# Minh Dao

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#### EDUCATION

## University of South Florida

Expected Graduation December 2026

Bachelor of Science in Computer and Information Science

Cumulative GPA: 3.91/4.00

- Awards: USF Green & Gold Presidential Scholarship (2023), Gold Prize in International Creative Papers Conference & Olympic (2022)
- Coursework: Machine Learning Specialization, Large Language Model Operations (LLMOps), Data Structures & Algorithms, Operating Systems, Parallel Computing, Database Design, Probability & Statistics, Linear Algebra

#### TECHNICAL SKILLS

Languages: Python, SQL, C/C++, JavaScript, TypeScript, HTML/CSS

Frameworks: PyTorch, TensorFlow, Langchain, FastAPI, React, Node.js, Express.js, Next.js, Vue.js, Flask Libraries: Scikit-learn, OpenAI, OpenCV, SciPy, Pandas, Numpy, Matplotlib, Huggingface, Shiny, SQLAlchemy

Tools: Git, AWS, Azure, Apache Airflow, MLflow, Databricks, DVC, Docker, ONNX, WandB, Postman, Netlify, Vercel

### EXPERIENCE

#### Moffitt Cancer Center

Feb. 2025 – Present

ML Research Intern

Tampa, Florida

- Designed a pipeline using **Python**, **Shiny**, and **Scikit-learn** to identify immune and pharmacokinetic predictors of post-transplant complications in **93 thalassemia patients**
- Boosted model accuracy by 30% through engineering 124 clinical features via univariate analysis with SciPy, validated separability with PCA/t-SNE, and identified 10 key predictors using permutation importance
- Implemented 9 Scikit-learn models to predict post-transplant immune rejection risk, achieving **0.89 F1** and 0.97 AUC with **AdaBoost**, and **0.80 F1** with **Logistic Regression**; tracked experiments and performance using **MLflow**
- Utilized **Pandas** and **Kaplan-Meier curves** to analyze post-transplant survival, revealing a statistically significant difference of **0.06 p-value** between patients with and without immune rejection

## **USF** Computational Biophysics Lab

Oct. 2024 – Present

Research Assistant

Tampa, Florida

- Developed an **AI model** for ion binding site prediction using **PyTorch**, **Biopython**, and **ESM2** Protein Language Model with few-shot learning; fine-tuned final **transformer** layer with single-layer **MLP**, achieving **0.8 F1** score
- Improved 15% data quality by recovering missing residues using internal libraries, removing 90%-similar sequences via CD-HIT, and addressing class imbalance with Focal Loss and ReduceLROnPlateau scheduler to prevent overfitting
- Accelerated preprocessing by 15% through parallelizing labeling and cleaning of 4,000 PDB files to JSON using multiprocessing; saved 10% GPU memory by enhancing data loader with dynamic batching based on sequence length

# FPT Software AI Center

May. 2024 – Sep. 2024

AI Engineer Intern

Hanoi, Vietnam

- Optimized a **text-image prompt** model using **PyTorch**, **Flask**, and **Stable Diffusion**; developed and deployed a client supporting dynamic user input with custom style templates on a high-traffic site serving **50K**+ users
- Enhanced **UNet** with a decoupled cross-attention layer for image features and integrated **object localization loss** to enable **multi-face ID** generation, boosting image quality, consistency, and identity preservation by up to **20**%
- Built a scalable image-text pipeline with **Apache Airflow** and **DVC** to preprocess **70K** images using BLIP-2 and Mask2Former; deployed the model as a low-latency microservice via **TorchServe** on **AWS ECS** with **API Gateway**

# PROJECTS

STEM Bot | Python, NextJS, FastAPI, Langchain, SQLAlchemy, PostgreSQL, Microsoft Azure, OpenAI

May. 2025

- Engineered a multi-modal STEM chatbot with real-time chat capabilities using NextJS and FastAPI
- Increased response relevance by 25% by deploying a RAG system with Langchain on Microsoft Azure, orchestrating OpenAI and Hugging Face models for text-image routing in Dockerized environments
- Reduced 40% message latency by parallelizing data processing across asynchronous pipelines with Redis Streams

GenFlow | Python, PyTorch, Numpy, Hugging Face, NVIDIA NeMo, ONNX, WandB

Oct. 2024

- Assisted Prof. Tassef Rahman in research on generative flow models for protein conformation design
- Improved 12% structure quality by fine-tuning Transformer blocks as a flow model using pre-trained ESM and a custom trunk decoder, integrating embedded structural features to guide the generation process.
- Enhanced model inference speed by 10% by integrating a D-Flow module that optimizes noise distribution