# Tang, Zijia (Thomas)

Email: zijia.tang@duke.edu Phone: +1 (858) 340-5757 Interest area: Artificial Intelligence, Large Language Model, Bioinformatics

#### **EDUCATION**

Duke University

Computer Science Major

Aug. 2024 – On going

GPA: 4.0 / 4.0

#### **PUBLICATION**

scPerb: Single-cell Perturbation via Style Transfer-based Variational Autoencoder | First Author | Published in Journal of Advanced Research (Impact Factor: 11.79)

- Designed and developed scPerb, a novel deep learning framework that accurately predicts single-cell transcriptional responses to gene or drug perturbations by leveraging a hybrid of style transfer and variational autoencoder (VAE) architectures. Reduced the need for expensive and labor-intensive wet-lab experiments.
- Demonstrated that scPerb achieves **state-of-the-art accuracy (99.5%)**, significantly outperforming existing models.
- Presented both a 5-minute oral talk and a poster presentation at the MCBIOS 2025 Conference, receiving positive feedback from domain experts for its innovation.
- Published in the high-impact, peer-reviewed *Journal of Advanced Research* (DOI: 10.1016/j.jare.2024.10.035).

PINet: Privileged Information Improves the Interpretability and Generalization of Structural MRI in Alzheimer's Disease | Independent Researcher | First Author | Presented at ACM-BCB 2024

- Proposed PINet, a novel hybrid deep learning architecture integrating Convolutional Neural Networks (CNNs) and Transformers, augmented with privileged information during training to enhance interpretability and generalizability in structural MRI analysis.
- Achieved **nearly 96% classification accuracy** in **early** Alzheimer's Disease detection, outperforming traditional deep learning baselines on benchmark datasets.
- Accepted as a rapid-fire paper by the ACM Conference on Bioinformatics, Computational Biology, and Health Informatics (ACM-BCB), awarded to the **top 25%** of submissions.
- DOI: 10.1145/3584371.3613000

Interpretation of interstitial lung diseases from magnetic resonance image using deep learning | Independent researcher | First Author | Published in AEIS 2022 Conference

- Developed ConvT, an end-to-end deep learning pipeline for differentiating Nonspecific Interstitial Pneumonia (NSIP) and Usual Interstitial Pneumonia (UIP) using non-invasive MRI scans, addressing diagnostic ambiguity in clinical radiology.
- Published at the **2022 International Conference on Advanced Engineering and Intelligent Systems (AEIS),** DOI: 10.1109/AEIS59450.2022.00027.

### **PROJECTS**

## Desktop Video for mac | Language: Swift, SwiftUI, AVFoundation

- Engineered a lightweight macOS application enabling users to set videos and images as dynamic desktop wallpapers across multiple monitors.
- Implemented support and features like HDR playback, real-time volume control, stretch-to-fit rendering, and seamless multi-screen synchronization, status persistence, and bookmark access to support sandboxed macOS environments.

## SSAPAutoCalendar | Language: Python, HTML

- Designed a backend utility to automatically retrieve, parse, and update academic calendar data from dynamic HTML pages hosted on university and high school servers.
- Implemented robust HTML parsing and scheduling logic to transform unstructured web content into structured, machine-readable formats.

### WORK AND RESEARCH EXPERIENCE

High school & Undergraduate Intern at Song Lab (Department of Health Outcomes & Biomedical Informatics at the University of Florida)

May 2023 – Present Undergraduate Intern at Yi Zhang's Lab (Department of Neurosurgery and Department of Biostatistics and Bioinformatics at Duke University)

Aug 2024 – Present