# **Tyler Jay Yang**

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### **EDUCATION AND HONORS**

Carnegie Mellon University, School of Computer Science Pittsburgh, PA

Expected May 2027

Bachelor of Science, Computer Science, GPA: 4.0 | Dean's List

Honors: 2x American Invitational Mathematics Examination (AIME), National Merit Scholarship Recipient

**Relevant Coursework:** Linear Algebra, Vector Calculus, Probability & Statistics, Data Structures & Algorithms, Discrete Math, Theoretical CS, Computer Systems, Deep Learning, Computational Biology, Quantum Computation

#### **EXPERIENCE**

# **Computational Chemistry Research Assistant**

February 2025 - Present

Mechanical and Artificial Intelligence Lab, Carnegie Mellon University

- Wrote transformer encoder and decoder from scratch in *PyTorch* for solubility estimation and retrosynthetic prediction from SMILES strings
- Integrated named-entity recognition and natural language processing tools for text mining of chemistry literature
- Developed synthetic data generation pipeline to generate >100,000 images to pretrain deep learning models for reaction diagram parsing

## **Machine Learning Intern**

December 2022 - August 2023

Hillman Cancer Center, University of Pittsburgh Medical Center

- Created *ImageJ Java scripts* to preprocess fluorescence microscopy images to train *MATLAB* image segmentation algorithm to isolate thousands of asymmetrically dividing T cell clusters
- Achieved 90% accuracy, outperformed Sartorius commercial software

### **PUBLICATIONS**

Shiqi Xu, **Tyler J. Yang**, Suhong Xu, Yi-Nan Gong. *Plasma membrane repair empowers the necrotic survivors as innate immune modulators.* Seminars in Cell & Developmental Biology, Volume 156, (2024) <a href="doi.org/10.1016/j.semcdb.2023.08.001">doi.org/10.1016/j.semcdb.2023.08.001</a>

### **PROJECTS**

## Mitigating Algorithmic Bias in Skin Cancer Detection

- Implemented per-epoch adaptive resampling in *Keras* to implement a debiasing variational autoencoder, achieving a **31% accuracy improvement** in cancer detection over traditional CNNs
- Automated hyperparameter search across 6 model architectures using Sklearn and Comet
- Processed and labeled **>10,000 images** using *Tensorflow* and Pandas, leveraging *HDF5* utilities to enable handling of big data with optimized memory usage and faster I/O operations

## **Quantum Algorithm Virtual Machine**

- Modeled probabilistic and quantum algorithms, simulating pbit and qubit state manipulations in C, upgraded simulation to support 100x more qubits using randomized Monte Carlo algorithms
- Designed libraries for qubit operations and string parsing, ensuring correctness with programming contracts and optimizing memory management using *Valgrind*

## **Sequence Analysis Toolkit**

- Developed pathfinding and optimization algorithms for genome assembly and sequence alignment
- Built Hidden Markov Models using *NumPy* and implemented the Viterbi algorithm for genome reconstruction
- Performed Principal Component Analysis (PCA) in R with *ggplot* to classify SARS-CoV-2 variants

#### **ACTIVITIES**

Board Member, Carnegie Mellon Computational Biology Undergraduate SocietyJanuary 2025 – PresentFLIQ Hackathon Planning Team, IDQuantique Innovation Track, Quantum CoalitionApril 2025 – May 2025Quantum Coalition Delegate, Carnegie Mellon Quantum Computing ClubAugust 2024 – PresentFutsal ClubOctober 2024 – PresentLab Supervisor, University of Memphis Satellite CampusAugust 2022 – May 2023

#### SKILLS

Languages: Python, C, C++, JavaScript, Java, Julia, R, MATLAB

Machine Learning and Big Data: Tensorflow/Keras, PyTorch, Sklearn, Pandas, NumPy, Matplotlib, Flask, SQL Other: Git, GitHub, LaTeX, BASH scripting, Blender, Arduino, SolidWorks CAD, 3-D Printing, SLURM