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.

EDUCATION

UCLA | Ph.D. Computer Science

2024 - 2029

Advisor: Dr. Sriram Sankararaman

I am currently developing improved deep generative models for synthetic data and imputation methods in genetics. I have also recently been exploring discrete diffusion and masked language modeling approaches for rare feature imputation.

UCLA | B.S. Computer Science

2020 - 2024

PROFESSIONAL EXPERIENCE

UCLA Computer Science | Sriram Lab

Jun 2022 - Present

Graduate Student Researcher

- Building deep generative models for genetic variation data (presenting at American Society of Human Genetics in Boston, Oct 2025)
- Developed kernelized, scalable machine learning/statistical software to understand human genetic architecture
- Projects: QuadKAST (co-first author), FAME, FLEX

Research Intern (Bruins in Genomics)

- Evaluated novel nonlinear explanation method for machine learning models (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)

- Computational modeling and inference, refined model built using approximate bayesian computation to detect blood cancer early
- 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
- Project: Metapipeline-DNA

PUBLICATIONS

FLEX | Preprint Oct 2025

 Liu, Z., Fu, B., Jeong, M., <u>Anand, P.,</u> Anand, A., Jang, S., Gorla, A., Zhu, J., Pajukanta, P., Palamara, P., Zaitlen, N., Border, R., & Sankararaman, S. (2025). Comprehensive gene heritability estimation reveals the genetic architecture of rare coding variants underlying complex traits. https://doi.org/10.1101/2025.10.07.681018

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

FAME | To appear in Nature Genetics

Sep 2023

• 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

Jan 2025 **Teaching Assistant**

UCLA CM146: Introduction to Machine Learning

Jan 2024 **Course Reader**

UCLA CM146: Introduction to Machine Learning

Jan 2023 Course Reader

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: Generative Models, Probabilistic Models, Hypothesis Testing, Cluster Computing

AWARDS

Warren Alpert Computational Biology and Al 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