

Prateek Anand

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My research interests are in developing novel AI/ML computational methods. Broadly, I am invested in deep learning, traditional machine learning, and statistical approaches that are scalable and interpretable. Current applications of my work are centered around genetics and published in RECOMB 2024 and Genome Research.

EDUCATION

UCLA | Ph.D. Computer Science

2024 - 2029

Advisor: Dr. Sriram Sankararaman

I am currently developing improved generative models for synthetic data and imputation methods in genetics.

UCLA | B.S. Computer Science

2020 - 2024

PROFESSIONAL EXPERIENCE

UCLA Computer Science | Sriram Lab

Jun 2022 - Present

Graduate Student Researcher

- Developing scalable machine learning/statistical software to understand human genetic architecture (relationship between genetics and traits)
- Projects: QuadKAST (co-first author) and FAME

Research Intern (Bruins in Genomics)

- Evaluated novel nonlinear explanation method for machine learning models on simulated genetic data (form of symbolic regression used to represent complex black box models)
- Presented at UCLA poster symposium

Stanford University School of Medicine | Curtis Lab

Jun 2023 – Jan 2024

Research Intern (Canary CREST Program for Early Cancer Detection)

- Early detection of blood cancer through computational modeling and inference, refined model built using approximate bayesian computation
- Presented at Stanford poster symposium

UCLA Jonsson Comprehensive Cancer Center | Boutros Lab

Jan 2021 - Apr 2022

Research Assistant

- Updated, maintained, and evaluated CI/CD HATCHet pipeline for use on lab cluster compute server (tool for quantifying/locating copy-number aberrations in human cancer samples)
- Project: Metapipeline-DNA

PUBLICATIONS

QuadKAST | Genome Research

Sep 2024

- Fu, B.*, Anand, P.*, Anand, A.*, Mefford, J., & Sankararaman, S. (2024). *A Scalable Adaptive Quadratic Kernel Method for Interpretable Epistasis Analysis in Complex Traits*.
<https://doi.org/10.1101/2024.03.09.584250>

Metapipeline-DNA | Preprint

- Patel, Y.*, Zhu, C.*, Yamaguchi, T. N.* et al. Anand, P., ... Boutros, P. C. (2024). *Metapipeline-DNA: A Comprehensive Germline & Somatic Genomics Nextflow Pipeline*.
<https://doi.org/10.1101/2024.09.04.611267>

- Fu, B.*, Pazokitoroudi, A.*, Xue, A., Anand, A., Anand, P., Zaitlen, N., & Sankararaman, S. (2023). *A Biobank-Scale Test of Marginal Epistasis Reveals Genome-Wide Signals of Polygenic Epistasis*. <https://doi.org/10.1101/2023.09.10.557084>

TEACHING

Teaching Assistant	Jan 2025
<ul style="list-style-type: none">• UCLA CM146: Introduction to Machine Learning	
Course Reader	Jan 2024
<ul style="list-style-type: none">• UCLA CM146: Introduction to Machine Learning	
Course Reader	Jan 2023
<ul style="list-style-type: none">• UCLA CM146: Introduction to Machine Learning	

SKILLS

Programming languages: Python, C++

Frameworks: Scikit-learn, PyTorch, Numpy, Pandas, Git

Courses: Machine Learning, Artificial Intelligence, Neural Networks and Deep Learning, Big Data Analytics, Software Engineering and Construction, Data Structures, Algorithms and Complexity, Linear Algebra, Statistics and Probability, Optimization, Machine Learning in Genetics, Algorithms in Bioinformatics

Other: Probabilistic Models, Hypothesis Testing, Cluster Computing

AWARDS

Warren Alpert Computational Biology and AI Fellow: Graduate Training Fellowship	2024
NCI Scholarship: Funding for Canary CREST Research Program	2023
NSF REU Scholarship: Funding for Bruins in Genomics Research Program	2022
UCLA Dean's Honors List: Winter 21, Spring 21, Fall 21, Winter 22, Fall 22, Winter 23	
Andy Grove Intel Scholarship: Education Scholarship for Academic Excellence	2020
Valedictorian: Homestead High School	2020