Enakshi Saha, PhD

Department of Biostatistics Harvard University E-mail: esaha@hsph.harvard.edu Homepage: http://www.enakshisaha.com Github: https://github.com/Enakshi-Saha

Employment

• University of South Carolina Assistant Professor at Arnold School of Public Health – Department of Epidemiology and Biostatistics	Columbia, SC August, 2024 - present
• Harvard University • Research Associate at Harvard T.H. Chan School of Public Health – Mentor: Dr. John Quackenbush	Boston, MA June, 2023 - present
Postdoctoral Fellow at Harvard T.H. Chan School of Public Health – Mentor: Dr. John Quackenbush	June, 2021 - June, 2023
• Argonne National Laboratory • National Science Foundation Mathematical Sciences Graduate Intern	Lemont, IL June - August, 2020
 Mentor: Dr. Carlo Graziani, Argonne National Laboratory & The Department of Astronomy & Astrophysics 	University of Chicago,

Education

• The University of Chicago	Chicago, USA
• PhD in Statistics	September, 2016 - June, 2021
 Thesis Title: Flexible Bayesian Methods for High-dimensional Data Advisor: Dr. Veronika Rockova, The University of Chicago Booth School of Business 	
• Indian Statistical Institute	Kolkata, India
• Master of Statistics(M.Stat)	2014 - 2016
 Specialization: Mathematical Statistics and Probability Dissertation: Some One-sample Tests for High Dimensional Low Sample Size Data (Supervised by Dr. Anil K. Ghosh, Stat-Math Unit, ISI Kolkata) 	
• Indian Statistical Institute	Kolkata, India
• Bachelor of Statistics(B.Stat) (Hons.)	2011 - 2014

Publications

2016 SARIMA modeling of the monthly average maximum and minimum temperatures in the eastern plateau region of India. Saha, Enakshi, Arnab Hazra, and Pabitra Banik. Mausam 67, no. 4: 841-848.

2017 Some high-dimensional one-sample tests based on functions of interpoint distances.

Saha, Enakshi, Soham Sarkar, and Anil K. Ghosh. Journal of Multivariate Analysis 161: 83-95.

2019 On theory for BART.
 Rockova, Veronika, and Enakshi Saha.
 22nd international conference on artificial intelligence and statistics (AISTATS), pp. 2839-2848. PMLR.

2021 **Dynamic sparse factor analysis.** Saha, Enakshi, Kenichiro McAlinn and Veronika Rockova Journal of Applied Econometrics (Revision invited) arXiv:1812.04187.

2021 Impact of the COVID-19 induced lockdown measures on PM2. 5 concentration in USA.

Ghosal, Rahul, and Enakshi Saha. Atmospheric Environment 254 (2021): 118388.

2023 The Network Zoo: a multilingual package for the inference and analysis of gene regulatory networks.

Ben Guebila M, Wang T, ... , Saha E, et al. Genome Biology 24, no. 1: 45.

2023 Gender difference in the effects of chronic diseases on daily physical activity patterns in older adults: analysis of objectively measured physical activity in NHATS 2021

Saha, Enakshi and Rahul Ghosal Annals of Epidemiology 86, 110-118.e4.

2023 Theory of Posterior Concentration for Generalized Bayesian Additive Regression Trees Saha, Enakshi

Electronic Journal of Statistics (revision submitted) arxiv:2304.12505.

2023 Gene regulatory Networks Reveal Sex Difference in Lung Adenocarcinoma Saha E, Marouen Ben Guebila, ..., Dawn DeMeo, John Quackenbush, Camila Lopes-Ramos. Journal of the National Cancer Institute (submitted) bioRxiv 2023.09.22.559001.

2023 BONOBO: Bayesian Optimized sample-specific Networks Obtained By OMICS data

Saha E, Viola Fanfani, ..., John Quackenbush. Genome Research (revision submitted), RECOMB 2024 (accepted) bioRxiv 2023.11.16.567119.

2024 Understanding the Effect of Aging on Lung Adenocarcinoma using Gene Regulatory Networks

Saha E, Marouen Ben Guebila, Viola Fanfani, Kate H. Shutta, Dawn L. DeMeo, John Quackenbush, Camila M. Lopes-Ramos *European Respiratory Journal (submitted)*.

2024 node2vec2rank: Large Scale and Stable Graph Differential Analysis via Multi-Layer Node Embeddings and Ranking Mandros P., Ian Gallagher, ..., Saha E, et al. *PNAS Computational Biology (submitted) bioRxiv 2024.06.16.599201.*

Working Papers

- 2024 SEAHORSE: Serendipity Engine Assaying Heterogeneous Omics-Related Sampling Experiments Saha E, Derrick DeConti, ..., John Quackenbush. http://seahorse.tm4.org/
- 2024 Single-sample Multiomic Associations Using Gaussian graphical models Saha E, Kate Hoff-Shutta, ... , John Quackenbush.

Teaching Experience

As Instructor:

Statistical Methods and Applications, Spring 2019, The University of Chicago

As Teaching Assistant (at The University of Chicago):

- Measure Theoretic Probability III, Spring 2021
- Applied Linear Statistical Methods, Fall 2020 (Remote)
- Statistical Methods and Applications, Winter 2020, Fall 2019, Fall 2018
- Mathematical Statistics II, Spring 2018
- Mathematical Statistics I, Winter 2018
- Statistical Theory and Method II, Winter 2017 and Spring 2017

Services

As Journal Reviewer:

- Statistics: Bayesian Analysis, Sankhya Series A
- Machine Learning: ICML, AISTATS
- Medicine: Annals of the American Thoracic Society, Journal of Applied Pharmaceutical Science

Invited Talks

• 28th Annual International Conference on Research in Computational Molecular Biology (RECOMB 2024)

Title: BONOBO: Bayesian Optimized sample-specific Networks Obtained By OMICS data

• Global Health Week 2024 at the Harvard T. H. Chan School of Public Health

Title: Gender difference in the effects of chronic diseases on daily physical activity patterns in older adults: analysis of objectively measured physical activity in NHATS 2021

• Pulmonary and Critical Care Medicine, Work-in-Progress, 2023-2024, Brigham and Women's Hospital

Title: Gene Regulatory Networks Reveal Sex Difference in Lung Adenocarcinoma

• 2023 Joint Statistical Meetings, Toronto, Canada

Title: Bayesian Optimized sample-specific Networks Obtained By assimilating OMICS data (BONOBO)

• Harvard University Network Science Meeting, 2020

Title: Bayesian Latent Variable Models for Dynamic Data

• University of Chicago Department of Statistics Consulting Seminar 2020

Title: Bayesian Latent Variable Modeling of COVID-19 Infection Rate

• 2020 Talk at Radix Trading, Chicago, Illinois

Title: Dynamic Sparse Factor Analysis and Its Role in Dynamic Portfolio Allocation

• LANS Summer Argonne Students' Symposium, 2020

Title: Bayesian Latent Variable Modeling of COVID-19 Infection Rate

• ORISE / DOE Ignite Off! 2020

Title: Bayesian Latent Variable Modeling of COVID-19 Infection Rate

- 2019 Joint Statistical Meetings, Denver, Colorado Title: On Theory for BART
- University of Chicago Department of Statistics Consulting Seminar 2019

Title: Role of Gene Tbx5 in Formation of Forelimb in Zebrafish

• University of Chicago Department of Statistics Consulting Seminar 2018

Title: Effect of Hunger on Moral Development in Children

- University of Chicago Department of Statistics Consulting Seminar 2017 Title: An Introduction to Bradley-Terry Model
- D.Basu Memorial Lecture (2014), ISI Kolkata

Title: Exact Distribution Free Multisample Run Tests Applicable to High Dimensional Data

Awards and Honors

- Recipient of Radix Trading Fellowship 2020-2021
- Finalist of ORISE / DOE Ignite Off Competition (2020)

Talk Title: Bayesian Latent-Variable Modeling of COVID-19 Infection Rate

- Winner of Senior Consultant Award, Department of Statistics, The University of Chicago (2018-19)
- Winner of Statistical Consulting Project, Department of Statistics, The University of Chicago (2017-18)
- Nominated for D.Basu Gold Medal for the most outstanding student in B.Stat, Indian Statistical Institute (2014)
- Recipient of Mukul Chaudhury Memorial Prize for the Highest Scoring Female Student in B.Stat, Indian Statistical Institute (2012-2013)
- Recipient of Prize Money for Academic Excellence in B.Stat and M.Stat, Indian Statistical Institute (2011-2015)

Statistical Consulting Experience

- Effect of Hunger on Moral Development in Children with Elizabeth Huppert & Jean Decety, Department of Psychology, The University of Chicago
- Role of Gene Tbx5 in Formation of Forelimb in Zebrafish

with Erin Boyle Anderson & Robert K. Ho, Department of Organismal Biology and Anatomy, The University of Chicago

• Moderated Mediation Analysis of Moral Communication and Moral Hypocrisy

with Elizabeth Huppert, Emma Levine & Jean Decety, Department of Psychology, The University of Chicago

Programming Skills

R, Python, MATLAB

References

• Dr. John Quackenbush

Henry Pickering Walcott Professor of Computational Biology and Bioinformatics; Chair of the Department of Biostatistics at the Harvard T.H. Chan School of Public Health; Professor in the Channing Division of Network Medicine, and Professor at the Dana-Farber Cancer Institute

Contact: johnq@hsph.harvard.edu

• Dr. Carlo Graziani

Computational Scientist at Argonne National Laboratory and Research Associate Professor at the Department of Astronomy & Astrophysics, University of Chicago

 $Contact: \ cgraziani@anl.gov$

• Dr. Rajarshi Mukherjee

Assistant Professor in the Department of Biostatistics at the Harvard T.H. Chan School of Public Health

Contact: ram521@mail.harvard.edu

• Dr. Dawn L. DeMeo

Associate Professor of Medicine at Harvard Medical School, and a senior respiratory genetics researcher in the Channing Division of Network Medicine.

Contact: redld@channing.harvard.edu

• Dr. Camila M. Lopes-Ramos

Instructor in Medicine, Department of Biostatistics at the Harvard T.H. Chan School of Public Health, Brigham and Women's Hospital and Harvard Medical School

Contact: camilaramos@hsph.harvard.edu