Ziqi (Amber) Tang

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9 One Bungtown Road, Cold Spring Harbor, NY

Education ____

Cold Spring Harbor Laboratory

Ph.D. Candidate in Biology

Fall 2019 - Current

Expected graduation May 2024

University of North Carolina at Chapel Hill

BS. in Biology; BS. in Computer Science

Fall 2015 - Spring 2019

Graduated with Distinction and Highest Honors

Research Experience _

Peter Koo Lab at Cold Spring Harbor Laboratory

Ph.D. candidate

April 2020 - Current

- · Assessed transfer learning with transformer-based language models for various RNA regulation prediction tasks
- Prototyped a multi-task convolutional network that predicts various RNA regulatory functions from primary sequence
- Developed a framework to comprehensively evaluate the generalization and interpretability of convolutional networks for predicting epigenomics data
- Deployed a deep convolutional network to model single-cell ATAC-seq data and performed model interpretability methods to uncover learned cell-type specific features

Camford Capital

Venture Fellow July 2023 - Oct 2023

- Prototyped algorithms for Deep Learning guided sequence design
- Participated in internal venture initiatives at the interface of Computation, Artificial Intelligence and Biomedicine

Terry Furey Lab at Department of Genetics, UNC Chapel-Hill

Research Assistant Fall 2017 - Spring 2019

- Analyzed ATAC-seq data from patients with Crohn's disease
- Investigated the role of allelic imbalance expression in inflammatory tissue

Dayan Network Neuroscience Lab at Department of Radiology, UNC Chapel-Hill

Research Assistant Fall 2017 - Spring 2019

- Differentiated prodromal Parkinson's disease patients from healthy individuals fMRI images using SVM
- Generated brain connectivity matrix for Parkinson's Disease patients

Publications and Conferences _____

Publications

- Toneyan, S.*, **Tang, Z.***, Koo, P., *Evaluating deep learning for predicting epigenomic profiles*, Nature Machine Intelligence, Dec 2022. [Link]
- Kawaguchi, R., Tang, Z., Fischer, S., Rajesh, C., Tripathy, R., Koo, P., Gillis, J., *Learning single-cell chromatin accessibility profiles using meta-analytic marker genes*, Briefings in Bioinformatics, Dec 2022. [Link]
- Majdandzic, A., Rajesh, C., **Tang, Z**., Toneyan, S., Labelson, E., Tripathy, R., Koo, P., *Selecting deep neural networks that yield consistent attribution-based interpretations for genomics*, Proceedings of the 17th Machine Learning in Computational Biology meeting, Dec 2022. [Link]
- Lee, N., Tang, Z., Toneyan, S., Koo, P., EvoAug: improving generalization and interpretability of genomic deep neural networks with evolution-inspired data augmentations, Genome Biology, May 2023. [link]

Conferences and Presentations

Poster presenter

- Building foundation models for regulatory genomics requires rethinking large language models; International Conference on Machine Learning Computational Biology Workshop 2023
- Benchmarking Binary and Quantitative Genomic Models; Biology of Genomes 2022
- Evaluating deep learning for predicting epigenomic profiles; Intelligent Systems for Molecular Biology 2022
- Evaluating deep learning for predicting epigenomic profiles; American Society of Human Genomics 2022

Professional Service

Diversity Initiative for the Advancement of STEM at CSHL

E-board member Spring 2022 - Current

· Organized seminars and discussions to support and increase the presence of underrepresented minorities in STEM fields

Reviewer

· Nature Genetics, Nature Methods, Bioinformatics, Machine Learning in Structural Biology workshop, NeurIPS 2022, International Conference on Research in Computational Molecular Biology 2024

Programm Committee

• Machine Learning in Computational Biology 2022

Additional Experience _____

CSHL Postbaccalaureate Research Education Program

Mentor

Summer 2023 - Current

• Provided mentoring to a PREP scholar, guiding their research project and supproting the preparation of graduate school applications.

CSHL Undergraduate Research Program

Mentor Summer 2021-2023

• Lead lectures in Programming Course, introducing methods in biological data analysis for 20+ undergraduate students

CIFAR Deep Learning + Reinforcement learning Summer School

Participant July 2021

• Participated in lecture and discussions about cutting-edge topics in machine learning

Computer Science department, UNC Chapel-Hill

Teaching Assistant Fall 2017 - Fall 2018

• Assisted teaching in Data Structure and Introduction to Algorithm for 100+ students

Technical Skills _____

- Coding in Python, C, java, SQL
- Frameworks including TensorFlow, PyTorch and Git
- Experience with developing machine learning models, evaluating their performance, and model interpretation for scientific discovery on high performance computing clusters (UGE, Slurm)