# ARJUN BIDDANDA

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#### RESEARCH INTERESTS

Population Genomics, Statistical Genetics, Medical Genetics, Ancient DNA

## **EMPLOYMENT**

2021 - Postdoctoral Research Associate · Department of Statistics · University of Oxford

Advisor: Pier Palamara

#### **EDUCATION**

2015 - 2020 PhD · Human Genetics · University of Chicago

Dissertation Title: Investigating the spatio-temporal structure of human genetic diversity

Advisor: John Novembre

2011 - 2015 BS.Eng · Computer Science (Cum Laude) · Cornell University

Advisor: Alon Keinan

## **PUBLICATIONS**

\* - indicates equal contribution

Arjun Biddanda, Daniel P. Rice, and John Novembre. Geographic patterns of human allele frequency variation: a variant-centric perspective. eLife, 2020.

Charleston W K Chiang, Joseph H Marcus, Carlo Sidore, Arjun Biddanda, Hussein Al-Asadi, Magdalena Zoledziewska, Maristella Pitzalis, Fabio Busonero, Andrea Maschio, Giorgio Pistis, Maristella Steri, Andrea Angius, Kirk E Lohmueller, Goncalo R Abecasis, David Schlessinger, Francesco Cucca, and John Novembre. Genomic history of the Sardinian population. Nature Genetics, 2018.

Peter de Barros Damgaard, Rui Martiniano, Jack Kamm, J Víctor Moreno-Mayar, ..., Arjun Biddanda, ..., Martin Sikora, Alan K Outram, Richard Durbin, and Eske Willerslev. The first horse herders and the impact of early Bronze Age steppe expansions into Asia. Science, 2018.

Yedael Y Waldman\*, Arjun Biddanda\*, Natalie R Davidson, Paul Billing-Ross, Maya Dubrovsky, Christopher L Campbell, Carole Oddoux, Eitan Friedman, Gil Atzmon, Eran Halperin, and Others. The genetics of Bene Israel from India reveals both substantial Jewish and Indian ancestry. PLoS One, 11(3):e0152056, 2016.

Yedael Y Waldman, Arjun Biddanda, Maya Dubrovsky, Christopher L Campbell, Carole Oddoux, Eitan Friedman, Gil Atzmon, Eran Halperin, Harry Ostrer, and Alon Keinan. The genetic history of Cochin Jews from India. Human Genetics, pages 1–17, 2016.

Feng Gao\*, Diana Chang\*, Arjun Biddanda\*, Li Ma, Yingjie Guo, Zilu Zhou, and Alon Keinan. XWAS: a software toolset for genetic data analysis and association studies of the X chromosome. Journal of Heredity, 106(5):666-671, 2015.

# PRESENTATIONS

Arjun Biddanda, Matthias Steinrücken, and John Novembre. Properties of two-locus genealogies and linkage disequilibrium in temporally stratified samples. Probabilistic Models in Genomics (Poster), 2021.

Arjun Biddanda, Matthias Steinrücken, and John Novembre. Linkage Disequilibrium in Ancient DNA: Theory and Applications. American Society of Human Genetics (Poster), 2019.

Arjun Biddanda, Matthias Steinrücken, and John Novembre. Linkage Disequilibrium in Ancient DNA: Theory and Applications. Midwest Population Genetics (Talk), 2019.

Arjun Biddanda and John Novembre. Inference and visualization of the geographic distribution for variant sets. American Society of Human Genetics (Poster), 2018.

HONORS / AWARDS

2020 Presidential Membership

Genetics Society of America

2019 Reviewer's Choice Abstract

American Society of Human Genetics

2017 Honorable Mention

NSF Graduate Research Fellowship

2015-2018 NIH Genetics and Regulation Training Grant

University of Chicago

2013 - 2015 Dean's List

Cornell University College of Engineering

TEACHING EXPERIENCE

Winter 2019 Guest Lecturer, University of Chicago

Computing Skills for Biologists

Winter 2018 Teaching Assistant, University of Chicago

HG 486: Fundamentals of Computational Biology

Summer 2017 Course Assistant, University of Chicago

Marine Biological Laboratory - Quantitative Approaches to Biology Bootcamp

Spring 2017 Teaching Assistant, University of Chicago

HG 469: Human Variation and Disease

2016 - 2018 Tutor, University of Chicago

Introduction to Statistics for Geneticists

2016 - present Instructor, Software Carpentry

2013 - 2015 Teaching Assistant, Cornell University

CS 3110: Functional Programming and Data Structures

INDUSTRY EXPERIENCE

2013 Software Engineering Intern

Guidewire Software

SERVICE

Summer 2020 Organizer

Genetics and Society Reading Group (Departmental Group)

2019-2020 Novembre - He - Stephens (NHS) Meeting Coordinator

University of Chicago

2019 - 2020 Tutor

Tutoring Chicago

2019 - Peer Review

eLife, Molecular Biology and Evolution, Genes

**MENTORSHIP** 

2020 - present Achyutha Menon (U. Chicago Undergraduate)

Project: Efficient storage and visualization of linkage disequilibrium across human populations

Sponsored by a College Summer Research Fellowship

COMPUTATIONAL

**SKILLS** 

Python, Bash, R, OCaml, C++, Java

Git, LTEX, Microsoft Office

SOCIETY

**Memberships** 

American Society of Human Genetics (ASHG) Genetics Society of America (GSA) 2018 -

2021 -