

Abhilesh Dhawanjewar

BIOINFORMATICIAN · MRC MITOCHONDRIAL BIOLOGY UNIT · UNIVERSITY OF CAMBRIDGE

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Education

Ph.D. in Evolutionary Biology

Aug 2015 - Jul 2022

UNIVERSITY OF NEBRASKA-LINCOLN, USA

Advisors: Dr. Kristi Montooth & Dr. Colin Meiklejohn

Integrated B.S.-M.S. Dual Degree

Aug 2010 - May 2015

INDIAN INSTITUTE OF SCIENCE EDUCATION AND RESEARCH, PUNE, INDIA

Advisor: Dr. M.S. Madhusudhan

Summary

I am a computational biologist with a strong interest in the molecular mechanisms underlying human disease, particularly disorders of the nervous system. My current work at the MRC Mitochondrial Biology Unit focuses on understanding the pathogenesis of neurodegenerative diseases driven by mitochondrial DNA (mtDNA) mutations. I leverage expertise in bioinformatics, single-cell genomics, and statistical modeling to build computational frameworks that dissect the fundamental processes governing the inheritance and effects of mtDNA mutations in neural tissues.

Building on a PhD in mitochondrial-nuclear coevolution, my research bridges evolutionary biology with clinically relevant questions. I use large-scale, multimodal datasets to investigate how mitochondrial dysfunction affects gene regulation, metabolism, and cell-type specific processes in the nervous system. Through this work, I aim to uncover mechanisms contributing to mitochondrial and neurodegenerative diseases, and to support the development of computational tools for their study.

Publications

JOURNAL ARTICLES

Burr S., Auckland K., Glynos A.* **Dhawanjewar A.S.***, Wei W., Ryall C., Hynes-Allen A.M., Prater M., Sczaniecka-Clift M., Prudent J., Chinnery P.F., & van den Amele J. (2025). MitoPerturb-Seq identifies common and gene-specific single-cell responses to mitochondrial DNA depletion and heteroplasmy *bioRxiv*, 2025.07.08.663208v1

Chandrasegaram R., Hynes-Allen A.M., Gao B., **Dhawanjewar A.S.**, Frison M., Petridi S., Chinnery P.F., Ma H. & van den Amele J.(2025). Single-molecule mitochondrial DNA imaging reveals heteroplasmy dynamics shaped by developmental bottlenecks and selection in different organs in vivo *bioRxiv*, 2025.01.24.634671v1

Dhawanjewar, A.S., Montooth K.L., & Meiklejohn, C.D. Mitochondrial OXPHOS genes exhibit higher levels of molecular compensation of human disease associated mutations relative to nuclear OXPHOS genes in mammals. *In preparation, manuscript available on request*

M. Florencia Camus & **Dhawanjewar, A.S.** (2023). Multilevel selection on mitochondrial genomes. *Current Opinion in Genetics & Development*, 80, 102050.

Dhawanjewar A.S.*, Roy A.A.* , & Madhusudhan M.S. (2020). A knowledge-based scoring function to assess the stability of quaternary protein assemblies. *Oxford Bioinformatics*, 36(12), 3739-3748.

Roy, A.A.* , **Dhawanjewar, A.S.***, Sharma, P., Singh, G., & Madhusudhan, M.S. (2019). Protein Interaction Z Score Assessment (PIZSA): an empirical scoring scheme for evaluation of protein-protein interactions. *Nucleic acids research*, 47(W1), W331-W337.

Montooth, K.L., **Dhawanjewar, A.S.**, & Meiklejohn, C.D. (2019). Temperature-sensitive reproduction and the physiological and evolutionary potential for Mother's Curse. *Integrative and comparative biology*, 59(4), 890-899.

Nelson, T.C., Jones, M.R., Velotta, J.P., **Dhawanjewar, A.S.**, & Schweizer, R.M. (2019). UNVEILING connections between genotype, phenotype, and fitness in natural populations. *Molecular ecology*, 28(8), 1866-1876.

* Equal contribution

WEB-SERVERS

Prediction of Stable Quaternary Protein Assemblies -

PIZSA (Protein Interaction Z-score Assessment) - <http://cospi.iiserpune.ac.in/pizsa/>

Research Experience

Cell- and Tissue-specificity of Mitochondrial Disease Mutations

MRC MITOCHONDRIAL BIOLOGY UNIT · UNIVERSITY OF CAMBRIDGE

2024-Present

UK

- Developed and applied computational frameworks to analyze multimodal single-cell data, aiming to understand cellular heterogeneity and tissue-specific effects driving mitochondrial neurodegenerative disease pathogenesis.
- Investigated nuclear genome contributions to mitochondrial disease by analyzing CRISPR-based screen data to identify key genetic modifiers.
- Built and implemented robust bioinformatic pipelines for diverse genomic datasets, including single-cell transcriptomics and protein-DNA interaction analysis (DamID-seq).

The Evolution of Sexually Antagonistic Variation in Fruit Flies

UNIVERSITY COLLEGE LONDON

2022-Present

UK

- Designed experiments implementing sex-limited selection in *Drosophila melanogaster* for experimental evolution
- Developed analytical and statistical tools using Approximate Bayesian Computation (ABC) to identify and characterize sexually antagonistic variation from genomic data

Mitochondrial-Nuclear Coevolution in Mammalian Genomes

UNIVERSITY OF NEBRASKA-LINCOLN

2021-2022

USA

- Compiled datasets and computed evolutionary rate correlations between mitochondrial genes and nuclear genes with different degrees of interaction for mammalian species.
- Nuclear genes interacting with mitochondrial genes exhibit stronger correlations in evolutionary rates, supporting the hypothesis of mito-nuclear coevolution.

Molecular Compensation in the Oxidative Phosphorylation System (OXPHOS)

UNIVERSITY OF NEBRASKA-LINCOLN

2019-2021

USA

- Curated and analyzed mitochondrial and nuclear protein sequences from 1200 mammalian species to identify potential compensating residues for disease-causing mutations using sequence, structural and phylogenetic analysis
- Mitochondrial genes exhibit a higher degree of compensatory evolution compared to nuclear genes, suggesting a higher degree of functional redundancy in the mitochondrial genome

Mitochondrial-Nuclear Interactions and the Thermal Sensitivity of Male Reproduction

UNIVERSITY OF NEBRASKA-LINCOLN

2016-2019

USA

- Performed G×G×E×E×E phenotypic assays to characterize the effects of a mitochondrial-nuclear incompatibility between *Drosophila melanogaster* and *Drosophila simulans* hybrid on thermal male sterility.
- Mitochondrial-nuclear incompatibility exacerbates thermal sensitivity of spermatogenesis that is further modulated by environmental cues such as temperature, diet and age of exposure.

Prediction of Stable Quaternary Assemblies Protein Interaction Z Score Assessment (PIZSA)

INDIAN INSTITUTE OF SCIENCE EDUCATION AND RESEARCH, PUNE

2013-2015

India

- Constructed knowledge-based statistical potentials trained over 4900 native three-dimensional protein structures to predict the stability of protein-protein interactions
- Extensively bench-marked across multiple test sets and is among the top 6 methods, outperforming 31 other statistical, physics, based and machine learning scoring schemes and deployed the algorithm as a web-server

Honors & Awards

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| Registration Waiver and Travel Support , EMBO Population genomics: Background and tools (€400) | 2024 |
| Blair Paxton Udale Fund for Life Sciences, The University of Nebraska Foundation (\$1900) | 2021 |
| Milton E. Mohr Fellowship, UNL Center for Biotechnology (\$1000) | 2021 |
| Blair Paxton Udale Fund for Life Sciences, The University of Nebraska Foundation (\$500) | 2020 |
| Milton E. Mohr Fellowship, UNL Center for Biotechnology (\$1000) | 2019 |
| Suzanne O. Prather Memorial Fund, University of Nebraska Foundation (\$1500) | 2019 |
| Runner-Up Best Poster Award, School of Biological Sciences, UNL (\$50) | 2019 |
| AAAS/Science Program for Excellence in Science, American Association for the Advancement of Science | 2019 |
| Jessie A. Lee Fund, School of Biological Sciences, UNL (\$2000) | 2018 |
| Best Poster Award, School of Biological Sciences, UNL (\$100) | 2018 |
| Conference Registration Award, Society for Molecular Biology and Evolution (\$450) | 2018 |
| Travel Grant, Society for Molecular Biology and Evolution (\$250) | 2017 |
| Runner-Up Best Poster Award, School of Biological Sciences, UNL (\$50) | 2017 |
| Mary D. Rogick Memorial Fund, School of Biological Sciences, UNL (\$1300) | 2017 |
| Travel Grant, Society for the Study of Evolution (\$500) | 2016 |
| Blair Paxton Udale Fund for Life Sciences, The University of Nebraska Foundation (\$1500) | 2016 |
| Rosemary Grant Award, Society for the Study of Evolution (\$2500) | 2016 |
| Travel Grant, The Indian Institute of Science Education and Research, Pune (\$1300) | 2014 |
| Travel Grant, The American Society of Naturalists (\$250) | 2014 |
| Working Internship, Max Planck Institute for Evolutionary Biology (\$3500) | 2013 |
| INSPIRE Scholarship, Department of Science and Technology, India (\$8000) | 2010 |
| National Talent Search Examination (NTSE) Scholar, NCERT, India (\$250) | 2006 |

Conference Presentations

INVITED TALKS

The Ethics of Using Genetic Tools for Conservation

UNVEIL SYMPOSIUM 2018

Jun 2018

Missoula, Montana, USA

Population Genomics of the Range-Expanding Populations of *Argiope bruennichi*

20TH INTERNATIONAL CONGRESS OF ARACHNOLOGY

Jul 2016

Golden, Colorado, USA

ORAL PRESENTATIONS

Faster Mitochondrial Evolution Drives Mitochondrial-Nuclear Coevolution

CELLS WITHIN CELLS SYMPOSIUM

Jan 2025

Cambridge, UK

Compensatory Evolution of Disease Associated Residues in the Oxidative Phosphorylation (OXPHOS) pathway *

SOCIETY FOR MOLECULAR BIOLOGY AND EVOLUTION MEETING

Jun 2020

Québec City, Canada

Environmental Modification of Mitochondrial-Nuclear Epistasis in Shaping Thermal Male Sterility in *Drosophila*

UNVEIL SYMPOSIUM 2018

Jun 2018

Missoula, Montana, USA

Mitochondrial-Nuclear Interactions and the Thermal Sensitivity of Male Reproduction

MITOCHONDRIAL GENOMICS AND EVOLUTION, AN SMBE SATELLITE MEETING

Sep 2017

Ein Gedi, Israel

POSTER PRESENTATIONS

The Evolution of Sexual Antagonism in Fruit Flies

EMBO POPGEN - POPULATION GENOMICS: BACKGROUND AND TOOLS

Jul 2024

Naples, Italy

Compensatory Evolution of Disease Associated Residues in the Mitochondrial Genome

2ND UNVEIL SYMPOSIUM 2019

Oct 2019

Lincoln, Nebraska, USA

Structural Compensation of Disease Associated Residues in the Mitochondrial Genome

EUROPEAN SOCIETY FOR EVOLUTIONARY BIOLOGY MEETING

Aug 2019

Turku, Finland

Mitochondrial Diseases and Compensated Pathogenic Deviations

SOCIETY FOR INTEGRATIVE AND COMPARATIVE BIOLOGY MEETING

Jan 2019

Tampa, Florida, USA

Genetic and Environmental Factors Underlying the Thermal Sensitivity of Male Reproduction

SOCIETY FOR MOLECULAR BIOLOGY AND EVOLUTION MEETING

Jul 2018

Yokohama, Japan

Mitochondrial-Nuclear Interactions and the Thermal Sensitivity of Male Reproduction

UNIVERSITY OF NEBRASKA-LINCOLN SPRING RESEARCH FAIR

Apr 2017

Lincoln, Nebraska, USA

Mitochondrial-Nuclear Interactions and the Thermal Sensitivity of Male Reproduction

58TH ANNUAL DROSOPHILA RESEARCH CONFERENCE

Mar 2017

San Diego, California, USA

Prediction of Protein-Protein Interactions through the use of Statistical Potentials

BIOPHYSICS PASCHIM MEETING

Mar 2015

Mumbai, India

Comparative Mitogenomic Analysis in the Range-Expanding Populations of *Argiope bruennichi*

QEVOLUTION2014, WORKSHOP ON QUANTITATIVE EVOLUTIONARY BIOLOGY

Sep 2014

Şirince, Turkey

* Conference cancelled due to COVID-19 concerns

Outreach

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| Panel Member at Imperial Lates: Future Cities, Imperial College London | 2025 |
| Technology Lead and Ecology Surveyor, The Ealing Beaver Project | 2022-Present |
| BioBlitz Organiser and Surveyor, Ascott Allotments, Ealing, London | 2023 |
| Organiser & Lightning Talks and Film Festival Master of Ceremonies, SciComm 2020 | 2020 |
| My Captain Discover Mentor, The Climber | 2018 |
| Scientists in Cars Getting Coffee, Film Festival, SciComm 2018 | 2018 |
| Master of Ceremonies, Lightning Talks and Film Festival, SciComm 2018 | 2018 |
| Boys and Girls Science Club, Park Middle School, Lincoln Community Learning Centers | 2016-2017 |
| Junior Sunday with a Scientist, Nebraska State Museum | 2017 |
| Sunday with a Scientist: Diversity of Life in Nebraska, Nebraska State Museum | 2017 |
| Sunday with a Scientist: Darwin Day, Nebraska State Museum | 2017 |
| Science Night Live Moderator, SciComm 2016 | 2016 |
| Sunday with a Scientist: Evolution on the Wing, Nebraska State Museum | 2016 |
| Junior Sunday with a Scientist, Nebraska State Museum | 2016 |
| Investigate: Show-and-tell Amblypygi, Nebraska State Museum | 2016 |
| Science Tutoring for less-privileged high school students, Pune, India | 2013-2015 |

Teaching

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| Guest Lecture, BIOL0011 - Evolutionary Genetics | Spring 2023 |
| Teaching Assistant, LIFE 120L - Fundamental Biology Lab I | Fall 2021 - Spring 2022 |
| Guest Lecture, BIOS 897 - Communicating Science Through Outreach | Spring 2017, 2018 |
| Teaching Assistant, LIFE 121L - Fundamental Biology Lab II | Fall 2020 |
| Teaching Assistant, LIFE 120L - Fundamental Biology Lab I | Spring 2019 |
| Teaching Assistant, LIFE 120L - Fundamental Biology Lab I | Fall 2016 - Fall 2017 |
| Teaching Assistant, BIOS 101L - General Biology Lab | Fall 2015 - Spring 2016 |

Professional Service

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| Postdoctoral Representative, Genetics, Evolution and Environment, UCL | 2022-Present |
| Organizing Committee, SciComm 2020: A Conference on Effective Science Communication | 2020 |
| Graduate Student Representative, UNL oSTEM Conference 2020 | 2020 |
| Workshop co-organizer: Ethics of Biotechnology Applications to Conservation Biology, UNVEIL Symposium 2018 | 2018 |
| Grad Student Volunteer, Strategic Vision Committee, School of Biological Sciences, UNL | 2018 |
| Vice President, Biology Graduate Students Association, UNL | 2017-2019 |
| Undergraduate Poster Judge, UNL Spring Research Fair | 2018-2021 |

Peer Review

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| Ecology and Evolution | Genetics |
| Journal of Evolutionary Biology | G3: Genes Genomes Genetics |

Memberships

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| Society for Molecular Biology and Evolution (SMBE) | 2018-2025 |
| European Society for Evolutionary Biology (ESEB) | 2019-2020 |
| The Society for Integrative and Comparative Biology (SICB) | 2019-2020 |
| American Association for the Advancement of Science (AAAS) | 2019-2020 |
| Genetics Society of America (GSA) | 2017-2019 |
| Society for the Study of Evolution (SSE) | 2016-2019 |
| International Society of Arachnologists (ISA) | 2016-2017 |
| American Society of Naturalists (ASN) | 2014-2015 |