Rahul Gupta

| | Education and training | | |
|--------------|--|--|--|
| - | Internal Medicine Internship , <i>Massachusetts General Hospital</i> , Boston, MA Stanbury Physician-Scientist Pathway | | |
| 2017–Present | MD Candidate , <i>Harvard Medical School</i> , Boston, MA Harvard-MIT Program in Health Sciences and Technology (HST) | USMLE STEP 1: 259 USMLE STEP 2: 271 | |
| 2019–2023 | Doctor of Philosophy , <i>Harvard Medical School</i> , Boston, MA Biological and Biomedical Sciences, Program in Genetics and Genomics Co-advised by Drs. Vamsi Mootha & Benjamin Neale | | |
| 2012–2016 | Bachelor of Science in Engineering, University of Pennsylvania, Philadel School of Engineering and Applied Science (SEAS) Majors: Chemical & Biomolecular Engineering, Biology; Minor: Chemistry | lphia, PA, Summa Cum Laude GPA: 3.96/4.00 MCAT Score: 524/528 | |
| | Selected publications and preprints | (* indicates co-first author) | |
| 2024 | Karczewski, KJ*, Gupta, R* , Kanai, M*, et al. Pan-UK Biobank GWAS improves discovery, analysis of genetic architecture, and resolution into ancestry-enriched effects. <i>medRxiv</i> 2024. | | |
| 2023 | Gupta, R , Kanai, M, Durham, TJ, et al. Nuclear genetic control of mtDNA copy number and heteroplasmy in humans. <i>Nature</i> 2023. | | |
| 2021 | Gupta , R , Karczewski, KJ, Howrigan, D, et al. Human genetic analyses of organelles highlight the nucleus, but not the mitochondrion, in age-related trait heritability. <i>eLife</i> 2021. | | |
| 2020 | Rath S*, Sharma, R*, Gupta, R* , et al. MitoCarta3.0: an updated mitochondrial proteome now with sub-organelle localization and pathway annotations. <i>Nucleic Acids Res.</i> 2020. | | |
| 2018 | Jha, P*, McDevitt, MT*, Gupta, R* , et al. Systems Analyses Reveal Physiological Roles and Genetic Regulators of Liver Lipid Species. <i>Cell Systems</i> 2018. | | |
| 2017 | Gupta, R* , Lan, M*, Mojsilovic-Petrovic, J, et al. The Proline/Arginine Dipeptide from Hexanucleotide Repeat Expanded C9ORF72 Inhibits the Proteasome. <i>eNeuro</i> 2017. | | |
| | PhD Dissertation | | |
| 2019–2023 | Harvard Program in Biological and Biomedical Sciences (BBS), Genetics and Genomics Concen- tration, <i>Graduate student</i> , Boston MA | | |
| Advisors: | Vamsi K. Mootha, MD, Benjamin M. Neale, PhD | | |
| Title: | Nuclear genetic control of mitochondrial function and its contribution to human disease: insights at biobank scale | | |
| | Used statistical genetics at scale to gain basic biological insights about Discovered that the nuclear genome influences the abundance of common Created metabolomic models in patients with rare disorders to gain insights Analyzed the contributions of variation in mitochondria-localizing gene Assessed phenotype heritability across diverse populations in UK Bioba Developed and deployed large-scale computational pipelines using Goop | non human mtDNA mutations sights into common disease s to disease heritability nk | |
| | Consulting experience | | |
| 2023–Present | Marea Theraputics | | |
| | Beryl Cummings, PhD | | |
| | • Created tools to analyze large-scale genotype and phenotype data to p | rioritize targets | |
| | - Developed pipelines to produce disease specific scherts for genetic dise | 2010/0/ | |

 $_{\odot}$ $\,$ Developed pipelines to produce disease-specific cohorts for genetic discovery

Honors and fellowships

- 2021–2025 NIH Ruth L. Kirschstein National Research Service Award (NRSA F30), National Institute on Aging
 - 2021 Paul and Daisy Soros Fellowship for New Americans Finalist
 - 2021 DoD NDSEG Fellowship Program Waitlist
 - 2019 Outstanding Student Poster Presentation, New England Society of General Internal Medicine
 - 2016 Rose Award, UPenn Center for Undergraduate Research and Fellowships
 - 2016 Neysa Cristol Adams Prize in Biology Honorable Mention, UPenn Department of Biology
 - 2016 Hugo Otto Wolf Memorial Prize, UPenn School of Engineering and Applied Science
 - 2016 A. Norman Hixson Prize in Chemical Engineering, UPenn Department of Chemical and Biomolecular Engineering, CBE
 - 2016 **Senior Delaware Valley Section Award**, Delaware Valley Section of the American Institute of Chemical Engineers, AIChE
 - 2016 **President's Innovation Prize Semifinalist**, *UPenn*, Diagnosis of malaria in low-resource settings with AmpliTest
 - 2015 Donald F. & Mildred Topp Othmer Scholarship Award, AIChE National
 - 2015 Junior Delaware Valley Section Award, Delaware Valley Section of AIChE
 - 2014 Donald F. Othmer Sophomore Academic Excellence Award, UPenn CBE
- 2014–2016 Tau Beta Pi, Pennsylvania Delta Chapter
- 2012–2014, Dean's List, UPenn School of Engineering and Applied Science

2015-2016

Other publications and preprints

(* indicates co-first author)

- 2024 Rath, SP, **Gupta, R**, Todres, E, Wang, H, Jourdain, AA, et al. Mitochondrial genome copy number variation across tissues in mice and humans. *PNAS* 2024.
- 2024 Rogers, RS, Sharma, R, Shah, HS, Skinner, OS, Guo, A, Panda, A, **Gupta, R**, et al. Circulating N-lactoyl-amino acids and N-formyl-methionine reflect mitochondrial dysfunction and predict mortality in septic shock. *Metabolomics* 2024.
- 2023 Wharrie, S, Yang, Z, Raj, V, Monti, R, **Gupta, R**, et al. HAPNEST: efficient, large-scale generation and evaluation of synthetic datasets for genotypes and phenotypes. *Bioinformatics* 2023
- 2023 Majara, L, Kalungi, A, Koen, N, Tsuo, K, Wang, Y, **Gupta, R**, et al. Low and differential polygenic score generalizability among African populations due largely to genetic diversity. *HGG Adv.* 2023.
- 2022 Tsuo K, Zhou, W, Wang, Y, Kanai, M, Namba, S, **Gupta, R**, et al. Multi-ancestry meta-analysis of asthma identifies novel associations and highlights the value of increased power and diversity. *Cell Genomics* 2022.
- 2021 The COVID-19 Host Genetics Initiative. Mapping the human genetic architecture of COVID-19. *Nature* 2021.
- 2018 Jha, P, McDevitt, MT, Halilbasic, E, Williams, EG, Quiros, PM, Gariani, K, Sleiman, MB, Gupta, R, et al. Genetic Regulation of Plasma Lipid Species and Their Association with Metabolic Phenotypes. *Cell Systems* 2018.

Talks

- 2025 **Gupta, R**, (2025). Uncovering mechanisms driving mtDNA variation across 750,000 people. Freedom Together Foundation Annual Meeting. Oral presentation, plenary session.
- 2024 **Gupta, R**, (2024). Biobank-scale analysis of mtDNA copy number and heteroplasmy. Mass. General Hospital Center for Genomic Medicine Seminar Series. Oral presentation.
- 2024 **Gupta, R**, (2024). Nuclear genetic control of mtDNA homeostasis revealed from >250,000 diverse human genomes. *All of Us* Office Hours Researcher Presentation, Invited oral presentation.
- 2023 **Gupta, R**, (2023). Nuclear genetic control of mtDNA homeostasis revealed from >250,000 human genomes. Keystone Meeting; Mitochondrial Dysfunction: From Ultra-rare Diseases To Aging, Oral presentation, plenary session.

- 2023 **Gupta, R**, (2023). Nuclear genetic control of mtDNA homeostasis revealed from >250,000 human genomes. EUROMIT. Invited oral presentation.
- 2023 **Gupta, R**, (2023). Nuclear genetic control of mtDNA homeostasis revealed from >250,000 human genomes. Broad Institute Board of Scientific Counselors Meeting. Oral presentation.
- 2022 **Gupta, R**, (2022). The genetic determinants of mitochondrial function and aging in humans. Howard Hughes Medical Institute Deconstructing and Decoding the Genome Conference, Virtual.
- 2022 **Gupta, R**, (2022). Assessing mitochondrial dysfunction at biobank scale. Harvard Program in Genetics and Genomics Seminar. Oral presentation.
- 2021 **Gupta, R**, (2021). Quantifying mitochondrial dysfunction at biobank scale. Mass. General Hospital Center for Genomic Medicine Seminar Series. Oral presentation.

Posters

- 2024 **Gupta, R**, Chau, G, Durham, T, et al. (2024). Pervasive human mitochondrial-nuclear interactions and their implications for disease identified across >600,000 individuals. American Society of Human Genetics.
- 2022 **Gupta, R**, Sharma, R, Surendran, P, et al. (2022). Quantifying mitochondrial dysfunction at biobank scale using insights from rare disease. American Society of Human Genetics.
- 2022 **Gupta, R**, Karczewski, KJ, Howrigan, D, et al. (2022). Organelles and aging: a human genetic approach. Harvard Program in Genetics and Genomics Symposium.
- 2021 **Gupta, R**, Karczewski, KJ, Howrigan, D, et al. (2021). Organelles and aging: a human genetic approach. American Society of Human Genetics. Virtual.
- 2021 **Gupta, R**, Karczewski, KJ, Howrigan, D, et al. (2021). Human genetic analyses of organelles highlight the nucleus, but not the mitochondrion, in age-related trait heritability. MGH Scientific Advisory Board Meeting. Virtual.
- 2019 **Gupta, R**, McGirr, S, Trainor, A (2019). An acute presentation of hereditary pancreatitis in a 24-year-old. New England Regional Society of General Internal Medicine.
- 2019 **Gupta, R**, Calvo S, Aguet F, Mootha VK (2019). Human Genetics of Mitochondrial DNA Copy Number Variation. Harvard Medical School Soma Weiss Student Research Day; Boston, MA.
- 2017 **Gupta, R**, McDevitt M, Jha P, et al. (2017). Identification and Bioinformatic Network Analysis of Lipid Quantitative Trait Loci. Morgridge Scientific Advisory Board Meeting; Madison, WI.
- 2016 **Gupta, R**, Lan M, et al. (2016). The Proline/Arginine Dipeptide from C9orf72 RAN Translation Leads to Cellular Degradation Pathway Dysfunction. Penn Symposium on Undergraduate Research in Biology; Philadelphia, PA.

Previous research and laboratory experience

2018–2019 Broad Institute of MIT and Harvard, MD Research Assistantship, Cambridge MA

- Advisor: Vamsi K. Mootha, MD
 - Topic: Characterizing human variation in mtDNA copy number (mtCN) across tissues and individuals
 - $_{\odot}$ Measured mtCN across \approx 10,000 human samples using multiplex quantitative PCR
 - o Systematically assessed genetic and transcriptomic correlates to mtCN across tissues
- 2016–2017 Morgridge Institute for Research, Research assistant, Madison WI
 - Advisor: David Pagliarini, PhD
 - Topic: Leveraging systems genetics strategies to understand the link between liver and serum lipid species and clinical and molecular traits
 - o Created a pipeline to nominate genes from quantitative trait loci for lipidomics in mice
 - o Integrated genetic, transcriptomic, proteomic, and metabolomic data to highlight genes of interest

2014–2016 Children's Hospital of Philadelphia, Undergraduate researcher, Philadelphia PA

Advisor: Robert Kalb, MD

- Topic: Proteasome-mediated toxicity of dipeptide products from C9ORF72 hexanucleotide repeat expansion (HRE) in amyotrophic lateral sclerosis
 - o Used molecular techniques to probe the impact of peptide products from C9ORF72 HRE

- o Assessed proteasomal and autophagic flux impairment in C9ORF72 HRE
- 2015–2016 University of Pennsylvania Departments of Chemical Engineering & Bioengineering, Interdisciplinary senior design project, Philadelphia PA
 - Advisor: Haim Bau, PhD
 - Topic: Diagnosis of malaria in low-resource settings with AmpliTest
 - o Created an electricity-free diagnostic platform for P. falciparum using isothermal amplification
 - $_{\odot}$ Generated a low-cost prototype performed validation using 3D-printing and computer-aided design
 - 2013 Cytec Industries (now SOLVAY) Analytical Laboratories, Internship, Stamford, CT

Manager: Min Wang, PhD

o Developed methods for HPLC, GC, and MS-based analysis of synthetic products

Teaching and leadership

| 2020–Present | Live-in Resident Tutor, New Vassar House, MIT, Cambridge, MA | |
|------------------------|--|--|
| | $_{\odot}~$ Helped guide and support a new community of ~ 40 undergraduate students | |
| | • Provided academic advising, addressed concerns among students, facilitated a safe living community | |
| 2018–Present | Statistics and Medical School Admissions Tutor, Cambridge Coaching, Cambridge, MA | |
| 2021, 2022 | Journal Club Facilitator, The YES for CURE Program, Cambridge, MA | |
| | \circ Ran a weekly cancer-focused journal club for high schoolers emphasizing paper reading skills | |
| 2017–2021 | Class Representative, Curriculum Committee, HST MD Program, Harvard Medical School, Boston, MA | |
| 2018-2020 | Non-resident Pre-medical Tutor, Kirkland House, Harvard University, Cambridge, MA | |
| 2018–2019 | Co-founder, MD HST Diversity Ambassadors , <i>HST MD Program, Harvard Medical School</i> , Boston, MA | |
| | o Identified steps in the HST application pipeline which exacerbated exclusion of minority groups | |
| | $_{\odot}$ Designed targeted materials to boost engagement with underrepresented schools and groups | |
| 2014–2015 | Teaching assistant, Introduction to Engineering, UPenn SEAS, Philadelphia, PA | |
| 2015 | Teaching assistant, Mass Balances, UPenn SEAS, Philadelphia, PA | |
| 2014–2016 | Founder and co-president of Access Engineering, UPenn SEAS, Philadelphia, PA | |
| | $_{\odot}$ Created the organization's vision to improve engineering outreach to underprivileged high schools through weekly hands-on labs | |
| | o Generated a 2-semester curriculum including Java, Arduino robotics, and computer-aided design | |
| | $_{\odot}$ Formed relationships with 5 high schools, recruiting $>$ 70 students for weekly lessons | |
| | \circ Won School of Engineering and Applied Sciences club of the year in 2016 | |
| 2014–2016 | New Initiatives Chair and Section Leader of Penn Sciences Across Ages , University of Pennsylvania, Philadelphia, PA | |
| | Hosted a weekly science class at Lea Day School | |
| | $_{\odot}$ Worked on expanding outreach efforts by contacting new schools and developing new programs | |
| 2014-2016 | Community Service Chair, UPenn Tau Beta Pi Engineering Honor Society, Philadelphia, PA | |
| | Skills | |
| Graduate coursework | Probability I (BST 230), Statistical Inference I (BST 231), Statistical Methods (BST 232), Bayesian Methodology (BST 249), Principles of Genetics (GEN 201) | |
| Programming | R, Python, MATLAB, Java, OCaml, M/Caché | |
| - | Biobank-scale genome-wide association studies, multi-ancestry heritability analyses, phenome-wide associ- ation analyses, cross-trait genotype and phenotype correlations | |
| Platforms | Google Cloud Platform, DNANexus, Amazon AWS, Terra | |

Tools ImageJ, SolidWorks, ImageStudio, Adobe Illustrator, Docker