

Gino Prasad

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Machine Learning | Computer Vision | Bioinformatics | Data Science

Education

PhD in Computer Science, UC San Diego

NSF Graduate Research Fellow (Fall 2025 – Present)

Summer 2023 – Spring 2027

Academic Advisor: [Vineet Bafna](#), Professor of Computer Science

GPA: 3.96 / 4.00

- **Focus:** Applied Machine Learning and Computer Vision

- **Relevant Courses:** Deep Learning, Generative AI, 3D Physics Simulation, LLM Modeling, Recommender Systems

Bachelor of Science, Bioinformatics: UC San Diego

Fall 2020 - Spring 2023

Major: Bioinformatics (B.S.), Minor: Computer Science

GPA 3.97 / 4.00

- **Relevant Courses:** Machine Learning, Optimization, Probability & Statistics, Physics (Optics/Fluids), Linear Algebra, Algorithms

Experience

Data Science & Machine Learning Intern

Jun 2025 - Aug 2025

Genentech Research & Development (gRED Computational Sciences)

- Implemented an **LLM agent system** with **LangChain** for natural language querying and analysis of large-scale cell-line screening datasets.
- Developed a **AI Computer Vision Model (VAE)** of **5M+** Cell Imaging Data samples of cancer cell lines to predict cell lifecycle stage.
- Trained Transformer Models to predict **drug dose-response** from gene expression and cell line embeddings.
- **Technical Skills:** Transformers, PyTorch, Scikit-Image, LLM Agents, LangChain, Deep Learning

Machine Learning Graduate Researcher

Jun 2022 - Current

UC San Diego Bafna Lab

- Developed interSeg, an **AI Computer Vision Model** for **classification** in biomedical imaging achieving high predictive accuracy.
- **InterSeg** predicts cancer subtypes in In-Vitro Cell Lines and Patient Tissue **Imaging Data**.
- Co-authored **EvidenceBench**, a benchmark for assessing **LLM reasoning** performance on biomedical papers.
- **Technical Skills:** PyTorch, TensorFlow, OpenCV, Skimage, Pandas

Machine Learning & Computational Biology Research Assistant

Oct 2021 - Jun 2023

UC San Diego Yeo Lab

- Built a **Convolutional Neural Network** for Spatial Transcriptomics Image data, to perform **semantic segmentation**.
- Used a **U-Net Architecture** to predict nuclei boundaries, bypassing the need for DAPI staining.
- **Technical Skills:** TensorFlow, Keras, NumPy, Pandas, PyTorch, PyLab, Python, Linux, Git.

Software Engineering Intern

Jun 2021 - Aug 2021

People Inc.

- Designed front-end software for People Inc., the largest digital publisher in the US, managing sites like Investopedia and Verywell Health.
- Developed cross-platform web applications in a collaborative environment using Agile/Scrum.
- **Technical Skills:** JavaScript, Vue, HTML, SASS, Maven, Database querying, APIs.

Phage Genomics Research Initiative

Oct 2020 - Jun 2021

UC San Diego Poglano Lab

- Created a **BLAST parser website** using Google App Engine and Python ([GitHub](#)), used by the UCSD professor and class.
- Queries the NCBI BLAST data to perform comparative genomic analysis of Bacteriophage genes with unknown functions.
- **Technical Skills:** Flask, Python, Google Cloud App Engine.

Publications

Rajkumar*, Prasad*, et al., Accurate Prediction of ecDNA in Interphase Cancer Cells using Deep Neural Networks.

bioRxiv Preprint, <https://doi.org/10.1101/2025.06.23.661188> (Co-First Author Publication)

Wang et al. (2025), EvidenceBench: A Benchmark for Extracting Evidence from Biomedical Papers.

Conference On Language Modeling, <https://doi.org/10.48550/arXiv.2504.18736>

Luebeck et al. (2025), AmpliconSuite enables discovery of extrachromosomal DNA in tumor genomes.

American Association for Cancer Research, <https://doi.org/10.1158/1538-7445.AM2025-3755>

Dehkordi et al. (2025), Breakage fusion bridge cycles drive high oncogene number with moderate intratumoural heterogeneity.
Nature Communications, <https://doi.org/10.1038/s41467-025-56670-8>

Lv et al. (2025), Spatial-Temporal Diversity of Extrachromosomal DNA Shapes Urothelial Carcinoma Evolution and Tumor-Immune Microenvironment.
Cancer Discovery, <https://doi.org/10.1158/2159-8290.CD-24-1532>

Luebeck et al. (2024), AmpliconSuite: Analyzing focal amplifications in cancer genomes.
Cancer Genetics, <https://doi.org/10.1016/j.cancergen.2024.08.015>

Mah et al. (2024), Bento: A toolkit for subcellular analysis of spatial transcriptomics data.
Genome Biology, <https://doi.org/10.1186/s13059-024-03217-7>

Chapman et al. (2023), Circular extrachromosomal DNA promotes inter- and intratumoral heterogeneity in high-risk medulloblastoma.
Nature Genetics, <https://doi.org/10.1038/s41588-023-01551-3>

Prichard et al. (2023), Identifying the core genome of the nucleus-forming bacteriophage family and characterization of *Erwinia* phage RAY.
Cell Reports, <https://doi.org/10.1016/j.celrep.2023.112432>

Achievements

June 2025	NSF Graduate Research Fellowship , Awarded for Outstanding Achievement in ML/Bioinformatics Research	National Science Foundation (NSF)
Oct 2024	Cancer Grand Challenge eDynamic Conference , Presented a Computer Vision Model for Cancer Imaging	Cancer Grand Challenge (NCI)
Oct 2024	Cancer Grand Challenge Future Leaders Speaker , Presented on AI applications in Cancer Research	Cancer Grand Challenge (NCI)
June 2023	Summa Cum Laude Honors , Awarded for Exceptional GPA.	UC San Diego
April 2023	Undergraduate Research Conference , Presented on Computer Vision methods for FISH Imaging.	UC San Diego
Jul 2020	UCSD BioScholars Honors Society Member , Awarded membership based on academic achievement.	UC San Diego

Skills

Machine Learning	Experience With Computer Vision, Generative AI, Transformer Architectures, and Convolutional Neural Networks .
Computer Vision	Research Focus in developing AI Models for Cancer Imaging and Tumor Modeling.
Programming	Python (PyTorch, JAX, TensorFlow, Keras, Pandas, NumPy), R, C++, Cloud Computing, Bash, Linux, Git, JavaScript, Java, SQL.

Mentoring

2023 - 2024	Data Science Capstone Mentor , Mentored 3 Data Science undergraduates in a Computer Vision capstone competition to semantically segment cell imaging data using deep learning.	UC San Diego
2023 - 2024	Early Research Scholars Program Mentor , Mentored 2 Computer Science undergraduates in a project to extract mutational signatures from The Cancer Genome Atlas (TCGA) genome sequencing data.	UC San Diego

Personal Projects

Autotune Implementation Using Phase Vocoder

github.com/GinoP123/AutotunePV.git	May 2023
• Created an autotuner from scratch using Phase Vocoders and Yin pitch prediction. • Able to autotune any audio clip to a specific major or minor scale using Hann window functions. • Examples of popular songs autotuned here .	

Custom Search Engine for Linux File System

github.com/GinoP123/FileSearch	Jul 2022
• Created a keyword-matching search engine, with caching, completely from scratch using dynamic programming. • Added learning capability by including popularity and relevance weights. • I personally use this tool all the time, and find it a huge time-saver for navigating in Linux.	