

# DOMINIQUE DANG

Cambridge, MA ◇ 781-980-3797 ◇ [ddang@mit.edu](mailto:ddang@mit.edu) ◇ [dom-dang.github.io](https://dom-dang.github.io)

## Education

Massachusetts Institute of Technology

May 2027

B.Sc. Computer Science & Molecular Biology

*Classwork:* Machine Learning in Molecular & Cellular Biology, Biostatistics, Algorithms, Probability & Statistics, Linear Algebra, Cell Biology

## Research & Work Experience

Novartis Biomedical Research Institute *Oncology Data Science Intern*

May 2025 – Aug. 2025

- 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
- Applied gene signature scores to single-cell RNA-seq data from human breast cancer tumors to identify malignant subpopulations with high engraftment potential

Undergraduate Research *Prof. Eliezer Calo - MIT*

Jan. 2025 – Aug. 2025

- Analyzed nanopore sequencing data to resolve tRNA splicing patterns and detect RNA modifications
- Designed and optimized bioinformatics pipelines incorporating tools such as NanoPlot (read QC), BWA (read alignment), and DESeq2 (differential expression/statistical analysis)

Undergraduate Research *Prof. Anders Hansen - MIT*

Jan. 2024 – Dec. 2024

- 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

## Publications

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

HackMIT *Logistics Director*

Feb. 2024 – present

- 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

MIT Museum *STEM Educator*

March 2023 – present

- Facilitated the Maker Hub and Learning Lab, encouraging and educating over 50 daily visitors on STEM topics through interactive activities and hands-on demonstrations

## Technical Skills

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

1<sup>st</sup> place, Innovation in Research: Data Science Received at Novartis Summer of Science Internship.

2025

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