# **DOMINIQUE DANG**

#### Cambridge, MA | 781-980-3797 | ddang@mit.edu | dom-dang.github.io

#### **EDUCATION**

Massachusetts Institute of Technology, Cambridge, MA

MAY 2027

**Bachelor of Science**: Computer Science & Molecular Biology

Relevant Courses: Biochemistry, Genetics, Cell Biology, Algorithms, Probability & Statistics, Linear Algebra,

**Fundamentals of Programming** 

# **RESEARCH & PROFESSIONAL EXPERIENCE**

Oncology Data Science Intern | Novartis Biomedical Research Institute

MAY — AUG 2025

- Built a predictive machine learning model in Python and R achieving 85% accuracy in forecasting tumorigenesis outcomes for human cancer cell lines in xenograft mouse models
- Engineered predictive features by integrating differential gene expression analysis (DESeq2) and gene set enrichment analysis (GSEA), identifying 200+ biomarkers that significantly correlated 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 Researcher** | MIT— Eliezer Calo Lab

IAN, 2025 — PRESENT

- Analyzing 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)

### **Genetics Medicine Scholar** | Eli Lilly

IAN. 2025

- Conducted disease pathology mapping and target identification using molecular profiling and preclinical models to discover novel therapeutic targets
- Designed and executed drug development strategies including preclinical study planning, therapeutic design, and regulatory submissions to advance therapies toward clinical approval

#### **Undergraduate Researcher** | MIT — Anders Hansen Lab

DEC. 2024 — FEB. 2025

 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

#### **Young Scholar's Program** | Northeastern University — Rouzbeh Amini Lab

JUNE— AUG. 2022

• Investigated mechanical properties of the tricuspid valve in porcine hearts through biaxial mechanical testing to analyze stress-strain relationships, contributing as second author to a **publication** 

# **LEADERSHIP & WORK EXPERIENCE**

**HackMIT** — Logistics Director

FEB. 2024 — PRESENT

• Led a 15-member subteam to organize HackMIT, the largest U.S. collegiate hackathon, enhancing operational efficiency and streamlining the review process for 3,000+ applications

#### **MIT Museum** — *STEM Educator*

MAR. 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

# **SKILLS & AWARDS**

**Programming & Data Science:** Python (pandas, scikit-learn), R (ggplot2, Seurat), MATLAB, Bash, Jupyter, Git **Bioinformatics:** FASTQ/FASTA, SAMtools, BLAST, IGV, StringDB, Scanpy

**Lab Skills:** PCR, gel electrophoresis, Gibson assembly, DNA purification, tissue culture, flow cytometry **Awards:** 1st Place – Innovation in Research: Data Science (Novartis), HackMIT Top Beginner Project