

- **Research interests:** Leverage statistical and computational methods in diverse multi-omics data to better understand and predict complex human diseases.

ACADEMIC |
POSITIONS

Tenure-track Assistant Professor

Human Genetics Center

Department of Epidemiology

University of Texas Health Science Center School of Public Health

Houston, TX

2025 – present

EDUCATION |
& TRAINING

DARTMOUTH COLLEGE, Hanover, NH

Bachelor of Arts: *magna cum laude*

Advisor: Dorothy Wallace, Ph.D.

2014 - 2018

Mathematics

Biology, with high honors

HARVARD MEDICAL SCHOOL, Boston, MA

Doctor of Philosophy

Advisor: Chirag Patel, Ph.D.

2018 – 2022

Bioinformatics and Integrative Genomics

BROAD INSTITUTE, Cambridge, MA

Research Fellow, Massachusetts General Hospital

Analytical and Translational Genetics Unit

Primary advisor: Alicia Martin, Ph.D.

Secondary advisor: Alexander Gusev, Ph.D.

2022 – 2024

MASSACHUSETTS GENERAL HOSPITAL, Boston, MA

NIH Precision and Genomic Medicine T32 Fellow

Center for Genomic Medicine

Directors: Jordan Smoller, MD, ScD, Heidi Rehm, PhD, FACMG

2023 – 2024

AWARDS |

ASHG Trainee Research Excellence Awards (Semi-finalist), American Society of Human Genetics

Early Career Investigator Program (ECIP) Award, World Congress of Psychiatric Genetics

Eliana Hechter Memorial Prize, The Broad Institute

“Reviewers’ Choice” Award (Top 10%), American Society of Human Genetics

Early Career Researcher Award, Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium

Early Career Researcher Scholarship, International Conference in Quantitative Genomics

“Reviewers’ Choice” Award (Top 10%), American Society of Human Genetics

Epstein Award (Semi-finalist), American Society of Human Genetics

2024

2024

2023

2023

2022

2020

2020

2020

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| UK Biobank Early-Career Researcher of the Year (Semi-finalist), UK Biobank | 2019 |
| Program in Quantitative Genomics (PQG) Student/Postdoc Travel Fund, Harvard School of Public Health | 2019 |
| Travel Fund, Harvard Medical School | 2018 |
| Christopher G. Reed Biologist Award, Dartmouth College Biology Department | 2018 |
| MAA Outstanding Presentation Award (Top 5%), Mathematical Association of America Mathfest | 2017 |
| Rufus Choate Scholar (Top 5% GPA), Dartmouth College | 2017 |
| First Place AWM Essay Contest, American Women in Mathematics Association, Math for America | 2017 |
| Mathematics Department Poster Award (1 st place), Dartmouth College. | 2016 |
| Wetterhahn Library Research Award, Dartmouth College | 2016 |
| James O. Freedman Presidential Scholar, Dartmouth College | 2016 |

GRANTS |

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|---|-----------|---------|
| Mass General Brigham (MGB) Precision and Genomic Medicine Training Grant T32; 5T32HG010464, PI: Jordan Smoller, Heidi Rehm | \$56,880 | 2023–24 |
| NSF Graduate Research Fellowship Program | \$102,000 | 2020–22 |
| NHGRI Institutional National Training Grant T32; 5T32HG002295-17, PI: Peter Park | \$73,464 | 2018–20 |
| AWS Cloud Credits, Amazon Web Services Research Grants Support for research through AWS promotional credits applicable to AWS services | \$2000 | 2018 |
| Student Experiential Learning Fund, Center for Professional Development and the Dartmouth Center for the Advancement of Learning Fund to support “students’ skill, confidence, and capacity building through unpaid or low- paid, intentional experiences” | \$5000 | 2017 |
| Hellman Family Foundation Research Grant, Dartmouth College Support for full-time leave term research given to “projects in which students are intellectually engaged and playing a key role in the project” | \$5,200 | 2017 |
| Neukom Travel Grant, William H. Neukom Institute for Computational Science Awarded to “excellent and deserving proposals” to students who presenting their research at conferences, workshops, or symposia | \$1,000 | 2017 |
| Rockefeller Mini Grant, Nelson A. Rockefeller Center for Public Policy and the Social Sciences | \$300 | 2017 |

PUBLICATIONS |

**Denotes co-primary authorship*

He Y, Lu W, Jee YH, Wang Y, Tsuo K, Qian DC, Diao JA, Huang H, Patel CJ, Byun J, Pasaniuc B, Atkinson EG, Amos, CI, Moll M, Cho MH, Martin “Multi-trait and multi-ancestry genetic analysis of comorbid lung diseases and traits improves genetic discovery and polygenic risk prediction”. *Nature Genetics* (under revisions). DOI: 10.1101/2024.08.25.24312558

He Y*, Groha S*, Taraszka K*, Lakhani CM, Braunstein L, Foulkes W, Polak P, King D, Tell R, White K, Zaitlen N, Patel CJ, Gusev A. “Genetic Ancestry and Population Differences in Somatic Alterations and Clinical Outcomes for Five Common Cancers”. *Nature Genetics*. (under revision)

Choi KW*, Tubbs J*, Lee H, He Y, Tsuo K, Yohannes M, Madsen E, Ghimire D, Hermosilla S, Ge T, Martin AR, Axinn W, Smoller JW. “Genetic Architecture and Socio-Environmental Risk Factors for Major Depressive Disorder in Nepal”. *Psychological Medicine* (2024) DOI: 10.1017/S0033291724001284

Diao JA, He Y, Khazanchi R, Tiako MJN, Witonsky JI, Pierson E, Rajpurkar P, Elhawary JR, Melas-Kyriazi L, Yen A, Martin AR, Levy S, Patel CJ, Farhat M, Borrell LN, Cho M, Silverman EK, Burchard EG, Manrai AK. “Implications of

Race Adjustment in Lung Function Equations". *New England Journal of Medicine* (2024) DOI: 10.1056/NEJMsa2311809

Wang Y*, He Y*, Shi Y, Qian DC, Gray KJ, Winn R, Martin AR "Boldly aspiring towards equitable benefits from genomic advances to individuals of ancestrally diverse backgrounds". *The American Journal of Human Genetics* (2024) DOI: 10.1016/j.ajhg.2024.04.002

Ciobanu O, He Y, Martin AR, Remick JS, Shelton JW, Eng TY, Qian CQ. "Patterns of under- and over-treatment in adjuvant radiotherapy for early-stage endometrial cancer based on molecular classification". *JAMA Oncology* (2024) DOI:10.1001/jamaoncol.2024.0104.

He Y, Martin AR. "We need more-diverse biobanks to improve behavioural genetics". *Nature Human Behavior* (2023). DOI: <https://doi.org/10.1038/s41562-0D23-01795-3>

He Y, Qian DC, Diao JA, Cho MH, Silverman EK, Gusev A, Manrai AK, Martin AR, Patel CJ "Prediction and stratification of longitudinal risk for chronic obstructive pulmonary disease across smoking behaviors". *Nature Communications* (2023). DOI: 10.1038/s41467-023-44047-8

He Y, Patel CJ. "Software Application Profile: PXStools -- an R package of tools for conducting exposure-wide analysis and deriving polyexposure risk scores". *International Journal of Epidemiology* (2022). DOI: 10.1093/ije/dyac216

He Y, Patel CJ. "Polygenic and polyexposure risks of type 2 diabetes predict chronic conditions in participants of the UK Biobank". *Acta Diabetologica* (2022). DOI: 10.1007/s00592-022-01864-5

He Y, Lakhani CM, Rasooly D, Manrai AK, Tzoulaki I, Patel CJ. "Comparisons of Polyexposure, Polygenic, and Clinical Risk Scores in Risk Prediction of Type 2 Diabetes". *Diabetes Care* (2020) DOI: 10.2337/dc20-2049

Tierney BT, He Y, Church GM, Segal E, Kostic AD, Patel CJ. "The Predictive Power of the Microbiome Exceeds That of Genome-Wide Association Studies in the Discrimination of Complex Human Disease". *bioRxiv* (2020). DOI: 10.1101/2019.12.31.891978

He Y, Kodali A, Wallace DL. "Predictive Modeling of Neuroblastoma Growth Dynamics in Xenograft Model after Anti-VEGF Therapy". *Bulletin of Mathematical Biology* (2018). DOI: 10.1007/s11538-018-0441-3

Qian DC, Jin JL, Titus AJ, He Y, Vaissie M, Li Y, Molfese DL, Salas R, Amos CI. "Genome-wide imaging association study implicates functional activity and glial homeostasis of the caudate in smoking addiction". *BMC Genomics* (2017). DOI: 10.1186/s12864-017-4124-5

PRESENTATIONS |

INVITED TALKS

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| Festival of Genomics & Biodata; Boston, MA | 06/2024 |
| "Integration of polygenic risk scores with social health for precision medicine" | |
| Feigin Rising Stars Symposium, Baylor College of Medicine; Houston, TX | 01/2024 |
| "Big data for precision medicine: an integrative gene & environment approach in global biobanks" | |
| Center for Genomic Medicine Seminar Series, Massachusetts General Hospital; Boston, MA | 11/2023 |
| "Comprehensive analyses of genetic and environmental factors in global biobanks" | |
| Faculty Retirement Conference, Dartmouth College; Hanover, NH | 06/2023 |
| "Prediction and stratification of disease risk through integrative multi-omics analysis of global biobanks" | |

SELECTED PLATFORM SPEAKER

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| ASHG Annual Meeting, Denver, CO | 11/2024 |
| "Multi-trait and multi-ancestry genetic analysis of comorbid lung diseases and traits improves genetic discovery and polygenic risk prediction" | |
| He Y, Lu W, Jee YH, Wang Y, Tsuo K, Moll M, Cho MH, Martin AR | |
| 7 th International Conference of Quantitative Genomics (ICQG7) | 07/2024 |

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| “Multi-trait and multi-ancestry polygenic risk score approach improves genetic discovery and risk prediction of respiratory diseases” | |
| He Y, Lu W, Jee YH, Wang Y, Tsuo K, Moll M, Cho MH, Martin AR | |
| STATGEN 2024: Conference on Statistics in Genomics and Genetics; Pittsburgh, PA | 05/2024 |
| “Multi-trait and multi-ancestry prediction of respiratory diseases”. | |
| He Y, Wang Y, Jee YH, Y Matthew M, Tsuo K, Cho MH, Martin AR | |
| IGES Annual Meeting; Paris, France | 08/2022 |
| “Polyexposure risk score offers greater predictive performance for COPD than polygenic score or smoking” | |
| He Y, Patel CJ | |
| ASHG Annual Meeting, Virtual | 10/2020 |
| “Comparisons of Polyexposure, Polygenic, and Clinical Risk Scores in Risk Prediction of Type 2 Diabetes” | |
| He Y, Lakhani CM, Rasooly D, Manrai AK, Tzoulaki I, Patel CJ | |
| UK Biobank Annual Meeting; London, UK | 07/2019 |
| “Poly-Exposure Scores for Disease Prediction” | |
| He Y, Lakhani CM, Manrai AK, Patel CJ | |
| NHGRI Research Training and Career Development Meeting; St. Louis, MO | 04/2019 |
| “Poly-Exposure Scores for Disease Prediction” | |
| He Y, Lakhani CM, Manrai AK, Patel CJ | |
| Society for Industrial and Applied Mathematics (SIAM) Annual Meeting; Pittsburgh, PA | 06/2017 |
| “Mathematical Model of Tumor Growth <i>In Vivo</i> ” | |
| He Y, Kodali A, Wallace, DI | |
| POSTER PRESENTATIONS | |
| Symposium in Artificial Intelligence for Learning Health Systems; Puerto Rico. | 05/2024 |
| “SBG-LUNG: Integrating raw spirogram, blood markers, and genetic data in a hybrid learning model for lung cancer prediction” | |
| He Y, Diao JA, Martin AR, Manrai AK, Qian DC, Patel CJ | |
| American Human Genetics Conference; Washington, D.C. | 11/2023 |
| “Comprehensive polygenic prediction of respiratory diseases: a cross biobank multi-trait and multi-ancestry approach” | |
| He Y, Wang Y, Tsuo K, Martin AR | |
| European Human Genetics Conference; Glasgow, Scotland, UK | 06/2023 |
| “Socioeconomic and environmental risk assessment of chronic obstructive pulmonary disease across smoking behaviors and populations” | |
| He Y, Martin AR, Patel CJ | |
| CHARGE Annual Meeting; Philadelphia, PA | 04/2022 |
| “PXStools: an R package for conducting exposure-wide analysis and deriving polyexposure risk scores” | |
| He Y, Patel CJ | |
| American Society of Human Genetics Annual Meeting; Houston, TX | 10/2018 |
| “Genetic Ancestry Differences in Oncogenic Mutations and Driver Genes” | |
| He Y, Gusev A | |

TEACHING |

COURSE DEVELOPMENT

Curriculum that I have designed (lectures, problems sets, etc.) and taught

Global Initiative for Neuropsychiatric Genetics Education in Research (GINGER) **2023**

Member of the core teaching team at GINGER which aims to build neuropsychiatric genetics research capacity in low and middle- income countries. Designed and recorded lectures on fundamental of genome wide association studies and polygenic risk scores.

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| Topics in Translational Biomedical Informatics (BMI 722) Tutorials | 2019 F, 2020 F |
| Harvard Medical School | |
| Designed and led introductory sessions for using R and Python to supplement students taking BMI 722. | |
| Constructed and graded homework for coding tutorials. | |

GUEST LECTURES

Invited lecturer

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| BMI 704: Data Science for Medical Decision Making , Harvard Medical School | 2021 |
| Lecture on the principals of constructing environmental risk score for disease prediction. | |
| BMI 722: Topics in Translational Biomedical Informatics , Harvard Medical School | 2020 |
| Introductory lecture on how to conduct genetic-enviroment interaction studies. | |
| Medlytics: Data Science for Health and Medicine , MIT Beaverworks Summer Institute | 2019 |
| Overview of current research in biomedical data science, especially research applications in healthcare. | |

TEACHING SUPPORT

Designed / graded problem sets, led office hours and discussion sections

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| BMI 704: Data Science for Medical Decision Making , Harvard Medical School | 2020 S, 2021 S |
| Head TA for 2021: managed team of 3 teaching fellows | |
| BMI 722: Topics in Translational Biomedical Informatics , Harvard Medical School | 2019 – 2021 F |
| Head TA for 2020, 2021: managed team of 3-4 teaching fellows | |
| BMI 706: Data Visualization , Harvard Medical School | 2021 S |
| High Impact Cancer Research Postgraduate Certificate Program , Harvard Medical School | 2019 – 2020 |
| Member of the teaching support team for a year-long program on cancer research for >50 students across the world | |
| Biology 2: Human Biology , Dartmouth College | 2017 F |
| Biology 13: Genetics , Dartmouth College | 2016 F |
| Learning fellow: received extensive training in teaching, learning, and pedagogy as well as strategies for assisting with cooperative learning in the classroom | |

TUTORING

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| Peer Tutor, Tutor Clearinghouse , Dartmouth College | 2014 – 2017 |
| Provided one-on-one peer tutoring for over 15 students in: Multivariable Calculus, Genetics, General Chemistry, Mathematical Biology | |

RESEARCH

Natalia Morado-Mata (current undergraduate student at The University of Texas at San Antonio)

Aashna Shah (current PhD student at Harvard University)

Faith Adams (current PhD student at Icahn School of Medicine)

Pauline Gabrieli, while as a master's student at Harvard University (current medical student at Harvard Medical School)

MENTORSHIP

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| Lily Wang, while as an PhD student at Harvard University (through DBMI) | 2021 |
| Andrea Liu, while as an undergraduate student at Harvard University (through WiSTEM) | 2020 |
| Jena Lorman, while as an undergraduate student at Harvard University (through WiSTEM) | 2019 |
| Theia Qu, while as an undergraduate4 student at Dartmouth College (through WISP) | 2017-2018 |
| Paula Mendoza, while as an undergraduate student at Dartmouth College (through WISP) | 2016-2018 |

SERVICE |

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| <i>Ad-Hoc Reviewer</i> | <i>Genome Biology</i> | 2025 |
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| <i>Ad-Hoc</i> Reviewer | <i>New England Journal of Medicine AI</i> | 2024 |
| <i>Ad-Hoc</i> Reviewer | <i>Journal of Translational Medicine</i> | 2023 |
| <i>Ad-Hoc</i> Reviewer | <i>Diabetes Care</i> | 2022 |
| <i>Ad-Hoc</i> Reviewer | <i>Acta Diabetologica</i> | 2021 |
| <i>Ad-Hoc</i> Reviewer | <i>Diabetologia</i> | 2021 |
| <i>Ad-Hoc</i> Reviewer | <i>European Heart Journal</i> | 2021 |
| Panelist | Harvard DBMI SIBMI Trainee Workshop | 2020, 2021 |
| Program Representative | Harvard Medical School and Harvard Medical Visiting Committee | 2021 |
| Program Representative | Harvard Integrative Life Sciences welcome event | 2020 |
| Co-Chair | Harvard Graduate Women in Science (HGWISE) mentoring program | 2019 – 2021 |
| Organizer | Monthly Research Seminar, Program in Bioinformatics and Integrative Genomics, Harvard Medical School | 2019 – 2020 |
| Planning Committee | Annual Retreat, Program in Bioinformatics and Integrative Genomics, Harvard Medical School | 2019 |
| Invited Panelist | Graduate Student Panel, Program in Bioinformatics and Integrative Genomics, Harvard Medical School | 2019 |
| Planning Committee | PhD Student Admissions Events, Program in Bioinformatics and Integrative Genomics, Harvard Medical School | 2018 – 2019 |

AFFILIATIONS |

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| Associate, Program in Quantitative Genomics | 2018 – 2022 |
| Member, Landry Cancer Biology Consortium | 2018 – 2022 |
| Member, Association for Women in Mathematics | 2016 – 2018 |
| Member, Gamma Sigma Alpha National Academic Greek Honor Society | 2016 – 2018 |