

Aaron S. Meyer

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EDUCATION

Massachusetts Institute of Technology, Cambridge, MA 2009–2014

Ph.D., Biological Engineering

- Thesis: Quantitative approaches to understanding signaling regulation of 3D cell migration

University of California, Los Angeles, CA 2005–2009

B.S., Bioengineering, magna cum laude

PROFESSIONAL EXPERIENCE

Associate Professor *Los Angeles, CA*

University of California, Los Angeles

2023–Present

Bioengineering Department

Bioinformatics Interdepartmental Graduate Program

Computational & Systems Biology Interdepartmental Program

Vice Chair of Graduate Studies *Los Angeles, CA*

University of California, Los Angeles

2024–Present

Bioengineering Department

Co-Director, Amgen Scholars Program *Los Angeles, CA*

University of California, Los Angeles

2024–Present

Assistant Professor *Los Angeles, CA*

University of California, Los Angeles

2017–2023

Principal Investigator & Research Fellow *Cambridge, MA*

Massachusetts Institute of Technology

2014–2017

Koch Institute for Integrative Cancer Research

Graduate Student Researcher *Cambridge, MA*

Massachusetts Institute of Technology

2009–2014

Department of Biological Engineering

Koch Institute for Integrative Cancer Research

Undergraduate Researcher *Los Angeles, CA*

University of California, Los Angeles

2006–2009

In the lab of Daniel Kamei

AWARDS & RECOGNITIONS

Diversity, Equity and Inclusion Award, Society For Biomaterials 2025

Outstanding Mentor Award, Bruins-In-Genomics 2022, 2024

Emerging Leader Award, Mark Foundation for Cancer Research 2023

Northrop Grumman Excellence in Teaching Award 2021

Milstein Abstract Award, Cytokine Society 2021

UCLA Hellman Fellow 2019

Career Awards at the Scientific Interface Finalist, Burroughs Wellcome Fund 2017

Fellowship, Terri Brodeur Breast Cancer Foundation 2017

Ten to Watch, Amgen Scholars Foundation 2016

Director's Early Independence Award, National Institutes of Health 2014

Highlighted by the NIH director's office ([link](#))

PUBLICATIONS

- [1] H. Kojima, T. A. Morinelli, Y. Wang, **J. L. Chin, A. S. Meyer**, Y.-C. Kao, K. Kadono, S. Yao, T. Torgerson, K. J. Dery, A. Bhat, E. F. Reed, F. M. Kaldas, D. J. van der Windt, D. G. Farmer, and J. W. Kupiec, "Group 1 innate lymphoid cells protect liver transplants from ischemia-reperfusion injury via an interferon- γ -mediated pathway," *American Journal of Transplantation*, Dec. 2024, doi: 10.1016/j.ajt.2024.11.035.
- [2] **A. A. Abraham, Z. C. Tan, P. Shrestha, E. R. Bozich, and A. S. Meyer**, "A multivalent binding model infers antibody Fc species from systems serology," *PLoS Computational Biology*, vol. 20, no. 12, p. e1012663, Dec. 2024, doi: 10.1371/journal.pcbi.1012663.
- [3] C. S. Movassaghi, K. A. Perrotta, M. E. Curry, A. N. Nashner, K. K. Nguyen, M. E. Wesely, M. A. Fillol, C. Lui, **A. S. Meyer**, and A. M. Andrews, "Machine-learning-guided design of voltammetry waveforms," *chemRxiv [Preprint]*, Dec. 2024.
- [4] **B. Orcutt-Jahns, J. R. L. Junior, E. Lin, R. C. Rockne, A. Matache, S. Branciamore, E. Hung, A. S. Rodin, P. P. Lee, and A. S. Meyer**, "Systems profiling reveals recurrently dysregulated cytokine signaling responses in ER+ breast cancer patients' blood," *npj Systems Biology and Applications*, no. 118, Oct. 2024, doi: 10.1038/s41540-024-00447-0.
- [5] **Z. C. Tan and A. S. Meyer**, "The structure is the message: preserving experimental context through tensor decomposition," *Cell Systems*, vol. 15, no. 8, pp. 679–693, Aug. 2024, doi: 10.1016/j.cels.2024.07.004.
- [6] **A. Ramirez, B. T. Orcutt-Jahns, S. Pascoe, A. Abraham, B. Remigio, N. Thomas, and A. S. Meyer**, "Integrative, high-resolution analysis of single cells across experimental conditions with PARAFAC2," *bioRxiv [Preprint]*, Jul. 2024, doi: 10.1101/2024.07.29.605698v1.
- [7] **E. C. Hung, E. Hodzic, Z. C. Tan, and A. S. Meyer**, "Censored Least Squares for Imputing Missing Values in PARAFAC Tensor Factorization," *bioRxiv [Preprint]*, Jul. 2024.
- [8] **J. L. Chin, Z. C. Tan, L. C. Chan, F. Ruffin, R. Parmar, R. Ahn, S. Taylor, A. S. Bayer, A. Hoffmann, J. Vance G Fowler, E. F. Reed, M. R. Yeaman, A. S. Meyer, and MRSA Systems Immunobiology Group**, "Tensor modeling of MRSA bacteremia cytokine and transcriptional patterns reveals coordinated, outcome-associated immunological programs," *PNAS Nexus*, p. pgae185, May 2024, doi: 10.1093/pnasnexus/pgae185.
- [9] **M. Creixell, S. D. Taylor, J. Gerritsen, S. Y. Bae, M. Jiang, T. Augustin, M. Loui, C. Boixo, P. Creixell, F. M. White, and A. S. Meyer**, "Dissecting signaling regulators driving AXL-mediated bypass resistance and associated phenotypes by phosphosite perturbations," *bioRxiv [Preprint]*, Oct. 2023.
- [10] **B. Orcutt-Jahns, P. C. Emmel, E. M. Snyder, S. D. Taylor, and A. S. Meyer**, "Multivalent, asymmetric IL-2–Fc fusions show enhanced selectivity for regulatory T cells," *Science Signaling*, Oct. 2023, doi: 10.1126/scisignal.adg0699.
- [11] S. R. Peyton, L. W. Chow, S. D. Finley, A. N. F. Versypt, R. Hill, M. L. Kemp, E. M. Langer, A. P. McGuigan, **A. S. Meyer**, S. K. Seidlits, K. Roy, and S. M. Mumenthaler, "Synthetic living materials in cancer biology," *Nature Reviews Bioengineering*, Oct. 2023, doi: 10.1038/s44222-023-00105-w.
- [12] A. Q. Terry, H. Kojima, R. A. Sosa, F. M. Kaldas, **J. L. Chin**, Y. Zhenga, B. V. Naini, D. Noguchi, J. Nevarez-Mejia, Y.-P. Jin, R. W. Busuttil, **A. S. Meyer**, D. W. Gjertson, J. W. Kupiec-Weglinski, and E. F. Reed, "Disulfide-HMGB1 signals through TLR4 and TLR9 to induce inflammatory macrophages capable of innate-adaptive crosstalk in human liver transplantation," *American Journal of Transplantation*, Aug. 2023, doi: 10.1016/j.ajt.2023.08.002.
- [13] **Z. C. Tan, A. Lux, M. Biburger, P. Varghese, S. Lees, F. Nimmerjahn, and A. S. Meyer**, "Mixed IgG Fc immune complexes exhibit blended binding profiles and refine FcR affinity estimates," *Cell Reports*, Jul. 2023, doi: 10.1016/j.celrep.2023.112734.

- [14] S. M. Gross, **F. Mohammadi**, C. Sanchez-Aguila, P. J. Zhan, **A. S. Meyer**, and L. M. Heiser, "Analysis and modeling of cancer drug responses using cell cycle phase-specific rate effects," *Nature Communications*, Jun. 2023, doi: 10.1101/2020.07.24.219907.
- [15] **J. L. Chin**, L. C. Chan, M. R. Yeaman, and **A. S. Meyer**, "Tensor-based insights into systems immunity and infectious disease," *Trends in Immunology*, vol. 44, no. 5, pp. 329–332, May 2023, doi: 10.1016/j.it.2023.03.003.
- [16] H. Yang, U. Y. Ulge, A. Quijano-Rubio, Z. J. Bernstein, J. David R. Maestas, J.-H. Chun, W. Wang, J.-X. Lin, K. M. Jude, S. Singh, **B. T. Orcutt-Jahns**, P. Li, J. Mou, L. Chung, Y.-H. Kuo, Y. H. Ali, **A. S. Meyer**, W. L. Grayson, N. M. Heller, K. C. Garcia, W. J. Leonard, D.-A. Silva, J. H. Elisseeff, D. Baker, and J. B. Spangler, "Design of cell-type-specific hyperstable IL-4 mimetics via modular de novo scaffolds," *Nature Chemical Biology*, pp. 1552–4469, Apr. 2023, doi: 10.1038/s41589-023-01313-6.
- [17] **C. Wilder**, D. Lefaudoux, R. Mathenge, K. Kishimoto, A. Z. Munoz, M. A. Nguyen, **A. S. Meyer**, Q. J. Cheng, and A. Hoffmann, "A stimulus-contingent positive feedback loop enables IFN- β dose-dependent activation of pro-inflammatory genes," *Molecular Systems Biology*, p. e11294, Mar. 2023, doi: 10.15252/msb.202211294.
- [18] P. Kulkarni, H. S. Wiley, H. Levine, H. Sauro, A. Anderson, S. T. C. Wong, **A. S. Meyer**, P. Iyengar, K. Corlette, K. Swanson, A. Mohanty, S. Bhattacharya, A. Patel, V. Jain, and R. Salgia, "Addressing the genetic/nongenetic duality in cancer with systems biology," *Trands in Cancer*, vol. 9, no. 3, pp. 185–187, Mar. 2023, doi: 10.1016/j.trecan.2022.12.004.
- [19] H. Kim, A. Wirasaputra, **F. Mohammadi**, A. N. Kundu, J. A. E. Esteves, L. M. Heiser, **A. S. Meyer**, and S. R. Peyton, "Live Cell Lineage Tracing of Dormant Cancer Cells," *Advanced Healthcare Materials*, vol. 12, no. 14, p. 2202275, Jan. 2023, doi: 10.1002/adhm.202202275.
- [20] **F. Mohammadi**, **S. Visagan**, S. M. Gross, **L. Karginov**, **J. Lagarde**, L. M. Heiser, and **A. S. Meyer**, "A lineage tree-based hidden Markov model to quantify cellular heterogeneity and plasticity," *Communications Biology*, Nov. 2022, doi: 10.1038/s42003-022-04208-9.
- [21] D. VanDyke, M. Iglesias, J. Tomala, A. Young, J. Smith, J. A. Perry, E. Gebara, A. R. Cross, L. S. Cheung, A. G. Dykema, **B. T. Orcutt-Jahns**, T. Henclová, J. Golias, J. Balolong, L. M. Tomasovic, D. Funda, **A. S. Meyer**, D. M. Pardoll, J. Hester, F. Issa, C. A. Hunter, M. S. Anderson, J. A. Bluestone, G. Raimondi, and J. B. Spangler, "Engineered human cytokine/antibody fusion proteins expand regulatory T cells and confer autoimmune disease protection," *Cell Reports*, Oct. 2022, doi: 10.1016/j.celrep.2022.111478.
- [22] **M. Creixell**, H. Kim, **F. Mohammadi**, S. R. Peyton, and **A. S. Meyer**, "Systems approaches to uncovering the contribution of environment-mediated drug resistance," *Current Opinion in Solid State and Materials Science*, Oct. 2022, doi: 10.1016/j.cossms.2022.101005.
- [23] **M. Creixell** and **A. S. Meyer**, "Dual data and motif clustering improves the modeling and interpretation of phosphoproteomic data," *Cell Reports Methods*, Feb. 2022, doi: 10.1016/j.crmeth.2022.100167.
- [24] A. Majumder, S. Hosseiniyan, M. J. Stroud, E. Adhikari, J. J. Saller, D. M. A. Smith, D. G. Zhang, S. Agarwal, **M. Creixell**, B. S. Meyer, M. F. Kinose, K. S. Bowers, B. Fang, P. A. Stewart, E. A. Welsh, T. A. Boyle, **A. S. Meyer**, J. M. Koomen, and E. B. Haura, " Integrated proteomics-based physical and functional mapping of AXL kinase signaling pathways and inhibitors define its role in cell migration," *Molecular Cancer Research*, Jan. 2022, [Online]. Available: <https://mcr.aacrjournals.org/content/early/2022/01/12/1541-7786.MCR-21-0275.abstract>
- [25] **Z. C. Tan**, **B. T. Orcutt-Jahns**, and **A. S. Meyer**, "A quantitative view of strategies to engineer cell-selective ligand binding," *Integrative Biology*, Dec. 2021, [Online]. Available: <https://academic.oup.com/ib/advance-article/doi/10.1093/intbio/zyab019/6470588?guestAccessKey=d1740482-b50d-426f-b177-621426c0a5bc>

- [26] **Z. C. Tan, M. Murphy, H. S. Alpay, S. D. Taylor, and A. S. Meyer**, “Tensor-structured decomposition improves systems serology analysis,” *Molecular Systems Biology*, vol. 17, no. 9, p. e10243, Sep. 2021, doi: 10.15252/msb.202110243.
- [27] **A. M. Farhat, A. C. Weiner, C. Posner, Z. S. Kim, S. M. Carlson, and A. S. Meyer**, “Modeling Cell-Specific Dynamics and Regulation of the Common Gamma Chain Cytokines,” *Cell Reports*, Apr. 2021, doi: 10.1016/j.celrep.2021.109044.
- [28] **Z. C. Tan and A. S. Meyer**, “A general model of multivalent binding with ligands of heterotypic subunits and multiple surface receptors ,” *Mathematical Biosciences*, p. 108714, 2021, doi: 10.1016/j.mbs.2021.108714.
- [29] **S. Y. Bae, N. Guan, R. Yan, K. Warner, S. D. Taylor, and A. S. Meyer**, “Measurement and models accounting for cell death capture hidden variation in compound response,” *Cell Death & Disease*, vol. 11, no. 4, p. 255, 2020, doi: 10.1038/s41419-020-2462-8.
- [30] C.-H. Lee, T. H. Kang, O. Godon, M. Watanabe, G. Delidakis, C. M. Gillis, D. Sterlin, D. Hardy, M. Cogné, L. E. Macdonald, A. J. Murphy, N. Tu, J. Lee, J. R. McDaniel, E. Makowski, P. M. Tessier, **A. S. Meyer**, P. Bruhns, and G. Georgiou, “An engineered human Fc domain that behaves like a pH-toggle switch for ultra-long circulation persistence ,” *Nature Communications*, vol. 10, no. 1, p. 5031, Nov. 2019, doi: 10.1038/s41467-019-13108-2.
- [31] **A. S. Meyer** and L. M. Heiser, “Systems biology approaches to measure and model phenotypic heterogeneity in cancer,” *Current Opinion in Systems Biology*, vol. 17, pp. 35–40, 2019, doi: 10.1016/j.coisb.2019.09.002.
- [32] K. Situ, B. A. Chua, **S. Y. Bae, A. S. Meyer**, and K. Morizono, “Versatile targeting system for lentiviral vectors involving biotinylated targeting molecules,” *Virology*, vol. 525, pp. 170–181, Dec. 2018, doi: 10.1016/j.virol.2018.09.017.
- [33] A. M. Claas, L. Atta, S. Gordonov, **A. S. Meyer**, and D. A. Lauffenburger, “Systems Modeling Identifies Divergent Receptor Tyrosine Kinase Reprogramming to MAPK Pathway Inhibition ,” *Cellular and Molecular Bioengineering*, Jul. 2018, doi: 10.1007/s12195-018-0542-y.
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- [35] **R. A. Robinett, N. Guan**, A. Lux, M. Biburger, F. Nimmerjahn, and **A. S. Meyer**, “Dissecting FcγR Regulation Through a Multivalent Binding Model,” *Cell Systems*, Jun. 2018, doi: 10.1016/j.cels.2018.05.018.
- [36] **A. Zweemer**, C. B. French, J. Mesfin, S. Gordonov, **A. S. Meyer**, and D. A. Lauffenburger, “Apoptotic Bodies Elicit Gas6-mediated Migration of AXL-expressing Tumor Cells,” *Molecular Cancer Research*, 2017, doi: 10.1158/1541-7786.mcr-17-0012.
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- [39] **S. Manole, E. J. Richards**, and **A. S. Meyer**, “JNK Pathway Activation Modulates Acquired Resistance to EGFR/HER2-Targeted Therapies,” *Cancer Research*, vol. 76, no. 18, pp. 5219–5228, Sep. 2016, doi: 10.1158/0008-5472.can-16-0123.
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Inhibitor Resistance ,” *Cancer Discovery*, vol. 6, no. 4, pp. 331–333, Apr. 2016, doi: 10.1158/2159-8290.cd-15-0933.

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- [42] M. A. Miller, M. L. Moss, G. Powell, R. Petrovich, L. Edwards, **A. S. Meyer**, L. G. Griffith, and D. A. Lauffenburger, “ Targeting autocrine HB-EGF signaling with specific ADAM12 inhibition using recombinant ADAM12 prodomain ,” *Scientific Reports*, vol. 5, p. 15150, Oct. 2015, [Online]. Available: <https://www.nature.com/articles/srep15150>
- [43] D. N. Riquelme, **A. S. Meyer**, M. Barzik, A. Keating, and F. Gertler, “Selectivity in Subunit Composition of Ena/VASP Tetramers,” *Bioscience Reports*, vol. 35, no. 5, Oct. 2015, doi: 10.1042/bsr20150149.
- [44] **A. S. Meyer**, **A. J. M. Zweemer**, and D. A. Lauffenburger, “The AXL Receptor Is a Sensor of Ligand Spatial Heterogeneity,” *Cell Systems*, vol. 1, no. 1, pp. 25–36, Jun. 2015, doi: 10.1016/j.cels.2015.06.002.
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- [50] Y.-S. Tsao, A. Merchant, **A. Meyer**, Z. Liu, M. Smith, D. Levitan, and E. Gustafson, “ Integrated Pathway Analysis of Genome-Wide Expression Changes Associated with Serum-Free Suspension Adaptation of an Antibody-Producing Chinese Hamster Ovary (CHO) Cell Line ,” *Animal Cell Technology: Basic & Applied Aspects*, vol. 16. in *Animal Cell Technology: Basic & Applied Aspects*, vol. 16. pp. 27–32, Mar. 2010. doi: 10.1007/978-90-481-3892-0_5.
- [51] F. Mashayekhi, **A. S. Meyer**, S. A. Shiigi, V. Nguyen, and D. T. Kamei, “Concentration of mammalian genomic DNA using two-phase aqueous micellar systems,” *Biotechnology and Bioengineering*, vol. 102, no. 6, pp. 1613–1623, Nov. 2009, doi: 10.1002/bit.22188.
- [52] S. A. Shiigi, **A. S. Meyer**, and D. T. Kamei, “Enhancing the Detection of Urinary Tract Infections Using Two-Phase Aqueous Micellar Systems,” *The UCLA USJ*, pp. 47–56, 2009.

RESEARCH SUPPORT

Contact PI on all grants unless indicated otherwise.

NIH NIAID U01-AI179524 (Co-I)

2024–2029

“Systems Analyses of Induction and Maintenance of Immunity to SARS-CoV-2 Vaccination in Kidney Transplant Recipients Receiving Mycophenolate Mofetil Immunotherapy”

| | |
|---|-----------|
| <i>NSF 2404470 (Co-I)</i> | 2024–2027 |
| “Multi-analyte Detection Enabled by Machine Learning-Guided Voltammetry” | |
| Division: CHE Research and Development | |
| <i>NIH NEI R01-EY011996 (Co-I)</i> | 2023–2027 |
| “Retinal Disease: Molecular Basis and Pathophysiology” | |
| <i>NIH NIAID U19-AI172713 (Co-PI)</i> | 2023–2028 |
| Systems Biology for Infectious Diseases Consortium | |
| “Systems Epigenomics of Persistent Bloodstream Infection” | |
| <i>Emerging Leader Award, Mark Foundation for Cancer Research</i> | 2023–2025 |
| “Tracking and Reactivating Humoral Immunity through Systems Serology” | |
| <i>SEEDS Grant</i> | 2023–2024 |
| Merck & Co., Inc. | |
| “Systematic and Receptor-Specific Dissection of Fc Receptor Functions” | |
| <i>NIH NIAID P01-AI120944 (Co-I)</i> | 2022–2027 |
| Transplant Immunology Program Project Grant | |
| “Innate-Adaptive Immunoregulation in Liver Transplant Ischemia/Reperfusion Injury” | |
| <i>COVID Relief Funds, Vice Chancellor for Research Office</i> | 2022 |
| <i>Administrative Supplement to U01-CA215709</i> | 2020–2021 |
| “Mechanistic Autoencoders for Patient-Specific Phosphoproteomic Models” | |
| <i>Grant</i> | 2020–2022 |
| Jayne Koskinas Ted Giovanis Foundation | |
| “Cell cycle-specific drug responses in breast cancer” | |
| <i>American Cancer Society, Research Scholar Grant (co-I)</i> | 2020–2023 |
| “Tissue-engineered models of glioblastoma for evaluating treatment responses” | |
| <i>NIH NIAID U01-AI148119</i> | 2019–2024 |
| Fc-Dependent Mechanisms of Antibody-Mediated Killing Consortium | |
| “Mapping the effector response space of antibody combinations” | |
| <i>UCLA Faculty Career Development Award</i> | 2019–2020 |
| <i>UCLA Hellman Fellow</i> | 2019–2020 |
| “Engineering anti-tumor antibody combinations for more effective and less toxic therapies” | |
| <i>Visterra, Inc. Research Agreement</i> | 2019–2021 |
| “IL-2 Receptor Binding Engineering” | |
| <i>Administrative Supplement to U01-CA215709</i> | 2018–2019 |
| “Cell lineage analysis to quantify heterogeneous cell cycle responses of cancer cells” | |
| <i>NIH NCI U01-CA215709</i> | 2017–2022 |
| Cancer Systems Biology Consortium | |
| “Precision Lung Cancer Therapy Design through Multiplexed Adapter Measurement” | |
| <i>Fellowship Grant</i> | 2017–2019 |
| Terri Brodeur Breast Cancer Foundation | |
| “Decoding the Role of TAM Receptors <i>In Vivo</i> Using More Specific and Potent Inhibitors” | |
| <i>AMIGOS Program Award</i> | 2016–2020 |
| Jayne Koskinas Ted Giovanis Foundation and Breast Cancer Research Foundation | |
| “Understanding the Role of Cell Plasticity in Mediating Drug Resistance” | |
| <i>GPU Grant</i> | 2016 |
| NVIDIA Corporation | |
| “Parameterizing Stochastic Cell Signaling Pathways Through Variability Fitting” | |

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| <i>Frontier Research Program Initiator Award</i> | 2015 |
| Koch Institute for Integrative Cancer Research | |
| “Multiplexed Tools for Probing Chemokine Receptor Activation State in Breast Cancer” | |
| <i>NIH Director’s Early Independence Award, DP5-OD019815</i> | 2014–2019 |
| “Adapter-Layer RTK Signaling: Basic Understanding & Targeted Drug Resistance” | |
| <i>Whitaker Fellowship</i> | 2013 |
| Massachusetts Institute of Technology | |
| <i>Repligen Fellowship in Cancer Research</i> | 2012 |
| Koch Institute for Integrative Cancer Research | |
| <i>Frontier Research Program Initiator Award</i> | 2011 |
| Koch Institute for Integrative Cancer Research | |
| “Global Growth Factor Reprogramming and Invasion By AXL Expression And Shedding In Breast Carcinoma” | |
| <i>Breast Cancer Research Predoctoral Fellowship</i> | 2010–2014 |
| Department of Defense, W81XWH-11-1-0088 | |
| “Molecular Regulatory Network Dysregulation in Breast Cancer Cell Migration & Invasion” | |
| <i>Graduate Research Fellowship</i> | 2009–2014 |
| National Science Foundation | |
| <i>Momenta Presidential Fellowship</i> | 2009 |
| Massachusetts Institute of Technology | |

TEACHING EXPERIENCE

| | |
|---|---|
| Instructor | <i>UCLA, Department of Bioengineering</i> |
| <i>Machine Learning & Data-Driven Modeling in Bioengineering</i> | 2018–Present |
| • Designed and lead project-based course tailored to the background of students in the program | |
| Instructor | <i>UCLA, Department of Bioengineering</i> |
| <i>Bioengineering Laboratory</i> | 2018–2024 |
| • Lab-based introduction to basics of experimental design and laboratory work in bioengineering | |
| Guest Speaker | <i>UCLA, Bioinformatics Interdepartmental Program</i> |
| <i>Bioinformatics 202</i> | 2023 |
| • Discussed our lab’s research and related topics in bioinformatics. | |
| Guest Lecturer | <i>UCLA, Molecular, Cell, and Developmental Biology</i> |
| <i>Fundamentals of Digital Imaging and Image Processing</i> | 2021, 2022 |
| • Led discussion of a paper from the lab used as a project within the class | |
| Discussion Leader | <i>UCLA, Microbiology, Immunology, & Molecular Genetics</i> |
| <i>Ethics and Accountability in Biomedical Research</i> | 2021, 2024 |
| • Led discussion of various ethics case studies | |
| Advisor & Instructor | <i>UCLA, Undergraduate Research Center</i> |
| <i>Integrated and Interdisciplinary Undergraduate Research Program</i> | 2019–2023 |
| • Advised program participants on developing research, presentation, and professional skills | |
| Team Mentor | <i>UCLA, Department of Bioengineering</i> |
| <i>Capstone Design</i> | 2017, 2018, 2019, 2020 |
| • Poster competition winning team: 2018, 2019 | |
| Guest Speaker | <i>UCLA, Department of Bioengineering</i> |
| <i>Introduction to Bioengineering</i> | 2017, 2019, 2020, 2023, 2024 |
| • Guest speaker to discuss research program and opportunities in bioengineering | |

Faculty of the Citizen Science Program*Citizen Science Program*

- Led a short course introducing students to the natural sciences and scientific method

Teaching Assistant*Thermodynamics of Biomolecular Systems**Bard College, Annandale-on-Hudson, NY*

2015–2016

MIT, Department of Biological Engineering

2010

CONFERENCE & INVITED PRESENTATIONS (LAST FIVE YEARS)

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| Maintaining Immunity After Immunization Kickoff Meeting, Invited Oral Presentation “Systems analyses of induction and maintenance of immunity to SARS-CoV-2 vaccination in KTx receiving MMF.” | Nov 2024 |
| UCLA, Bioengineering, Invited Dept. Seminar “Building an integrative view of immunity with tensors.” | Oct 2024 |
| Massachusetts Institute of Technology, Biological Engineering, Invited Oral Presentation “Experiment structure is the message: building an integrative view of immunity with tensors.” | Sep 2024 |
| UCLA Bruins-In-Genomics Summer Program, Invited Oral Presentation “Experiment structure is the message: building an integrative view of immunity with tensors.” | Jul 2024 |
| Fc Mechanisms of Cell Killing Workshop, Invited Oral Presentation “Mapping the effector response space of antibody combinations” | Jun 2024 |
| NIAID Systems Biology Consortium Webinar, Invited Oral Presentation “Tensor Modeling of Clinical Outcomes in <i>S. aureus</i> Bacteremia.” | May 2024 |
| Cytokine Based Drug Development Summit, Invited Oral Presentation “New Cytokine Targeting Strategies Enabled by Multivalent Cis-Targeted Complexes.” | May 2024 |
| Tracer Precision Health Workshop, Invited Oral Presentation “Mechanistic, integrative, and high-resolution dissection of single-cell studies with PARAFAC2.” | Apr 2024 |
| Systems Immunology in Transplantation: Advances, Challenges, and Opportunities, Invited Oral Presentation “Integration of multi-modal and multidimensional analysis.” | Mar 2024 |
| Cancer Systems Biology Program, Invited Oral Presentation “Analysis and modeling of cancer drug responses using cell cycle phase-specific rate effects.” | Dec 2023 |
| Systems Biology Consortium for Infectious Diseases, Invited Oral Presentation “Developing integrative signatures across omics, studies, and diseases with tensor-based analysis.” | Sep 2023 |
| UCLA Bioinformatics Retreat, Invited Oral Presentation “Building the tensor learning universe.” | Jul 2023 |
| Antibodies & Complement, Selected Oral Presentation “Cancer systems serology reveals active humoral immunity but disrupted Fc-elicited interactions.” | Jun 2023 |
| CSBC Annual Meeting, Invited Oral Presentation “Phosphoproteomic Analysis of AXL Identifies YAP as a Key Regulator of Resistance.” | Mar 2023 |
| SIAM Conference on Mathematics of Data Science, Invited Podium Presentation “Mechanistic and Data-Driven Dissection of Cell Communication Through Tensor Methods.” | Sep 2022 |
| UCLA Musculoskeletal Devices & Tech. Development Group, Invited Seminar “Uncovering immunologic mechanisms of MRSA persistence by tensor-mediated data integration.” | Sep 2022 |

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| Southern California Systems Biology Conference, Invited Podium Presentation "Mechanistic and Data-Driven Dissection of Cell Communication Through Tensor Methods." | Apr 2022 |
| American Assoc. for Cancer Research Annual Meeting, Invited Podium Presentation "Systems approaches for identifying cell states and pathways modulating therapy response." | Apr 2022 |
| Johns Hopkins Univ., Institute for Comp. Medicine, Invited Seminar "Mechanistic and Data-Driven Dissection of Cell Communication Through Tensor Methods." | Feb 2022 |
| Cellular & Molecular Bioengineering, Selected Oral Presentation "Rapid Prototyping of Multivalent And Multi-Specific Drugs To Overcome The Limited Selectivity Of IL-2 Toward Regulatory T Cells" | Jan 2022 |
| Biomedical Engineering Society Annual Meeting, Invited Podium Presentation "Tensor Factorization-Based Data Fusion Improves Predictions and Interpretation of MRSA Outcome." | Oct 2021 |
| Buffalo Quantitative Systems Pharmacology Symposium, Invited Speaker "Deeply profiling pharmacodynamic response with single cell dynamics." | Jul 2021 |
| CSHL Systems Immunology, Selected Oral Presentation "Developing a Mechanistic View of Mixed IgG Antibody Immune Effector Responses." | Apr 2021 |
| University of Massachusetts, Mol & Cell Biol Program, Invited Seminar "Mixture models of cell populations and signaling to understand heterogeneous drug response." | Mar 2021 |
| International Conference on Biomolecular Engineering, Selected Oral Presentation "Developing a Mechanistic View of Mixed IgG Antibody Immune Effector Responses." | Jan 2021 |
| Vanderbilt University, QSBC Center, Invited Center Seminar "Mixture models of cell populations and signaling to understand heterogeneous drug response." | Oct 2020 |
| Tufts University, Dept. of Bioengineering, Invited Dept. Seminar "Linking Statistical and Mechanistic Models for Drug Development." | Mar 2020 |
| Univ. of Calif., Los Angeles, Immunogenetics Center, Invited Speaker "Using models with incomplete information to study and engineer antibody effector response." | Jan 2020 |
| Biomedical Engineering Society Annual Meeting, Selected Oral Presentation "A Binding Model Predicts In Vivo Effector Cell-Elicited Killing Across Multiple Disease Models." | Oct 2019 |
| Xencor, Inc., Invited Oral Presentation "Computational molecular models for immune engineering." | Jul 2019 |
| CSBC West Coast Meeting, Selected Oral Presentation "Hidden Markov models on a tree as a general approach to single cell plasticity analysis." | May 2019 |
| Antibodies & Complement, Selected Oral Presentation "A Multivalent Binding Model Predicts FcγR Regulation and Effector Cell-Elicited Killing." | May 2019 |

RESEARCH SUPERVISION

Postdoctoral Fellows

- Catera Wilder, Ph.D. (Assistant Professor, UCSF) 2018–2022
- Song Yi Bae, Ph.D. (Senior Scientist, Astrin Biosciences) 2016–2019
- Edward Richards, Ph.D. (Senior Scientist, Dragonfly Therapeutics)
 - American Cancer Society Postdoctoral Fellowship
- Annelien Zweemer, Ph.D. (Assistant Professor, Leiden University) 2014–2017

Ph.D. Students

- Meera Trisal 2023–Present
- Michelle Loui 2022–Present
 - SURF Fellowship, UCLA Graduate Division
- Andrew Ramirez 2021–Present
 - NSF Graduate Research Fellowship
 - Cota Robles Fellowship
 - UCLA EDI Student Leadership Award
- Jackson Chin 2020–2025
 - Best Poster Award, QC Bio Retreat, 2022
- Brian Orcutt-Jahns (Postdoctoral Associate, Genentech) 2019–2024
 - Best Poster Award, CSBC Junior Investigator Meeting
 - Best Poster Award, Cytokine Society
 - Best Presentation Award, Los Angeles Bioscience Ecosystem Summit
 - Outstanding Ph.D. Award, Department of Bioengineering
- Cyrus Tan 2019–2024
 - Dissertation Year Fellowship, UCLA Graduate Division
- Farnaz Mohammadi (Postdoctoral Associate, Genentech) 2018–2023
 - Dissertation Year Fellowship, UCLA Graduate Division
- Marc Creixell (Scientist, Calico Life Sciences) 2018–2023
 - JCCC Fellowship

M.S. Students

- Het Desai 2023–2024
- Manmeet Bains 2023–2024
- Enio Hodzic (Machine Learning Algorithm Engineer, Adaptive Dynamics) 2021–2023
- Madeleine Murphy (Computational Biologist, Broad Institute) 2020–2022

Undergraduate Students

- Jamie Stickelmaier 2021–2023
- Ethan Hung (Amgen Scholar, Berkeley) 2021–2024
- Eva Hunter 2021–2022
- Hakan Alpay (Frontend Engineer, Facebook) 2021
- Luka Karginov (NCI CSBC Summer Scholar; Ph.D., Biological Engineering, MIT) 2020–2021
- Aditya Sivakumar 2020–2021
- Eli Snyder (M.D., University of Hawaii) 2020–2021
- Peter Emmel 2019–2022
- Amanda Tsao (M.D., University of Southern California) 2019–2021
- JC Lagarde 2019–2022
- Sumedha Kanthamneni (Google) 2019–2022
- Heather Carmen Mercieca (Amgen Scholar) 2019
- Linnet Chang (Analyst, Accenture) 2018–2021
- Stephen Lees (Ph.D., Biomedical Engineering, UVA) 2018–2021
- Zoe Kim (Engineer, GaN Corporation) 2018–2020
- Micah Bryant (M.S., Mechanical Engineering, UCSD) 2018–2020
- Robby Theisen (Ph.D., Biomedical Engineering, University of Michigan) 2018–2020
- Alison Tran (Biosciences Account Manager, Thermo Fisher Scientific) 2018–2020
- Willie Wu (Software Engineer, Rivian) 2018–2019
- Katrina Warner (Amgen Scholar; Ph.D., Biomedical Sciences, Harvard) 2018
- Donya Khashayar (Transfer Student Summer Research Program) 2018
- Rui Yan (Ph.D., ICME, Stanford)
 - Cathy Bank Scholarship2017–2019
- Ali Farhat (M.D./Ph.D., U Illinois) 2017–2019

- Rose Hills Foundation Scholar
- Adam Weiner (Ph.D., Tri-Institute CompBio) 2017–2019
 - Internet Research Initiative Award
- Ning Guan (Ph.D., Systems Biology, Harvard) 2015–2017
- Ryan Robinett (Ph.D., Comp. Sci., U. Chicago) 2015–2017
 - National Science Foundation Graduate Research Fellowship

SERVICE

Profession

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| Co-Chair, <i>Modeling Working Group, Systems Biology for Infectious Diseases Consortium</i> | 2023–Present |
| Webmaster, <i>BME Underrepresented Needs In Technology & Engineering (UNITE)</i> | 2022–Present |
| Co-Organizer, <i>BME UNITE Webinar Series</i> | 2021–Present |
| Chalk Talk Mentor, <i>BME Underrepresented Needs In Technology & Engineering (UNITE)</i> | 2024–Present |
| Organizer, <i>Sys. Immunol. Approaches in Transplantation Tolerance and Rejection Conference</i> | 2024–2025 |
| Poster Judge, <i>Immunology LA Conference</i> | 2024 |
| Ad Hoc External Tenure Case Evaluator, <i>University of California, San Diego</i> | 2024 |
| Ad Hoc Reviewer, <i>Cell</i> | 2024 |
| Ad Hoc Reviewer, <i>La Matematica</i> | 2024 |
| Reviewer, <i>National Centre of Competence in Research, Swiss National Science Foundation</i> | 2024 |
| Ad Hoc Reviewer, <i>Genome Medicine</i> | 2024 |
| Ad Hoc Study Section, <i>NCI Human Tumor Atlas Network</i> | 2024 |
| Ad Hoc Reviewer, <i>Metabolomics</i> | 2023 |
| Ad Hoc Reviewer, <i>Cancer Gene Therapy</i> | 2023 |
| Ad Hoc Reviewer (4x), <i>Science Signaling</i> | 2020–2023 |
| Ad Hoc Reviewer (3x), <i>Science Advances</i> | 2020–2023 |
| Track Chair: Computational & Systems Biology, <i>Biomedical Engineering Society Annual Meeting</i> | 2023 |
| Abstract Reviewer, <i>UC Systemwide Bioengineering Symposium</i> | 2023 |
| F31 Co-Sponsor, <i>Mollie Harrison (Advisor Stephanie Seidlits, University of Texas, Austin)</i> | 2022 |
| Ad Hoc Reviewer, <i>Cancer Immunology Research</i> | 2022 |
| Co-Chair, <i>Resource & Data Sharing Working Group, Cancer Systems Biology Consortium</i> | 2022–2024 |
| Ad Hoc Reviewer, <i>iScience</i> | 2022 |
| Poster Judge, <i>Cellular & Molecular Bioengineering Meeting</i> | 2022 |
| Example U01 proposal, <i>NIH National Institute of Allergy and Infectious Diseases</i> | 2021 |
| Reviewer, <i>Australia Medical Research Future Fund</i> | 2022 |
| Local Organizing Committee, <i>Southern California Systems Biology Conference</i> | 2022 |
| Ad Hoc Reviewer, <i>Soft Matter</i> | 2022 |
| Ad Hoc Reviewer, <i>FEBS Letters</i> | 2021 |
| Abstract Reviewer, <i>Biomedical Engineering Society Annual Meeting</i> | 2021, 2022 |
| Financial Officer, <i>Association of Cancer Systems Biologists</i> | 2021–Present |
| Session Co-Chair, <i>Biomedical Engineering Society Annual Meeting</i> | 2020 |
| Volunteer Speed Interviewer, <i>Biomedical Engineering Society Annual Meeting</i> | 2020 |
| Volunteer Resume Reviewer, <i>Biomedical Engineering Society Annual Meeting</i> | 2020 |
| Member, <i>BME Underrepresented Needs In Technology & Engineering (UNITE)</i> | 2020–Present |
| Panelist, <i>Amgen Scholars Summer Science Series</i> | 2020 |
| Ad Hoc Reviewer, <i>PLOS Biology</i> | 2020, 2021 |
| Ad Hoc Reviewer, <i>Cancer Research</i> | 2020 |
| External Reviewer, <i>Ming Hsieh Institute, USC</i> | 2020 |
| Ad Hoc Reviewer, <i>Cell Systems</i> | 2020 |
| Ad Hoc Reviewer, <i>APL Bioengineering</i> | 2020 |
| Ad Hoc Reviewer, <i>Integrative Biology</i> | 2019 |
| Ad Hoc Reviewer, <i>Scientific Reports</i> | 2019 |
| Ad Hoc Reviewer, <i>PNAS</i> | 2019 |
| Ad Hoc Reviewer, <i>Current Opinion in Systems Biology</i> | 2019 |

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| Co-Chair, Association of Cancer Systems Biologists | 2017–2021 |
| Ad Hoc Reviewer, <i>PLOS Computational Biology</i> | 2018, 2024 |
| Interviewee, <i>Prescriber Magazine</i> | 2017 |
| Ad Hoc Reviewer, <i>WIREs Systems Biology and Medicine</i> | 2017 |
| Ad Hoc Remote Reviewer, <i>Irish Research Council</i> | 2017 |
| Ad Hoc Reviewer, <i>Cell Reports</i> | 2017, 2023 |
| Graduate Research Fellowship Program Review Panelist, <i>National Science Foundation</i> | 2016–2017 |
| Meeting Organizer & Member, Association of Early Career Cancer Systems Biologists | 2015–2016 |
| Ad Hoc Reviewer, <i>Biomedical Engineering Society Annual Meeting</i> | 2016 |
| Ad Hoc Reviewer, <i>Drug Discovery Today</i> | 2016 |
| Ad Hoc Reviewer, <i>Molecular Cell</i> | 2015 |
| Member, <i>Biomedical Engineering Society</i> | 2010–Present |
| Coordinator, <i>MIT Biological Engineering Graduate Student Board</i> | 2010–2013 |
| Ad Hoc Reviewer, <i>Oncogene</i> | 2013 |
| Ad Hoc Reviewer, <i>Nature</i> | 2013 |
| Member, <i>MIT Biological Engineering Retreat Organizing Committee</i> | 2010–2012 |
| Ad Hoc Reviewer, <i>J. Cell Biol.</i> | 2011–2012 |

School & Campus

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|---|------------------------|
| Bioengineering Representative, HSSEAS GPU Ad Hoc Committee | 2024 |
| Member, <i>Hiring Committee, Hispanic Serving Institution STEM Faculty Director</i> | 2024 |
| Member, HSSEAS Strategic Planning Committee | 2024 |
| Chair, <i>Campus Response to the Climate Crisis Special Committee, Faculty Senate</i> | 2023–Present |
| Ad Hoc Member, <i>Executive Board, Faculty Senate</i> | 2023–Present |
| Faculty Speaker, <i>UCLA Life Sciences Webinar Series</i> | 2023 |
| Dean's Prize Poster Judge, <i>Undergraduate Research and Creativity Showcase</i> | 2023 |
| Poster Judge, <i>Jonsson Cancer Center Annual Retreat</i> | 2023 |
| Reviewer, <i>Tau Beta Pi Chapter Excellence Scholarship</i> | 2021–2023 |
| Selection Committee, <i>Faculty Career Development Award, Office of Equity, Diversity and Inclusion</i> | 2022 |
| Faculty Participant, <i>Coffee Chat Series, Computational & Systems Biology</i> | 2022 |
| Member, <i>Minors Committee, Computational & Systems Biology</i> | 2021–2023 |
| Panelist, <i>Graduate School Panel, Computational & Systems Biology</i> | 2021 |
| Faculty Representative, <i>Samueli Engineering Grad School Info Session</i> | 2020 |
| Faculty Representative, <i>Annual Biomedical Research Conference for Minority Students</i> | 2018, 2020 |
| Curriculum Advisory Committee, <i>Computational & Systems Biology</i> | 2020–Present |
| Written Qualifying Exam Evaluator, <i>Bioinformatics IDP</i> | 2020 |
| Mentor, <i>B.I.G. Summer</i> | 2020–2024 |
| Member, <i>SPUR "Life of a Faculty Member" Panel</i> | 2020 |
| Ad Hoc Member, <i>HSSEAS Faculty Executive Committee</i> | April 2020 |
| Application Reviewer, <i>Amgen Scholars Program</i> | 2020, 2022, 2023, 2024 |
| Reviewer, <i>Graduate Division's Faculty Review Committee</i> | 2020 |
| Faculty Volunteer, <i>Society of Women Engineers Recruitment Dinner</i> | 2019, 2020 |
| Member, <i>HSSEAS SEASnet Review Committee</i> | 2019 |
| Faculty Advisor, <i>Tau Beta Pi</i> | 2017–Present |
| Faculty Volunteer, <i>Amgen Scholars Symposium</i> | 2018–2024 |
| Member, <i>HSSEAS Awards Committee for Outstanding Student Awards</i> | 2018 |
| Faculty Speaker, <i>UCLA Engineering Alumni Reunion</i> | 2018 |

Department

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| Member, <i>Ad Hoc Promotion Committee</i> | 2023, 2024 |
| Poster Judge, <i>Bioengineering Research Day</i> | 2024 |
| Equity, Diversity, & Inclusion Panel Member, <i>Biomedical Engineering Society</i> | 2024 |
| Member, <i>Ad Hoc P&T Committee</i> | 2023 |
| Judge, <i>Biomedical Engineering Society BioHack</i> | 2022–2023 |

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| Member, <i>Diversity, Equity and Inclusion Committee</i> | 2021–2024 |
| Member, <i>Bioengineering Hiring Search Committee</i> | 2021–2022, 2023–2024, 2024–2025 |
| Member, <i>Teaching Facility & Shared Equipment Committee</i> | 2021–2024 |
| Member, <i>Strategic Planning Committee</i> | 2020 |
| Co-Chair, <i>Bioengineering and Computational Medicine Joint Hiring Search</i> | 2019–2020 |
| Field Chair, <i>Biosystem Science and Engineering</i> | 2019–Present |
| Co-Chair, <i>Graduate Admissions Committee</i> | 2019–Present |
| Member, <i>Undergraduate Curriculum Committee</i> | 2019–Present |
| Member, <i>Bioengineering Alumni Committee</i> | 2018–Present |
| Chair, <i>Department of Bioengineering Seminar Series</i> | 2018–2019 |
| Member, <i>Publicity Committee</i> | 2017–2018 |

PATENTS & DISCLOSURES

A.S. Meyer. "Methods of Identifying and Correcting Tumor Humoral Immune Dysregulation." U.S. patent application PCT/US24/32940, 2024.

Orcutt-Jahns, B., P.C. Emmel, A.S. Meyer. "Multi-specific engineered cytokines." U.S. patent application PCT/US24/32940, 2023.

A.S. Meyer. "Altering cytokine specificity through binding valency." U.S. patent application PCT/US22/35711, 2022.

Miller, M.A., M.J. Oudin, A.S. Meyer, L.G. Griffith, F.B. Gertler, D.A. Lauffenburger. "Methods of Reducing Kinase Inhibitor Resistance." US patent application 14/690,001, 2015.

THESIS COMMITTEE MEMBERSHIP

Daniel Bradbury, Bioengineering (Ph.D.)

Advisor: Daniel Kamei

2017–2020

Giovanni Valdez, Bioengineering (Ph.D.)

Advisor: Grace Xiao

2018–2021

Hiromi Miwa, Bioengineering (Ph.D.)

Advisor: Dino Di Carlo

2019–2022

Mark van Zee, Bioengineering (Ph.D.)

Advisor: Dino Di Carlo

2019–2022

Rob Dimatteo, Bioengineering (Ph.D.)

Advisor: Dino Di Carlo

2019–2021

Alexander Wickstrom, Bioengineering (M.S.)

Advisor: Jonathan Kao

2019–2019

Hector E Muñoz, Bioengineering (Ph.D.)

Advisor: Dino Di Carlo

2019–2020

Wei-Chia Elizabeth Luo, Bioengineering (Ph.D.)

Advisor: Gerard Wong

2020–2025

Mohammadali Alidoost, Bioengineering (Ph.D.)

Advisor: Jennifer Wilson

2021–2024

Cameron S. Movassaghi, Chemistry (Ph.D.)

Advisor: Anne M. Andrews

2021–2024

Felis Doyeon Koo, Bioengineering (Ph.D.)

Advisor: Dino Di Carlo

2021–2023

Favour Esedebé, Bioinformatics (Ph.D.)

Advisor: Tom Graeber

2021–2025

Connor Razma, Bioinformatics (M.S.)

Advisor: Alexander Hoffmann

2022–2023

Mai Tran, Earth, Planetary & Space Sci. (Ph.D.)

Advisor: William Newman

2022–2023

Nilay Shah, Computer Science (M.S.)

Advisor: Bolei Zhou

2022–2023

Rayo Suseno, Bioengineering (M.S.)

Advisor: Jennifer Wilson

2022–2023

Helen Huang, Bioinformatics (Ph.D.)
Advisor: Alexander Hoffman
2022–Present

Emily Bozich, Bioengineering (Ph.D.)
Advisor: Jennifer Wilson
2023–Present

Seth Hilliard, Comp. & Quant. Medicine (Ph.D.)
Advisor: Andrei Rodin (City of Hope)
2023–2023

Michael Mellody, Bioengineering (Ph.D.)
Advisor: Dino Di Carlo
2023–Present

Ahmed Ali, Bioengineering (M.S.)
Advisor: Jennifer Wilson
2023–2024

Alejandro Miron Jabalera, Bioengineering (Ph.D.)
Advisor: Tzung Hsiai
2024–Present

Calvin Lee, Bioengineering (M.S.)
Advisor: Roel Ophoff
2024–Present

Yuyang Han, Chemistry (Ph.D.)
Advisor: Anne Andrews
2024–Present

Shawn Liu, Bioengineering (M.S.)
Advisor: Jennifer Wilson
2023–2024

Jingwen Sun, Chemistry & Biochemistry (Ph.D.)
Advisor: Chong Liu
2023–Present

Frances Nicklen, Bioengineering (Ph.D.)
Advisor: Daniel Kamei
2023–Present

Citra Soemardy, Bioengineering (Ph.D.)
Advisor: Dino Di Carlo
2023–Present

Shivani Kumar, Bioengineering (M.S.)
Advisor: Mireille Kamariza
2024–2024

Katarina Reid, Bioengineering (M.S.)
Advisor: Joseph DiStefano
2024–2025

Dexter Lai, Bioengineering (Ph.D.)
Advisor: Jun Park
2024–Present