# **Michael Sellers Cuoco**

PhD Student, Bioinformatics and Systems Biology

**Curriculum Vitae** 

July 2020

University of California, San Diego

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### **Education**

May 2016 - May 2018 Postbaccalaureate coursework

**Harvard Extension School** 

Sep 2012- May 2016 BS in Cellular and Molecular Biology

**Trinity College** 

➤ Minor: Models and Data

➤ Major GPA: 3.62

Cambridge, Massachusetts

Hartford, Connecticut

### **Honors and Awards**

2017 Spot Award Cambridge, Massachusetts

**Broad Institute** 

➤ Awarded to nominees to acknowledge and demonstrate appreciation and recognition for their exceptional contributions.

2014 TriBeta National Biology Honors Society

Hartford, Connecticut

**Trinity College** 

➤ 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

Hartford, Connecticut

**Trinity College** 

➤ 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**

Aug 2016 – Aug 2020 Research Associate Cambridge, Massachusetts

Aviv Regev Lab, Broad Institute, MIT & Harvard University

➤ 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 investigation by pooled genetic perturbation experiments.

May 2014- Aug 2015 Undergraduate Student

Boston, Massachusetts

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

• 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.

Sep 2012- Dec 2013 Undergraduate Student

Hartford, Connecticut

SEA-PHAGES, Genomics Research Program, Trinity College

Jun 2011 – Jul 2011 High school Student

Boston, Massachusetts

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

### **Professional Training**

2018 Harvard Biotech Incubator

Boston, Massachusetts

Harvard Biotech Club

➤ 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

Boston, Massachusetts

Harvard Biotech Club

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

Heatlhcare Innovation & Commercialization Short Course

Boston, Massachusetts

Hartford, Connecticut

Harvard Biotech Club

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

### **Teaching**

2017

2018 CodeRATS Cambridge, Massachusetts

**Broad Institute** 

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

2015-2016 Teaching Assistant Hartford, Connecticut

**Trinity College** 

➤ Hosted study sessions on topics in genetics

2015-2016 **Tutor** 

**Trinity College** 

➤ Tutored biology and genetics students one-on-one by request

## **Publications**

#### **Published**

Jun 24, 2020 A single-cell landscape of high-grade serous ovarian cancer.

B Izar, I Tirosh, EH Stover, I Wakiro, <u>MS Cuoco</u>, I Alter, C Rodman, R Leeson, MJ Su, P Shah, M Iwanicki, SR Walker, A Kanodia, JC Melms, S Mei, JR Lin, CBM Porter, M Slyper, J Waldman, L Jerby-Arnon, O Ashenberg, TJ Brinker, C Mills, M Rogava, S Vigneau, PK Sorger, LA Garraway, PA Konstantinopoulos, JF Liu, U Matulonis, BE Johnson, O Rozenblatt-Rosen, A Rotem, A Regev *Nature medicine* doi.org/10.1038/s41591-020-0926-0

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, MS Cuoco, 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, <u>MS Cuoco</u>, 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 (

MS Cuoco listed at co-investigator)

eLife doi.org/10.7554/eLife.06416

#### **Preprints**

# Jun 23, 2020 RAAS blockade, kidney disease, and expression of ACE2, the entry receptor for SARS-CoV-2, in kidney epithelial and endothelial cells

A Subramanian, K Vernon, M Slyper, J Waldman, MD Luecken, K Gosik, D Dubinsky, M Cuoco, K Keller, J Purnell, L Nguyen, D Dionne, O Rozenblatt-Rosen, A Weins, Human Cell Atlas Lung Biological Network, A Regev, A Greka *BioRxiv* doi.org/10.1101/2020.06.23.167098

### 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, Pl 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, <u>MS Cuoco</u>, 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**

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.
Jun 2018	Annual Broad Institute-Israel Science Foundation Symposium; Cambridge, Massachusetts  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 Boston, Massachusetts

Dana-Farber Cancer Institute

➤ Escorted patients to appointments across the Longwood Medical Area

# **Skills**

statistical modelling, data science, reproducible researchAnalyticalR (advanced), Bash, Matlab, PythonProgrammingtidyverse, Rmarkdown, blogdownPackagesGit, Docker, TravisTools