Jai Mehta

☑ jmm@utk.edu 🔗 volweb2.utk.edu/~jmehta2/ in mehtajm 🕠 mehtajm

Education

University of Tennessee

Aug. 2023 - May 2027

BS in Honors Computer Science, Chemistry

GPA: 4.00/4.00

Minors in Pre-Health Professions, Molecular Biophysics

May 2023

Wilson Central High School Salutatorian, High School Diploma

GPA: 4.55/4.00

Coursework

Computer Science Courses: Honors Algorithms, Data Structures, Computer Organization, Discrete Mathematics, Honors Probability and Statistics, Linear Algebra, Honors Calculus I & II

Medical Courses: Cellular and Molecular Biology, General Chemistry (with labs), Organic Chemistry I & II (with labs), Biochemistry, Honors Physics I & II (Calculus-Based)

Experience

Researcher and Software Developer

Osaka, Japan

Systems Immunology Lab, Osaka University Immunology Frontier Research Center

Dec. 2024 - Present

- Building a Clonotype to Paratope converter using IMGT and MiXCR inputs that takes a V gene name, J gene name, and CDR3b sequence, and provides the complete amino acid sequence. [GitLab]
- Using MAFFT to number antibodies and adaptive immune repertoires with unprecedented accuracy and speed. Wrote sequence parsers to build insertions for sequences using compact maps in C++.

Teaching Assistant

Knoxville, TN

Introductory Computer Science in C++, COSC 102

Aug. 2024 - Present

- Hosted weekly office hours to assist students with labs for a cohort of over 200 students.
- Assisted in topics related to learning Linux, basic data structures, dynamic memory, recursion, file I/O, searching algorithms, and binary logic.

Research Member

Knoxville, TN

Emrich Lab

Aug. 2024 - Present

- Utilizing scRNA-seq data to further understand cell state plasticity.
- Building a Graph Neural Network (GNN) to predict pathogenicity predisposition from genome sequences through graph k-mer representation.

Research Intern

Oak Ridge, TN

Oak Ridge National Lab, Center for Molecular Biophysics

Nov. 2023 - Present

- Tested an in-house Machine Learning model against other models to evaluate Area Under the Receiver Operating Characteristic Curve (AUC-ROC) scores regarding prediction of T Cell Receptor to epitope binding specificity.
- Currently using AlphaFold-2 to predict MHC to peptide binding affinity.
- Created a parser to symmetrize Poly(Hexamethylene Adipamide) (Nylon66) Hydrolase Nyl50 tetramer in Python.

Research Intern

Memphis, TN

Yu Lab, St. Jude Children's Research Hospital

July 2024 - Aug. 2024

- Evaluated a multi-informational clustering algorithm using scATAC-seq data with scRNA-seq labels to
 observe misclassification levels between cell phenotypes. Utilized Euclidean and Cosine distance estimation
 methods.
- Tested "Memory-like" and "Effector-like" cell states defined by a Gaussian Mixture Model using Joint Embedding, Differential Expression, and Gene Set Enrichment Analysis on clinical CAR T data.

• Combined clinical CAR T data, ran through Gaussian Mixture Model, and tested to observe similarities between individual clinical datasets and combined data.

Presentations and Talks

Defining Effector and Memory-like Cell State In in-vivo CAR T Cells from Multi-center Clinical Trials

- o St. Jude Intern Presentation Day, Memphis, TN, Aug. 2024.
- o UTK Undergraduate Research and Fellowships Discovery Day, Knoxville, TN, Sep. 2024.
- o SASE National Conference, Computational Sciences Division, Boston, MA, Oct. 2024.

Evaluating a Multi-Informational Clustering Algorithm using scATAC-seq and scRNA-seq

 $\circ\,$ St. Jude Intern Presentation Day, Memphis, TN, Aug. 2024.

Panelist for Student Panel for Undergraduate Research, Knoxville, TN, Nov. 2024.

Grants

University of Tennessee Honors Grant Recipient o Awarded \$750 to pursue research endeavors.	2025
Selected Awards	
Tau Beta Pi Honors	2024
Cook Grand Challenge Engineering Honors	2023
Chancellor's University Honors	2023
National Merit Scholar	2023
Distinguished Tennessean Scholar	2023
Ned McWherter's Scholar	2023
John Tummins Memorial Scholarship	2023
National Youth Science Camp Attendee	2023
Eagle Scout	2022

Programming Languages: C++, Python, R, Assembly, Java, Bash,

Skills: SciKit-Learn, ScanPy, Seurat, MacOS, Windows, Linux, Raspberry Pi, Arduino, Vim, Jupyter Notebook, GitHub, LaTeX, Single Cell Analysis, Machine Learning, High Performance Computing, Google Colab, AlphaFold, PyTorch.