Rodman, JA Roth, SA Bender, B Kumar, JW Rocco, PA Fernandes, CC Mader, H Keren-Shaul, A Plotnikov, H Barr, A Tsherniak, O Rozenblatt-Rosen, V Krizhanovsky, SV Puram, A Regev, I Tirosh

BioRxiv doi.org/10.1101/807552

Aug 28, 2019 The enteric nervous system of the human and mouse colon at a single-cell resolution

E Drokhlyansky, CS Smillie, N Van Wittenberghe, M Ericsson, GK Griffin, D Dionne, MS Cuoco, MN Goder-Reiser, T Sharova, AJ Aguirre, GM Boland, D Graham, O Rozenblatt-Rosen, RJ Xavier, A Regev BioRxiv doi.org/10.1101/746743

Aug 4, 2019 Opposing immune and genetic forces shape oncogenic programs in synovial sarcoma

L Jerby, C Neftel, ME Shore, MJ McBride, B Haas, B Izar, HR Weissman, A Volorio, G Boulay, L Cironi, AR Richman, LC Broye, JM Gurski, CC Luo, R Mylvaganam, L Nguyen, S Mei, Jc Melms, C Georgescu, O Cohen, JE Buendia-Buendia, MS Cuoco, D Labes, DR Zollinger, JM Beechem, P Nielsen, I Chebib, G Cote, E Choy, I Letovanec, S Cherix, N Wagle, PK Sorger, AB Haynes, JT Mullen, I Stamenkovic, MN Rivera, C Kadoch, O Rozenblatt-Rosen, ML Suva, N Riggi, A Regev BioRxiv doi.org/10.1101/724302

Apr 12, 2019 Acquired FGFR and FGF alterations confer resistance to estrogen receptor (ER) targeted therapy in ER+ metastatic breast cancer

P Mao, O Cohen, KJ Kowalski, JG Kusiel, JE Buendia-Buendia, <u>MS Cuoco</u>, P Exman, SA Wander, AG Waks, U Nayar, J Chung, S Freeman, O Rozenblatt-Rosen, VA Miller, F Federica Piccioni, DE Root, A Regev, EP Winer, NU Lin, N Wagle *BioRxiv* doi.org/10.1101/605436

Publications are updated programmatically each week.

Presentations

Talks

Apr 2020	The cellular origins of drug resistance in cancer Regev Lab Staff Meeting; Cambridge, Massachusetts
Oct 2018	CRISPR screening for regulators of cancer immune checkpoint inhibitor resistance Regev Lab Science Days Retreat; Cambridge, Massachusetts
May 2017	Understanding the mechanisms of drug resistance in melanoma Regev Lab Staff Meeting; Cambridge, Massachusetts

May 2016	In vitro modeling and analysis of chromosome 8p arm-level deletion using CRISPR-Cas9. Trinity College Biology Department; Hartford, Connecticut
Aug 2015	In vitro modeling and analysis of chromosome 8p arm-level deletion using CRISPR-Cas9. Meyerson Lab Group Meeting; Boston, Massachusetts
Aug 2014	Genome engineering to generate models of chromosome arm-level aneuploidies. Meyerson Lab Group Meeting; Boston, Massachusetts
Nov 2012	The role of the FANCD2 gene in Fanconi Anemia and DNA repair. Concord-Carlisle High School STEM series; Concord, Massachusetts
Posters	
Dec 2019	Metabolic switching underlies the ability of cancer persister cells to cycle under drug treatment. Annual Broad Institute Retreat; Boston, Massachusetts
Feb 2019	Targeting the root of non-genetic cancer relapse using an expressed barcode library. Annual Klarman Cell Observatory Retreat; Cambridge, Massachusetts
Dec 2018	Discovering the master regulators of immune checkpoint inhibitor resistance in melanoma with Perturb-Seq. Annual Broad Institute Retreat; Boston, Massachusetts
Jul 2018	Single-cell RNA-Seq of melanoma ecosystems reveals sources of T cell exclusion linked to immunotherapy clinical outcomes. Annual Broad Institute-Israel Science Foundation Symposium; Cambridge, Massachusetts
Jun 2018	The Center for Cancer Precision Medicine enables exploration of immunotherapy resistance in melanoma at the single-cell level. Annual Dana-Farber / Harvard Cancer Center Genetics Retreat; Boston, Massachusetts
May 2016	In vitro modeling and analysis of chromosome 8p arm-level deletion using CRISPR-Cas9. Trinity College Annual Spring Research Symposium; Hartford, Connecticut
May 2013	Review of integrase-mediated site-specific recombination in mycobacteriophage species. Trinity College Annual Spring Research Symposium; Hartford, Connecticut
Service	

2018 **Patient Ambassador** Dana-Farber Cancer Institute

Boston, Massachusetts

➤ Escorted patients to appointments across the Longwood Medical Area

Skills

statistical modelling, data science, reproducible research Analytical R (advanced), Bash, Matlab, Python Programming tidyverse, Rmarkdown, blogdown Packages Git, Docker, Travis Tools