## UTHSAV CHITRA

Eric and Wendy Schmidt Center, Broad Institute of MIT and Harvard

https://uthsavc.github.io

# EDUCATION/ACADEMIC TRAINING

### Broad Institute of MIT and Harvard

July 2024 - present

Postdoctoral fellow, Eric and Wendy Schmidt Center

**Princeton University**, Princeton, New Jersey

Sept 2018 - May 2024

Ph.D., Computer Science

Brown University, Providence, Rhode Island

Sept 2013 - May 2017

Sc.B. Mathematics, A.B. Computer Science, A.B. Applied Math

GPA: 4.0/4.0

### RESEARCH INTERESTS

Computational genomics, machine learning, spatial biology, graphs and networks, genetic interactions.

#### **Publications**

Anomaly Detection in Spatial Transcriptomics via Spatially Localized Density Comparison.

Gary Hu, Julian Gold, Uthsav Chitra, Sunay Joshi, Benjamin J. Raphael.

In submission to ISMB 2025.

GASTON-Mix: a unified model of spatial gradients and domains using spatial mixture-of-experts.

Uthsav Chitra, Shu Dan, Fenna Krienen, Benjamin J. Raphael.

In submission to ISMB 2025.

Spatial metabolic gradients in the liver and small intestine.

Laith Samarah, Clover Zheng, Xi Xing, Won Dong Lee, Amichay Afriat, **Uthsav Chitra**, Michael MacArthur, Wenyun Lu, Connor Jankowski, Cong Ma, Craig Hunter, Benjamin J. Raphael, Joshua Rabinowitz. Under review at *Nature*.

Mapping the topography of spatial gene expression with interpretable deep learning.

**Uthsav Chitra**, Brian J. Arnold, Hirak Sarkar, Cong Ma, Sereno Lopez-Darwin, Kohei Sanno, Benjamin J. Raphael.

Nature Methods, in press. Accepted to RECOMB 2024.

Quantifying higher-order epistasis: beware the chimera.

Uthsav Chitra\*, Brian J. Arnold\*, Benjamin J. Raphael.

Nature Communications, in press.

A latent variable model for evaluating mutual exclusivity between driver mutations in cancer.

Ahmed Shuaibi\*, Uthsav Chitra\*, Benjamin J. Raphael.

RECOMB Satellite Workshop on Computational Cancer Biology (RECOMB-CCB), 2024. Best Paper Award.

A count-based model for delineating cell-cell interactions in spatial transcriptomics data.

Hirak Sarkar\*, Uthsav Chitra\*, Julian Gold, Benjamin J. Raphael.

Bioinformatics (2024). Accepted to ISMB 2024.

Belayer: Modeling discrete and continuous spatial variation in gene expression from spatially resolved transcriptomics.

Cong Ma\*, Uthsav Chitra\*, Shirley Zhang, Benjamin J. Raphael.

Cell Systems (2022). Accepted to RECOMB 2022.

NetMix2: Unifying network propagation and altered subnetworks.

Uthsav Chitra\*, Tae Yoon Park\*, Benjamin J. Raphael.

<sup>\*</sup> denotes joint first authorship

Journal of Computational Biology (2022). Accepted to RECOMB 2022.

Quantifying and Reducing Bias in Maximum Likelihood Estimation of Structured Anomalies.

Uthsav Chitra, Kimberly Ding, Jasper C. H. Lee, Benjamin J. Raphael.

International Conference on Machine Learning (ICML) 2021.

NetMix: A network-structured mixture model for reduced-bias estimation of altered subnetworks.

Matthew A Reyna\*, Uthsav Chitra\*, Rebecca Elyanow, Benjamin J. Raphael.

Journal of Computational Biology (2021). Accepted to RECOMB 2020.

Analyzing the Impact of Filter Bubbles on Social Network Polarization.

Uthsav Chitra and Christopher Musco.

ACM International Web Search and Data Mining Conference (WSDM) 2020.

Also appeared at KDD WISDOM 2019 workshop.

Random Walks on Hypergraphs with Edge-Dependent Vertex Weights.

Uthsav Chitra and Benjamin J. Raphael.

International Conference on Machine Learning (ICML) 2019.

Committee Selection is More Similar Than You Think: Evidence from Avalanche and Stellar.

Tarun Chitra and Uthsav Chitra.

Manuscript, 2019.

### Honors and Awards

Rising Stars in Data Science, UChicago/UC San Diego/Stanford Data Science Institutes	2024	
Best Paper Award, RECOMB Satellite Workshop on Computational Cancer Biology	2024	
Siebel Scholarship	2022	
• Award of \$35,000 given annually to 85 top graduate students worldwide in computer science,		
bioengineering, and business.		
Best Reviewer Award, International Conference on Machine Learning (ICML)	2021, 2022	
NSF Graduate Research Fellowship	2020	
Jerome Stein Memorial Award, Brown University Applied Math Department	2017	
• Given to the top two students who "show outstanding potential in an interdisciplinary area that involves		
applied mathematics."		
Phi Beta Kappa, Brown University (elected junior year, top 2% of class)	2016	
Top 200, William Lowell Putnam Math Competition	2015	
First Place, Brown University Hartshorn-Hypatia Math Examination	2013	
Semi-finalist, Siemens Competition (research project in number theory)	2012	

### Teaching

# Instructor/Curriculum Developer, Princeton Prison Teaching Initiative

2019-2023

- Taught college-accredited math classes at NJ state prisons.
- Developed and taught first-ever Java programming course for NJ state prisons.

### Teaching Assistant/Grader, Brown University

- MATH 1560: Number Theory
- CSCI 1570: Design and Analysis of Algorithms
- CSCI 1450: Probability in Computing
- CSCI 0530: Linear Algebra for CS
- MATH 1530: Abstract Algebra

Spring 2016, Spring 2017

Fall 2015, Fall 2016

Spring 2015

Fall 2014

Spring 2014

Counselor, Program in Mathematics for Young Scientists (PROMYS)

Summer 2014

• Guided students through daily number theory problem sets, mentored a group project, and aided seminars in abstract algebra.

• Assisted online, real-time math classes in algebra, number theory, combinatorics, and geometry.

# Talks

Mapping the topography of spatial gene expression with interpretable dee	p learning.
Computational and Systems Biology (CSB) seminar, MIT	November 2024
Rising Stars in Data Science, UC San Diego	November 2024
Eric and Wendy Schmidt Center meeting, Broad Institute	October 2024
Chen lab meeting, Broad Institute	July 2024
Conference on Research in Computational Molecular Biology (RECOMB)	May 2024
Campbell Lab Meeting, UToronto	$April\ 2024$
Single Cell Analyses, Cold Spring Harbor Laboratory (poster)	$November\ 2023$
Rutgers-Princeton Cancer Research Symposium (poster)	October 2023
NCI Junior Investigator (JI) Annual Meeting	$August\ 2023$
Wang Lab Meeting, Broad Institute	July 2023
Belayer: Modeling discrete and continuous spatial variation in gene expresersolved transcriptomics	ssion from spatially
Computational and Systems Biology (CSB) seminar, MIT	November 2024
Rising Stars in Data Science, UC San Diego	November 2024
Chen lab meeting, Broad Institute	July 2024
Campbell Lab Meeting, UToronto	April 2024
Knowles/Azizi Lab Meeting, Columbia	September 2023
Pe'er Lab Meeting, MSKCC	August 2023
Wang Lab Meeting, Broad Institute	July 2023
Sankararaman/Pimentel Lab Meeting, UCLA	April~2023
Pe'er Lab Group Meeting, Columbia	$April\ 2023$
Rutgers-Princeton Cancer Research Symposium (poster)	$October\ 2023$
NCI Junior Investigator (JI) Annual Meeting	August~2023
Brigham Women's Hospital Advanced Biomedical Computation Series	March 2023
NCI Spring School on Algorithmic Cancer Biology	March 2023
NetMix2: Unifying network propagation and altered subnetworks	
Conference on Research in Computational Molecular Biology (RECOMB)	May 2022
NetMix: A network-structured mixture model for reduced-bias estimation	of altered subnetworks
Knowles/Azizi Lab Meeting, Columbia	$September\ 2023$
Pe'er Lab Meeting, MSKCC	$August\ 2023$
Conference on Research in Computational Molecular Biology (RECOMB)	June~2020
Princeton University Generals Exam	May 2020
Analyzing the Impact of Filter Bubbles on Social Network Polarization	
ACM International Web Search and Data Mining Conference (WSDM)	February 2020
KDD WISDOM Workshop	August 2019
Random Walks on Hypergraphs with Edge-Dependent Vertex Weights	
SIAM Conference on Discrete Mathematics	June 2022
Princeton University Generals Exam	May 2020
International Conference of Machine Learning (ICML)	June 2019
Students Mentored	

Jairam Hathwar, Princeton CS undergraduate

Kohei Sanno, Princeton CS undergraduate

Clover Zheng, Princeton CS PhD student

Sunay Joshi, Princeton Math undergraduate

Ahmed Shuaibi, Princeton QCB PhD student

• Won Best Paper Award at RECOMB-CCB workshop.

Kimberly Ding, Princeton CS undergrad
• Fall 2019: Recommender Systems with Hypergraph Random Walks

- Spring 2020: Maximum Likelihood Estimation of Structured Anomalies
- Senior Thesis 2020-2021: Spatial-NetMix: Less Biased and More Flexible Anomaly Detection
  - Received the "Outstanding Computer Science Senior Thesis Prize"

Shirley Zhang, Princeton CS undergrad/alumni

Madelyne Xiao, Princeton CS PhD student

Summer 2020, 2021-2022

2022

2019-2021

• Received an NSF Graduate Research Fellowship

## SERVICE/OUTREACH

### Conference Reviewing

Computational biology: RECOMB 2020 poster session, RECOMB 2023, ISMB 2023, RECOMB 2024, ISMB 2024

Machine learning: ICML 2021 (**Top 10% Reviewer**), NeurIPS 2021, ICML 2022 (**Top 10% Reviewer**), ICML 2023, TMLR, ICML 2024 AccMLBio workshop.

# Program Committee

ISMB 2025.

## Journal Reviewing

Bioinformatics, Bioinformatics Advances, Frontiers in Big Data, Computational and Structural Biotechnology Journal.

Member, Princeton COS Graduate Student Committee	2022-2023
Member, Princeton Graduate Engineering Council	2021-2023
Member, Princeton COS Ad Hoc Committee	2021
Officer, Brown Math Departmental Undergraduate Group	2015-2017
Mentor, Brown Matched Advising Program for Sophomores	2016-2017

### Work Experience

#### Software Engineer, Facebook

2017-2018

• Built infrastructure, machine learning models, and data pipelines for improving ad quality.

## Software Engineering Intern, Facebook

Summer 2016

• Reduced upload time for video ads by 20%.

Hobbies/interests: Bouldering, biking, crosswords and other puzzles.