ISHAAN GUPTA

Passionate about unraveling the complexities of the genome, its variation and evolution, particularly implications in the immune system, using Bioinformatics Algorithms and ML

EDUCATION

Ph.D, Computer Science (Research area: Bioinformatics), UC San Diego, La Jolla (2023 – present)
Charles Lee Powell Fellow, Performing Genomics + Immunology + AI research under Dr. Pavel Pevzner

BS, Computer Science: specialization in Bioinformatics, UC San Diego, La Jolla (2019 – 2022)

bs, computer science : specialization in bioinformatics, oc san biego, La Jolla (2)

GPA: 3.92, Magna Cum Laude with CS Honors, IEEE Eta Kappa Nu Honors Society

Experience

Research Assistant (Dr. Pavel Pevzner)

04/2021 - 08/2023

San Diego, CA
8105296351

i3gupta@ucsd.edu

i3gupta@ucsd.edu

github.com/IshaanSD

in linkedin.com/in/IgAI

UCSD CSE Department, San Diego, CA

- · Applying my **Undergraduate Honors Thesis** (2022) to find and analyze novel antibody genes in primates and human haplotypes
- · Optimized <u>UniAligner</u> (**C++** aligner for highly-repetitive regions) reduced runtime **by 80%** for 1M bp; added <u>interactive dotplots</u> feature
- · Developed Nanopore-based protein sequencing pipeline by applying Signal Processing and ML (Clustering, PCA) approaches

<u>Software Engineer I</u> 06/2022 – 12/2022

Illumina (Systems Integration), San Diego, CA

- · Improved traceability of LIMS software central to Illumina's high-throughput Genotyping and Methylation pipelines (Agile SDLC)
- · Developed and deployed highly scalable microservices in Java 7/8 (Spring+JDBC) integrated using APIs and AMQP
- · Found and fixed critical bugs in legacy code, troubleshot software integration issues, automated tests, curated database for support

Bioinformatics and Machine Learning Intern

07/2021 - 09/2021

Abterra Biosciences, San Diego, CA

- · Achieved >95% performance in Proteomics task for Antibody sequencing by experimenting multiple Deep learning approaches
- · Optimized data structures and used Data Manipulation, parallel processing and ML libraries in Java 8 (DL4J)

Machine Learning Project Consultant

11/2020 - 03/2021

Model Medicines, La Jolla, CA

- · Developed **BigQuery**-powered **data pipeline** supporting **SQL queries** and **neural networks** for **screening** COVID-19 drug candidates.
- · Made 3 web scrapers to read scientific journals and evaluate novelty of potential drugs using search keywords and their synonyms

Undergraduate Research Assistant

03/2020 - 03/2021

Mali Lab, Bioengineering UCSD, San Diego, CA

- · Performed research on Immune-orthogonal epitopes for safe and efficient CRISPR/Cas9 gene therapies
- · Built Variants Analysis pipeline for pooled library screens and web scraper using Bash and Python scripts and Command line tools

Projects

Autocorrelation for ecDNA hubs (Graduate-level course project)

Analyzed spatial organization of extrachromosomal DNA, and confirmed formation of hubs that help co-promoting oncogenic expression by visualizing *Autocorrelation* function for *image processing* of nuclei-stained cell images

• GANs for Cancer Image Augmentation

Implemented Generative Adversarial Network using PyTorch to mimic mammograms with tumor and generate augmented data

COVID Mutations Analysis (https://youtu.be/C27B4mYRpXg)

Used NCBI database, BLAST CLI, and JSON parser to find and analyze phylogeny of mutations that increase COVID-19 infectivity Led project on *phylogeny of Sars-CoV-2 variants* to understand the evolution of different strains at *geographic* level

Notes2Map (Python API + JS Web app)

Developed web app to process lecture notes and transcripts and generate Mind Map of keywords for better visualization Used *Flask* to make *REST APIs* handling *NLP* and *Graph Algorithms* in Python and communicate with *ReactJS UI* app.

Mentoring/Leadership

• Vice President, Undergraduate Bioinformatics Club

08/2021 - 07/2022

Managing various club events for helping Bioinformatics students socialize, prepare for industry/academic positions

• Data Science Student Society Workshop Chair and Bioinformatics Bootcamp Chair

08/2020 - 07/2021

Developed 10 dry lab workshops on Bioinformatics skills, ML, Shell, Python, R, SQL; hosted labs on an AWS EC2 instance

SKILLS

- Java (Stream, lambda, DL4J, JUnit), OOP, Spring, JDBC, AMQP
- Python (Pandas, PyTorch, TensorFlow, SciPy), C++, Shell, R, C
- SQL (MySQL, Postgres), Stored Proc, MongoDB, Cassandra (CQL)
- HTML, CSS, JS (React, Node.js, Express), Web components, puppeteer
- Agile (Scrum/Kanban) practices, Git, CI/CD pipeline, Docker
- Hadoop, Apache Spark (MLlib), Data Pipeline: Airflow, Kafka, RabbitMQ
- AWS: Redshift, S3, EC2, SQS, Elastic Beanstalk
- Machine learning: CNN, GANs, NLP, Recommender Systems
- Bioinformatics: scRNA analysis, plink GWAS, ChIP Seq, Bioconductor

COURSEWORK

- Algorithms for Computational Biology (Graduate-level)
- Deep Learning, Recommender Sys, Computer Vision
- Software Engineering, Parallel Computing (CUDA)
- Adv. Bioinformatics Lab, Biomolecular Big Data

CERTIFICATIONS

- Programming Foundations: Design Patterns
- Spring Data; Effective Intg. Testing with Spring Boot
- Spark and Hadoop; NoSQL Databases
- SQL for Data Science; CNNs;