

# Xu Zhang

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## EDUCATION

### New York University

New York, NY

*The Master in Computer Science : GPA: 3.67*

*September 2024 – May 2026*

### The University of Texas at Austin

Austin, TX

*B.S. in Mathematics (with honor): GPA: 3.95* • 75th Annual Natural Sciences College Scholars (2023)

*January 2021 – May 2024*

## **KEYWORDS: Multimodal Learning, Vision-Language Model, Embodied Learning**

## SKILLS

**Programming Languages:** C, UNIX/Bash, Python, Java, MATLAB.

**Machine Learning:** Pytorch, Hugging Face, Scikit-learn, Numpy, Pandas, R.

**Other:** LaTeX, Diamond Sutra, I Ching, Windmill, Thomas Flare, Bowtie2, PCR reactions.

## RELATED COURSES TAKEN

MACHINE LEARNING, PREDICTIVE ANALYTICS, MATH STATISTICS, REAL ANALYSIS I&II, NUMERICAL ANALYSIS

## RELATED PROJECTS

Reimplemented MViT, YOLOv6 with few-shot learning, Prototypical Network, Alexnet

## RESEARCH&INTERNSHIP EXPERIENCE

### Capgemini Automobile Manufacturer LLM-based Market Analysis Pipeline

Shanghai, China

*AI Engineering Assistant (NLP)*

*June 2024 – August 2024*

- Co-developed a FastAPI server for hosting fine-tuned LLMs to extract key insights from market opinion on car parts in the client's products.
- Led the experiments on evaluating how vLLM enhances models' performance (including Qwen, Chatglm, etc.) and manages multi-thread requests through experiments on the FastAPI server we developed.
- Discovered and verified the crucial observation that using vLLM will result in outputs different from original models given fixed parameters due to vLLM's different implementation of attention kernel.

### Gene Expression Study in Binge-like Alcohol-Drinking Mice Across Brain Regions

Austin, TX

*Research Assistant (Supervised by Professor Dhivya Arasappan)*

*February 2023–June 2024*

- Conducted concordance and discordance analyses to identify genes with closely aligned expression levels across multiple brain regions.
- Executed pathway analysis to identify responsible pathways for differentially expressed concordant genes and those with closely aligned expression levels in multiple brain regions.
- Presented the work at Heart of Texas Research Conference.

## CONFERENCES & EVENTS

### Heart of Texas Undergraduate Research Conference

*Presenter*

*April 20, 2024*

- Presented our research on cross-brain-regions gene expression profiling in binge-drinking mice (High Drinking in the Dark line).
- Demonstrated a novel statistical method I developed to identify genes with similar expression patterns across brain regions and mouse lines by utilizing concordant pairs and a novel notion of distance.
- Presented a way to use permutation importance to select the most influential features from machine learning models for sample group prediction tasks, in order to proceed with further biological analysis.