Parul Johri

School of Life Sciences

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EDUCATION	
2018 - Present	Postdoctoral Researcher , Arizona State University, Tempe, AZ Advisor: Jeffrey D. Jensen
2012 – 2018	PhD, Evolution, Ecology and Behavior Program Major: Evolution; Minor: Bioinformatics Indiana University, Bloomington, IN Advisor: Michael Lynch
2009 – 2012	Master's in Biology (By Research) Tata Institute of Fundamental Research, Mumbai, India
2006 - 2009	B.Sc. (Honours) Mathematics St. Stephen's College, Delhi University, Delhi, India

RESEARCH INTERESTS

Population genetics, Statistical inference, Molecular evolution.

PUBLICATIONS

PEER-REVIEWED

- <u>Parul Johri*</u>, Brian Charlesworth*, Emma K. Howell, Michael Lynch[§], Jeffrey D. Jensen[§]. Revisiting the notion of deleterious sweeps. *Genetics*, 2021. (Highlighted by *Genetics*; *In press*)
- **Parul Johri**§, Kellen Riall, Hannes Becher, Laurent Excoffier, Brian Charlesworth, Jeffrey D. Jensen§. The impact of purifying and background selection on the inference of population history: problems and prospects. *Molecular Biology and Evolution*, 2021. 38(7): 2986-3003.
- **Parul Johri**§, Brian Charlesworth, Jeffrey D. Jensen§. Towards an evolutionarily appropriate null model: jointly inferring demography and purifying selection. *Genetics*, 2020. 215: 173-192. (<u>Highlighted by *Genetics*</u>)
- <u>Parul Johri</u>*,§, Georgi K. Marinov*,§, Thomas G. Doak, Michael Lynch. Population genetics of *Paramecium* mitochondrial genomes: recombination, mutation spectrum, and efficacy of selection. *Genome Biology and Evolution*, 2019. 11(5): 1398–1416.
- Parul Johri[§], Sascha Krenek, Georgi K. Marinov, Thomas, G. Doak, Thomas U. Berendonk, Michael Lynch. Population genomics of *Paramecium* species. *Molecular Biology and Evolution*, 2017. 34(5): 1194-1216.

- Matthew S. Ackerman, <u>Parul Johri</u>, Ken Spitze, Sen Xu, Thomas G. Doak, Kimberly Young, Michael Lynch. Estimating seven coefficients of pairwise relatedness using population-genomic data. *Genetics*, 2017. 206:105-118.
- Casey L. McGrath, Jean-Francois Gout, <u>Parul Johri</u>, Thomas G. Doak, Michael Lynch. Differential retention and divergent resolution of duplicate genes following whole-genome duplication. *Genome Research*, 2014. 24(10): 1665-75.

IN REVIEW/ PREPRINTS

- <u>Parul Johri</u>, Wolfgang Stephan, Jeffrey D. Jensen[§]. Soft selective sweeps: addressing new definitions, evaluating competing models, and interpreting empirical outliers. (*major revision in Plos Genetics*)
- Parul Johri, Charles F. Aquadro, Mark Beaumont, Brian Charlesworth, Laurent Excoffier, Adam Eyre-Walker, Peter D. Keightley, Michael Lynch, Gil McVean, Bret A. Payseur, Susanne P. Pfeifer, Wolfgang Stephan, Jeffrey D. Jensen. Statistical inference in population genomics. bioRxiv 2021.10.27.466171; doi: https://doi.org/10.1101/2021.10.27.466171, 2021. (Invited Perspective, Genetics)
- Ana Yansi Morales-Arce*, **Parul Johri***, Jeffrey D. Jensen§. Inferring the distribution of fitness effects in influenza A virus and human cytomegalovirus. (*Accepted, pending revision in Heredity*)
- Jean-Francois Gout, <u>Parul Johri</u>, Olivier Arnaiz, Thomas G. Doak, Simran Bhullar, Arnaud Couloux, Fréderic Guérin, Sophie Malinsky, Linda Sperling, Karine Labadie, Eric Meyer, Sandra Duharcourt, Michael Lynch. Universal trends of post-duplication evolution revealed by the genomes of 13 *Paramecium* species sharing an ancestral whole-genome duplication. *bioRxiv* 573576; doi: https://doi.org/10.1101/573576, 2019.

IN PREPARATION

- <u>Parul Johri</u>, Ryan N. Gutenkunst, Kirk E. Lohmueller, Adam Eyre-Walker, Jeffrey D. Jensen. On the prospect of achieving accurate joint estimation of selective effects together with population history. (*Invited Perspective, Genome Biology and Evolution*)
- **Parul Johri**§, Jean-Francois Gout, Michael Lynch. A population-genetic lens into the process of gene duplicate loss after whole-genome duplication. (*In preparation*)

^{*}These authors contributed equally.

[§] Corresponding authors.

PRESENTATIONS

INVITED

- 2021 EvoLunch seminar, Institute of Science and Technology, Vienna, Austria
- 2021 Department of Biology, Carleton University, Ottawa, Ontario, Canada
- **2021** EVOLTREE conference: Genomics and Adaptation in Forest Ecosystems (Keynote speaker), Birmensdorf, Switzerland
- 2021 Jawaharlal Nehru Centre for Advanced Scientific Research, Bangalore, India
- **2021** International Laboratory for Human Genome Research, National Autonomous University of Mexico, Mexico
- 2020 Department of Biology, University of North Carolina, Chapel Hill, NC
- 2020 Center for Evolution and Medicine, Arizona State University, Tempe, AZ

CONTRIBUTED (SELECTED)

- **2021** *Population Genetics Group*, Liverpool, England. Joint inference of demography and purifying selection.
- **2020-** Arizona Population Genetics Conference, Tempe, Arizona. Effects of fixation and segregation of deleterious mutations.
- **2019-** *Arizona Population Genetics Conference*, Tempe, Arizona. Joint estimation of demography and purifying selection.
- **2019 -** Annual meeting of the Society for the Study of Evolution (SSE), Providence, Rhode Island. Joint estimation of demography and purifying selection.
- **2019 -** Annual Meeting of the Society for Molecular Biology and Evolution (SMBE), Manchester, UK. Mechanistic basis of loss and preservation of whole-genome duplicates.
- **2019**, **2017** Evolution Research Symposium, Arizona State University, Tempe, Arizona.
- **2018-** *Arizona Population Genetics Conference*, Tucson, Arizona. *Paramecium* population genomics: constraints on non-coding regions and whole-genome duplicates.
- **2017** Annual meeting of the Society for the Study of Evolution (SSE), Portland, Oregon. Evolution and population-genetics of mitochondrial genomes in Paramecium species.
- **2016-** The Allied Genetics Conference (TAGC), Orlando, Florida. Population genomics of Paramecium species.
- **2016-** Annual Meeting of the Society for Molecular Biology and Evolution (SMBE), Queensland, Australia. Population genomics of Paramecium species.
- **2015-** Midwest Protozoology Meeting, Peoria, Illinois. Genetic variation in Paramecium.

ACADEMIC AWARDS/SCHOLARSHIPS:

2018-2020	Early Career Reviewer at Genetics, Genetics Society of America
2018, 2016	Young Investigator Travel Award, Society for Molecular Biology and Evolution.
Spring 2017	College of Arts and Sciences Travel Award, Indiana University.
Spring 2014	Departmental Fellowship, Indiana University.
2009-2012	Annual Departmental Fellowship, Tata Institute of Fundamental Research, India.
2008-2009	Medal for the most <i>Outstanding Student</i> in Mathematics, Department of Mathematics, St. Stephen's College, Delhi University, India.
Summer 2009	Summer Research Fellowship, Indian Academy of Sciences, Bangalore, India. [Awarded annually nationwide to 100 students (undergraduate and graduate) in Biology.]
Summer 2008	Summer Research Fellowship, Jawaharlal Nehru Centre for Advanced Scientific Research (JNCASR), Bangalore, India. [Awarded annually nationwide to 30 students (undergraduate and graduate) in Biology.]

PROFESSIONAL SERVICE

ORGANIZATION

2021- Co-organizer (with Jeffrey D. Jensen) of the symposium entitled "The effects of selection at linked sites and population history on levels and patterns of genomic variation" in the annual meeting of the *Society of Molecular Biology and Evolution*.

REVIEWER FOR

Genetics; G3: Genes, Genomes, Genetics; Genome Biology and Evolution; Bioinformatics; Molecular Ecology; Evolution; Ecology and Evolution; Journal of Molecular Evolution; BMC Genomics.

MEMBERSHIP IN SCIENTIFIC SOCIETIES

Society for Molecular Biology and Evolution (SMBE), 2012 – Present

Genetics Society of America (GSA), 2014 – Present

Society for the Study of Evolution (SSE), 2016 – Present

TEACHING AND MENTORING

UNDERGRADUATES SUPERVISED

Kellen Riall, August 2019 – June 2021 [Current position: PhD student at the University of Chicago]

Emma Howell, Spring 2019 – Summer 2020 [Current position: PhD student at the University of Wisconsin-Madison]

CLASSROOM TEACHING

Spring 2015	Head teaching assistant, Evolution (L318), Department of Biology, Indiana
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University.

Spring 2013 Head teaching assistant, **Biology Laboratory** (L113), Department of Biology,

Indiana University.

Fall 2012 Associate teaching assistant, **Biology Laboratory** (L113), Department of Biology,

Indiana University.

OTHER RESEARCH EXPERIENCES

Junior Research Scholar	Deflagellation in <i>Chlamydomonas reinhardtii</i> - the underlying signalling
2010-2012	mechanisms. Advisor: Prof. B. J. Rao, Tata Institute of Fundamental Research,

Mumbai, India.

Junior Research Scholar

2009 - 2010

Predicting multiple origins of replication in bird mitochondrial genomes using Monte Carlo Markov models. Advisor: Prof. B. J. Rao, Tata Institute of Fundamental Research, Mumbai; Co-advisor: Dr. Neeraja Krishnan, Indian

Institute of Science, Bangalore, India.

Summer Research Fellow

May-July, 2008

Mathematical modelling of the neuronal networks in the saccadic eye system. Advisor: Dr. Aditya Murthy, National Brain Research Centre, Gurgaon, India.

Summer Research Fellow

May-July, 2007

Culture of human endothelial cells in microfluidic channels. Advisor: Dr. Kaustubh Rao, National Centre for Biological Sciences, Bangalore, India.