

# DOMINIQUE DANG

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## Education

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Massachusetts Institute of Technology B.Sc. Computer Science & Molecular Biology	May 2027
<i>Classwork:</i> Machine Learning in Molecular & Cellular Biology, Biostatistics, Algorithms, Probability & Statistics, Linear Algebra, Cell Biology	

## Research & Work Experience

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Novartis Biomedical Research Institute <i>Oncology Data Science Intern</i>	May 2025 – Aug. 2025
<ul style="list-style-type: none"><li>Developed a predictive ML model in Python/R that achieved 85% accuracy in forecasting tumorigenesis outcomes by integrating differential expression (DESeq2) and enrichment (GSEA) features, uncovering 200+ biomarkers significantly associated with tumor formation</li><li>Applied gene signature scores to single-cell RNA-seq data from human breast cancer tumors to identify malignant subpopulations with high engraftment potential</li></ul>	

Undergraduate Research <i>Prof. Eliezer Calo - MIT</i>	Jan. 2025 – Aug. 2025
<ul style="list-style-type: none"><li>Analyzed nanopore sequencing data to resolve tRNA splicing patterns and detect RNA modifications</li><li>Designed and optimized bioinformatics pipelines incorporating tools such as NanoPlot (read QC), BWA (read alignment), and DESeq2 (differential expression/statistical analysis)</li></ul>	

Undergraduate Research <i>Prof. Anders Hansen - MIT</i>	Jan. 2024 – Dec. 2024
<ul style="list-style-type: none"><li>Characterized and validated bidirectional gene promoters to develop a novel tool for gene co-regulation, by utilizing Fluorescence-Activated Cell Sorting (FACS) and MATLAB to analyze protein expression levels and identify significant patterns and trends</li></ul>	

## Publications

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J. Clarin, **Dang, Dominique**, L. Santos, and R. Amini. Mechanical characterization of porcine tricuspid valve anterior leaflets over time: Applications to ex vivo studies. *ASME Open Journal of Engineering*, 2:021032, 05 2023. ISSN 2770-3495. doi:[10.1115/1.4062477](https://doi.org/10.1115/1.4062477).

## Leadership & Extracurriculars

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HackMIT <i>Logistics Director</i>	Feb. 2024 – present
<ul style="list-style-type: none"><li>Led a 15-member subteam to organize HackMIT, one of the largest U.S. collegiate hackathon, enhancing operational efficiency and streamlining the review process for 3,000+ applications</li></ul>	

MIT Museum <i>STEM Educator</i>	March 2023 – present
<ul style="list-style-type: none"><li>Facilitated the Maker Hub and Learning Lab, encouraging and educating over 50 daily visitors on STEM topics through interactive activities and hands-on demonstrations</li></ul>	

## Technical Skills

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Programming	Python (pandas, scikit-learn), R (Seurat, ggplot2), MATLAB, Bash, Git
Bioinformatics	RNA-seq analysis, DESeq2, FASTQ/FASTA, SAMtools, IGV, Scanpy
Lab Techniques	PCR, gel electrophoresis, Gibson Assembly, flow cytometry

## Awards

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1 <sup>st</sup> place, Innovation in Research: Data Science	Received at Novartis Summer of Science Internship.	2025
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Genetic Medicine Scholar	2025
Selected as 1 of 12 MIT sophomores for Eli Lilly's competitive research fellowship, contributing to disease pathology mapping, therapeutic target identification, and early-stage drug development strategy design (pre-clinical study planning, therapeutic design, and regulatory submissions).	

Top Beginner Hack	Received at HackMIT, one of the largest collegiate hackathons in the US.	2023
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