

Adelaide Tovar, Ph.D.

Postdoctoral Fellow

Gilbert S. Omenn Department of Computational Medicine & Bioinformatics

University of Michigan

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EDUCATION

2015 – 2021 **Ph.D., Genetics & Molecular Biology**
The University of North Carolina at Chapel Hill, Chapel Hill, NC

2011 – 2015 **S.B. in Course 7 (Biology)**
Massachusetts Institute of Technology, Cambridge, MA

RESEARCH TRAINING

04/2021 – present **Postdoctoral Fellow**, Gilbert S. Omenn Department of Computational Medicine and Bioinformatics, University of Michigan
Advisors: Drs. Steve Parker, Ph.D. and Jacob Kitzman, Ph.D.
Project: Developing and applying novel approaches to screen disease-associated noncoding variants

08/2015 – 02/2021 **Graduate Student**, Department of Genetics, The University of North Carolina at Chapel Hill
Advisor: Dr. Samir Kelada, M.P.H., Ph.D.
Dissertation: “Dissecting respiratory responses to ozone exposure with genetics and genomics”

08/2013 – 07/2015 **Undergraduate Research Assistant**, Department of Biological Engineering, Massachusetts Institute of Technology
Advisors: Drs. Darrell Irvine, Ph.D. and Gregory Szeto, Ph.D.
Thesis: “Developing an improved immunotherapy for late-stage cancers by engineered immunomodulation”

06/2013 – 08/2013 **Undergraduate Summer Researcher**, Department of Biology, University of North Texas
Advisor: Dr. Pamela Padilla, Ph.D.
Project: Identifying gene-by-diet interactions involved in responses to anoxia/hypoxia in *C. elegans*

FUNDING

09/2024 – 08/2029 **MOSAIC Pathway to Independence Award (K99/R00)**, NIH/NHGRI
K99 HG013676
Title: *Using the continuum of genetic causality to investigate trans regulatory mechanisms*

02/2023, 09/2023 **UROP Supplemental Funding**, University of Michigan

09/2022 – 08/2025 **Postdoctoral Diversity Enrichment Program**, Burroughs Wellcome Fund
Title: *Dissecting context-specific gene regulatory grammars*

07/2022 – 06/2024 **REACH Loan Repayment Program**, NIH/NIDDK
L70 DK134031
Title: *Annotating the environmental and cellular context of diabetes-associated variants with high-throughput sequencing and reporter assays*

05/2022 – 08/2024	Opportunity Pool Funding , NIDDK/Accelerating Medicines Partnership Program for Common Metabolic Diseases <i>A consortium-wide modular MPRA resource to assess cellular contexts of T2D variants</i>
09/2021 – 08/2023	T32 Postdoctoral Training Program in Basic Diabetes Research , NIH/NIDDK <i>Annotating the environmental and cellular context of diabetes-associated variants</i>
08/2020 – 05/2021	Dissertation Completion Fellowship , UNC Graduate School
11/2018, 09/2019	Graduate Student Travel Scholarship , International Mammalian Genome Society
02/2019	Transportation Grant , UNC Graduate School
10/2016 – 10/2019	Research Supplement to Promote Diversity in Health-Related Research , NIH/NIEHS <i>Administered under parent grant R01 ES024965 awarded to PI Kelada</i>

HONORS AND AWARDS

2025	Semi-finalist , ASHG/Charles J. Epstein Trainee Award for Excellence in Human Genetics Research
2025	Emerging Scholar in Genome Sciences , University of Virginia Department of Genome Sciences
2023	Postdoctoral Leadership Academy , SACNAS
2023	Rising Star in Genetics and Genomics , University of Utah Department of Human Genetics
2022	Human Genetics Scholar , ASHG
2019, 2020	Outstanding Poster Presentation Award , International Mammalian Genome Society
2020	Fellow , Yale Ciencia Academy for Career Development Fellow
2019	First Place Poster Presentation Award , UNC Initiative for Maximizing Student Development
2019	Minority Trainee Development Scholarship , American Thoracic Society
2015	S. Klein First Prize in Technical Writing , MIT
2015	Award Winner , Wellcome Image Awards (received jointly with Gregory Szeto and Jeffrey Wyckoff)
2015	Winner , MIT Koch Institute for Integrative Cancer Research Image Awards (received jointly with Gregory Szeto and Jeffrey Wyckoff)

PREPRINTS AND PUBLICATIONS

*Equal contribution is highlighted with **

1. Nishino K, Kitzman JO, Parker SCJ, **Tovar A**. Functional dissection of metabolic trait-associated gene regulation in steady state and stimulated human skeletal muscle cells. *bioRxiv*. 2024 Nov; 625886. [10.1101/2024.11.28.625886](https://doi.org/10.1101/2024.11.28.625886)
2. Varshney A, Manickam N, Orchard P, **Tovar A**, Zhang Z, Feng F, Erdos MR, Narisu N, Ventresca C, Nishino K, Rai V, Stringham HM, Jackson AU, Tamsen T, Gao C, Koues OI, Welch JD, Burant CF, Williams LK, Jenkinson C, DeFronzo RA, Norton L, Saramies J, Lakka TA, Laakso M, Tuomilehto J, Mohlke KL, Kitzman JO, Koistinen HA, Liu J, Boehnke M, Collins FS, Scott LJ, Parker SCJ. Population-scale skeletal muscle single-nucleus multi-omic profiling reveals extensive context specific gene regulation. *bioRxiv*. 2023 Dec; 571696. [10.1101/2023.12.15.571696](https://doi.org/10.1101/2023.12.15.571696). *In revision at Nature Genetics*.
3. **Tovar A**, Kyono Y, Nishino K, Bose M, Varshney A, Parker SCJ, Kitzman JO. Using a massively parallel reporter assay to discover context-dependent regulatory activity in type 2 diabetes-linked noncoding

regions. *bioRxiv*. 2023 Oct; 561391. [10.1101/2023.10.08.561391](https://doi.org/10.1101/2023.10.08.561391). *In review at HGG Advances*.

4. **Tovar A**, Smith GJ, Thomas JM, McFadden KM, Kelada SNP. A locus on chromosome 15 contributes to acute ozone-induced lung injury in Collaborative Cross mice. *Am J Respir Cell Mol Biol*. 2022 Nov;67(5):528-538. [10.1165/rcmb.2021-0326OC](https://doi.org/10.1165/rcmb.2021-0326OC)
5. **Tovar A***, Crouse WL*, Smith GJ, Thomas JM, Keith BP, McFadden KM, Moran TP, Furey TS, Kelada SNP. Integrative phenotypic and genomic analyses reveal strain-dependent responses to acute ozone exposure and their associations with airway macrophage transcriptional activity. *Am J Physiol Lung Cell Mol Physiol*. 2022 Jan 1;322(1):L33-L49. [10.1152/ajplung.00237.2021](https://doi.org/10.1152/ajplung.00237.2021)
6. Smith GJ, **Tovar A**, McFadden KM, Moran TP, Wagner JG, Harkema JR, Kelada SNP. A model of ozone-induced non-atopic asthma from the Collaborative Cross. *Am J Respir Cell Mol Biol*. 2021 Dec;65(6):672-674. [10.1165/rcmb.2020-0577LE](https://doi.org/10.1165/rcmb.2020-0577LE)
7. Smith GJ, **Tovar A**, Kanke M, Wang Y, Deshane JS, Sethupathy P, Kelada SNP. Ozone-induced changes in the murine lung extracellular vesicle small RNA landscape. *Physiol Rep*. 2021 Sep;9(18):e15054. [10.14814/phy2.15054](https://doi.org/10.14814/phy2.15054)
8. Lauder milk LT, **Tovar A**, Homstad AK, Thomas JM, McFadden KM, Tune MK, Cowley DO, Mock JR, Ideraabdullah F, Kelada SNP. Baseline and innate immune response characterization in the *Zfp30* knockout mouse strain. *Mamm Genome*. 2020 Aug;31(7-8):205-214. [10.1007/s00335-020-09847-z](https://doi.org/10.1007/s00335-020-09847-z)
9. **Tovar A***, Smith GJ*, Thomas JM, Crouse WL, Harkema J, Kelada SNP. Transcriptional profiling of the murine airway response to acute ozone exposure. *Toxicol Sci*. 2020 Jan 1;173(1):114-130. [10.1093/toxsci/kfz219](https://doi.org/10.1093/toxsci/kfz219)
10. Weiser M, Simon JM, Kochar B, **Tovar A**, Israel JW, Robinson A, Gipson GR, Schaner MS, Herfarth HH, Sartor RB, McGovern DPB, Rahbar R, Sadiq TS, Koruda MJ, Furey TS, Sheikh SZ. Molecular classification of Crohn's disease reveals two clinically relevant subtypes. *Gut*. 2018 Jan;67(1):36-42. [10.1136/gutjnl-2016-312518](https://doi.org/10.1136/gutjnl-2016-312518)

SELECTED PRESENTATIONS

Invited

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|---------|---|
| 09/2025 | 6th Annual Emerging Scholars in Genome Sciences , University of Virginia, Charlottesville, VA
Title: <i>Functional annotation of over 25,000 type 2 diabetes-associated variants across contexts</i> |
| 06/2024 | Diabetes Genetic Convergence – Rare to Common Variants in Networks , American Diabetes Association's 84 th Scientific Sessions, Orlando, FL
Title: <i>Exploring regulatory convergence in rare and common diabetes with massively parallel reporter assays</i> |
| 05/2023 | Rising Stars in Genetics and Genomics Symposium , University of Utah, Salt Lake City, UT
Title: <i>Using massively parallel reporter assays to uncover context-specific gene regulation in type 2 diabetes</i> |
| 06/2020 | Science in a Snapshot Seminar Series , Genetics Society of America, virtual
Title: <i>Gene-environment interactions underlie respiratory responses to the air pollutant ozone</i> |

Platform

10/2025	American Society of Human Genetics Annual Meeting , Boston, MA Title: <i>Functional annotation of over 25,000 type 2 diabetes-associated variants</i>
07/2025	Gordon Research Conference on Human Genetics and Genomics , Portland, ME Title: <i>Assessing cellular contexts of type 2 diabetes-associated variants at scale</i>
04/2025	Accelerating Medicines Partnership for Common Metabolic Disease Parliament Meeting , Philadelphia, PA Title: <i>Assessing cellular contexts of type 2 diabetes-associated variants at scale</i>
11/2024	American Society of Human Genetics Annual Meeting , Denver, CO Title: <i>Assessing cellular contexts of type 2 diabetes-associated variants at scale</i>
07/2023	Gordon Research Seminar on Human Genetics and Genomics , Waterville Valley, NH. Title: <i>Using massively parallel reporter assays to discover context-specific regulatory grammars</i>
03/2023	Accelerating Medicines Partnership for Common Metabolic Disease Parliament Meeting , Boston, MA Title: <i>A consortium-wide modular MPRA resource to assess cellular contexts of diabetes-associated variants</i>
06/2022	American Diabetes Association's 82nd Scientific Sessions , New Orleans, LA Title: <i>A modular massively parallel reporter assay uncovers context-specific activity of diabetes-associated regulatory elements</i>
09/2021	American Society of Human Genetics Annual Meeting , virtual Title: <i>A modular massively parallel reporter assay uncovers context-specific allelic activity of GWAS variants</i>
04/2020	The Allied Genetics Conference , virtual. Title: <i>A GxE QTL on Chr. 15 underlies susceptibility to air pollution-induced lung injury in mice</i>
05/2019	American Thoracic Society International Conference , Dallas, TX Title: <i>Investigating susceptibility to ozone-induced lung inflammation and injury using the Collaborative Cross genetic reference population</i>

MENTORSHIP

2023 – present	Mentor , Lovelyn Epelle, Ph.D. student in Human Genetics, University of Michigan
2023 – present	Mentor , Benjamin (Ben) Li, MSTP student in Bioinformatics, University of Michigan
2023 – present	Mentor , Miranda Jefferds, B.S. student in Biochemistry, University of Michigan
2023 – present	Mentor , Gabriella (Ellie) Bloss, B.S. student in Biochemistry, University of Michigan
2022 – 2025	Mentor , Hailey McMillen, B.S. in Biology, Health, and Society, University of Michigan <i>Currently: Medical assistant at University of Colorado Anschutz Medical Campus</i>

2022 – 2025	Mentor , Kirsten Nishino, B.S. in Molecular, Cellular, and Developmental Biology and B.S. in Life Science Informatics, University of Michigan <i>Currently: Ph.D. student in Genetics, Genomics, and Systems Biology at the University of Chicago</i>
2023 – 2024	Mentor , Sanjana Yadav, B.S. student in Computer Science, University of Michigan
2023	Mentor , Ronak Arun, B.S. in Biology, University of Michigan
2022 – 2023	Mentor , Elli Gasper, B.S. student in Biology, University of Michigan
2021 – 2022	Mentor , Maya Bose, Ph.D. student in Bioinformatics, University of Michigan
2019 – 2021	Mentor , Morgan Nalesnik, Ph.D. in Toxicology & Environmental Medicine
2018 – 2020	Mentor , Daniel Vargas, B.S. in Biology, UNC <i>Currently: Software engineer at The College Board</i>
2018 – 2019	Mentor , Jessica Bustamante, B.A. in Biology, UNC <i>Currently: Prenatal Genetic Counselor at Prisma Health</i>
2017	Mentor , Abbott Ndukwe, B.S. in Psychology, UNC <i>Currently: Project Manager at Cisco</i>

UNIVERSITY AND PROFESSIONAL SERVICE

Ad Hoc Reviewer

Genetics, Cell Genomics, eLife, PLoS Genetics

Abstract Reviewer

ASHG (2022, 2025), ADA (2023, 2024), ABRCMS (2022)

Application Reviewer

Intersections Science Fellows Symposium (2021), SACNAS (2021 – 2023)

Moderator/Session Chair

ASHG (2022, 2023), ADA (2024)

11/2024 – present	Committee Member , Atlas of Variant Effects Alliance Variant Effects Seminar Series
01/2022 – present	MPRA Working Group Chair , Accelerating Medicines Partnership for Common Metabolic Diseases
07/2019 – 08/2020	Chair , UNC Academic and Research-Intensive Careers Cohort
05/2019 – 05/2020	Family Leader , UNC Department of Genetics Network
05/2018 – 05/2020	Advisor , UNC University Career Services Pre-Graduate Educational Advising Program (PGEAP)
05/2018 – 05/2020	Peer Mentor , UNC BBSP First-Year Group
05/2019 – 08/2019	Member , UNC PGEAP Recruitment & Training Committee

TEACHING EXPERIENCE AND TRAINING

01/2024 – 04/2024	Postdoctoral Short Course on College Teaching in STEM , University of Michigan
08/2017 – 05/2020	Academic Coach/Tutor , UNC <i>GNET 621: Introduction to Genetic Analysis, GNET 632: Advanced Molecular Biology, BCB 720: Advanced Statistical Modeling</i>
01/2017 – 05/2017	Teaching Assistant , UNC GNET 632: Advanced Molecular Biology
11/2015 – 08/2016	Tutor , Tutor.com
09/2014 – 05/2015	Tutor , MIT Department of Biology
09/2014 – 12/2014	Tutor , InstaEDU
08/2013 – 05/2014	AP Biology Instructor , MIT Educational Studies Program

OUTREACH

08/2021 – 06/2022	Co-Chair, Education & DEI , Michigan DNA Day
08/2017 – 05/2018	Teacher , UNC DNA Day CONNECT
04/2016 – 04/2018	Ambassador , North Carolina DNA Day
04/2015	Volunteer , Cambridge Science Festival