Pavitra Ganesan Dass

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EDUCATION

Northeastern University, Boston, MA, US

Master of Science in Bioinformatics; GPA (3.66/4.00)

University of Mumbai, Mumbai, India May. 2023

Bachelor of Pharmacy; GPA (8.14/10)

TECHNICAL SKILLS

Programming & Pipelines: R (Seurat, DESeq2, ggplot2, dplyr), Python, Bash, Linux

 $\textbf{NGS \& Bioinformatics} : Bulk\ RNA-seq,\ scRNA-seq,\ STAR,\ Salmon,\ DESeq2,\ GSEA/MSigDB,\ Seurat,\ Bioconductor,\ SAM tools,\ Salmon,\ DESeq2,\ GSEA/MSigDB,\ Seurat,\ Bioconductor,\ SAM tools,\ Salmon,\ Sa$

GATK, BWA, IGV

Machine Learning & Statistics: Logistic Regression, Random Forest, kNN, PCA, UMAP, ensemble learning, ROC-AUC, F1

Visualization & QC: ggplot2, pheatmap, EnhancedVolcano, pathway dot plots, heatmaps, FastQC, MultiQC

Domain Interests: Cancer immunology (melanoma, myeloma), neuroinflammation (Alzheimer's microglia), viral rebound (HIV/SIV)

Communication: Lab presentations, teamwork across disciplines, proposal submissions

PROFESSIONAL EXPERIENCE

Center for Virology and Vaccine Research | Beth Israel Deaconess Medical Center, Harvard Medical School | Boston, MA, US
Bioinformatics Research Intern

July. 2025 - Present

- Analyzing baseline **PBMC RNA-seq and plasma cytokine profiles** from 20 SIV-infected rhesus macaques to identify predictive biomarkers of viral rebound following ART interruption
- Implementing **multi-omics pipelines** (DESeq2, GSEA/MSigDB, limma, Cytoscape/STRING) to uncover immune–metabolic pathways linked to rebound susceptibility
- Developing **predictive models** (logistic regression, random forest with cross-validation in scikit-learn) to integrate transcriptomic and cytokine features into a baseline rebound "signature"
- Conducting **single-sample and pathway-level analyses** (SLEA, correlation with viral load trajectories) to connect baseline immune programs with rebound dynamics.
- Designing visualizations and reproducible workflows (PCA, UMAP, heatmaps, pathway dot plots) to distill complex analyses into actionable biological insights.
- Translating findings into HIV cure research strategies; authored and presented a \$5K ResearchHub microgrant, externally peer-reviewed

Athena Drug Delivery Solutions Pvt Ltd | Ambernath, Maharashtra, India

Internship
Collaborated with teams to ensure quality control and microbiological testing compliance

Jun. 2022 - July. 2022

- Conadorated with teams to ensure quanty control and interoblological testing compliance
- Applied problem-solving to improve processes and maintained detailed records with attention to detail
- Communicated effectively to report findings and worked both independently and in teams to streamline laboratory operations

PROJECTS

Single-cell RNA-seq: Tumor Immune Profiling | Northeastern University

April. 2025

Dec. 2025

• Replicated scRNA-seq workflow (Seurat in R); performed QC, normalization, clustering, and pathway analysis to identify immune cell signatures in lung cancer samples.

Heart Failure Detection Using Machine Learning | Northeastern University

April. 2025

• Built Decision Tree, Logistic Regression, and Random Forest models on clinical data; applied preprocessing, feature engineering, and evaluation (ROC-AUC, F1) to improve predictive accuracy

Bulk RNA-Seq: Gene Expression & Pathway Analysis | Northeastern University

Feb. 2025

• Processed RNA-seq data (Salmon, DESeq2) and identified regulatory gene changes; performed pathway enrichment (GSEA) to highlight disease-relevant immune networks

Genome Assembly and Variant Calling for SRR2584868 | Northeastern University

Jan. 2025

• Designed an NGS pipeline (Trimmomatic, FastQC, STAR, bcftools) for genome assembly, alignment, and variant calling; visualized genetic variation in IGV and documented workflow on GitHub

CERTIFICATIONS

• Data Science & Analytics: DataCamp (Tidyverse, tidymodels, dplyr, ggplot2, SQL)

Nov. 2024 Dec. 2022

• Bioinformatics & Life Sciences: Ethical Edufabrica (Cancer Biology, Molecular Biology, Pharmacovigilance, Vaccine Technology)

E 1 2022

• AWS Academy: Machine Learning Foundations

Feb. 2022