

# Tian XIA

## Machine Learning Scientist

☎ +1 609 375 5853

@ tianx@bcm.edu

in linkedin.com/in/tian-xia-7a6639148

🏠 Personal website

🐙 github.com/no1summer

📍 2300 Old Spanish Trl, Houston, TX 77054 - USA

I am currently a cross-disciplinary researcher with expertise in machine learning, imaging analysis, and computational biology. My Ph.D. research focused on developing quantitative measurements for 3D+time images. It equips me with the knowledge and experience of image acquisition, processing and analysis in both temporal, spatial and frequency domain. With a solid background in math, physics, computer science and biology, I have the skills and expertise to develop and apply Machine Learning/ Deep Learning to address biological questions and beyond. I have developed several models that have been deployed in both industrial environment or for academical purpose, in a wide range of regression, classification, clustering, segmentation, and more.

## SKILLS

Programming	Python, R, MATLAB, C, Bash, SQL, Git, LaTeX
Data Science	Pandas, Scikit-Learn, PyTorch, AnnData, Seurat
Imaging Analysis	OpenCV, Numpy, Scipy, ImageJ, Scikit-Image, Matlab Image Processing Toolbox
Data Visualization	Matplotlib, Seaborn, ggplot2
Research	Data Science, Machine Learning, Deep Learning, Imaging Processing and Analysis, Computational Biology
Soft Skills	Critical Thinking, Active Learning, Time Management, Communication

## EXPERIENCE

Jun 2024 Jan 2024	<b>Merck - CBGX   CAMBRIDGE SITE - Co-op/intern</b> <i>Merck Research Laboratories – MENTOR : DR. REBECCA SENFT</i> <ul style="list-style-type: none"><li>➢ Fine tuning self-supervised <b>transformer</b>-based <b>deep learning</b> image feature extraction method DI-NOv2 for classifying cells with different treatment</li><li>➢ Classify RNA-seq data of patients' for different locations with <b>machine learning</b></li><li>➢ Developed a <b>machine learning</b> pipeline for determining the suitable marker for Optical Pooled <b>CRISPR Screening</b> (OPS)</li><li>➢ Established a CellProfiler and Cellpose (deep learning) pipeline to segment and quantify the cell images, which is integrated with Nextflow and deployed to High Performance Clustering (HPC).</li><li>➢ Experience with processing <b>scRNA-seq</b> data from fastq file</li><li>➢ Build and train <b>GPT2</b> from scratch and generate Shakespeare-like dialogue</li><li>➢ Explore different methods for integration of <b>multiomic/multimodality</b> data including scRNA-seq, ATAC-seq and imaging data.</li><li>➢ Fine tuning <b>scGPT</b> for multiomic/multimodality integration</li></ul> <div>Multiomic/Multimodality IntegrationMachine LearningDeep LearningLLMScikit-learnPytorchNextflowHPC</div>
Nov 2024 Dec 2019	<b>Baylor College of Medicine - LARINA'S LAB   IMAGING SCIENCE - Graduate Student</b> <i>Department of Integrative Physiology – MENTOR : DR. IRINA LARINA</i> <ul style="list-style-type: none"><li>➢ Constructed an <b>3D segmentation</b> procedure by fine-tuning pretrained <b>3D Swin Transformer</b> to quantify follicle volumes during the mouse ovulation process.</li><li>➢ Build an <b>semi-supervised deep learning</b> model for 3D segmentation, by introducing cross-attention block and mutual learning to the classic V-net backbone.</li><li>➢ Developing AI coach for tennis player based on <b>pose recognition</b> and <b>graph convolution network</b> from self-recording videos.</li><li>➢ Developed a quantitative imaging method of cilia metachronal wave in mouse fallopian tube with optical coherence tomography in vivo (Published at <b>Optica</b>, IF=10.4).</li><li>➢ Established a dynamic image signal processing procedure to track spermatozoa movement toward the egg (Invited oral presentation at <b>SPIE, 2022</b>).</li></ul> <div>Computer VisionDeep LearningObject DetectionSegmentationPose RecognitionCNNRNNGCNFourier TransformPhase Analysis</div>

Dec 2023 Jan 2020	<b>Rice University - CLASS   MACHINE LEARNING - Visiting Student</b> <i>Department of Computer Science – DATA SCIENCE PROJECT, STATISTICAL MACHINE LEARNING</i> <ul style="list-style-type: none"> <li>➢ Constructed a <b>machine learning pipeline</b> to identify genomic signatures in age-related macular degeneration. Reduced the number of feature from &gt;18000 to &lt;100 using <b>feature selection</b> techniques, including <i>minimum Redundancy Maximum Relevance, Random Forest, Generalized Linear Model, Principle Component Analysis, Statistical Test</i>.</li> <li>➢ Built a Python package for identifying the possible genes related to the disease from machine learning feature selection perspective.</li> <li>➢ Created a <b>classification network</b> (customized Resnet50 with Ensemble strategy) to distinguish fine-grained food images.</li> <li>➢ Construct a <b>generalized linear model</b> to identify individuals with the high risk of stroke with more than 90% accuracy.</li> </ul> <div> <span>Data Science</span> <span>Machine Learning</span> <span>Computational Biology</span> <span>Pandas</span> <span>Scikit-learn</span> <span>R</span> <span>Version Control</span> <span>Data Visualization</span> </div>
Apr 2019 Sep 2018	<b>Massachusetts Institute of Technology - WEINBERG'S LAB   CANCER BIOLOGY - Research Assistant</b> <i>Department of Biology – MENTOR : DR. ROBERT WEINBERG</i> <ul style="list-style-type: none"> <li>➢ Help build a genetically defined syngeneic mouse model of ovarian cancer (Published at <b>Cancer Discovery</b>)</li> </ul> <div> <span>Molecular Biology</span> <span>Gene Editing</span> <span>CRISPR</span> <span>Drug Testing</span> </div>
Jul 2018 May 2018	<b>Princeton University - KANG'S LAB   CANCER BIOLOGY - Research Assistant</b> <i>Department of Biology – MENTOR : DR. YIBIN KANG</i> <ul style="list-style-type: none"> <li>➢ Study the phenotype of mir200 knockout in the mouse model by Immunohistochemistry (IHC) imaging.</li> </ul> <div> <span>Immunohistochemistry Imaging</span> <span>Statistical Analysis</span> </div>
Feb 2018 Jul 2017	<b>Zhejiang University - SHAO'S LAB   CANCER BIOLOGY - Research Assistant</b> <i>School of Medicine – MENTOR : DR. JIMIN SHAO</i> <ul style="list-style-type: none"> <li>➢ Study the IL-6, p-stat3, Fra-1, Nanog pathway in the progression and metastasis of colon cancer by immunofluorescence (IF) imaging. (Published at <b>Oncogene</b>)</li> </ul> <div> <span>Immunofluorescence Imaging</span> <span>Cancer Research</span> </div>

## EDUCATION

2024	<b>Baylor College of Medicine</b> Ph.D. Quantitative & Computational Biosciences Concentration : Image Processing and Analysis, Machine/Deep Learning, Multimodality Integration
2019	<b>Zhejiang University</b> B.S. Pharmaceutical Sciences, GPA 3.97 Concentration : Molecular Biology, Cancer Biology, Computational Biology, Drug Discovery

## PUBLICATIONS

**In vivo volumetric depth-resolved imaging of cilia metachronal wave with dynamic optical coherence tomography** 2023  
[Tian Xia](#), Kohei Umezu, Deirdre Scully, Shang Wang, Irina Larina

 [Optica](#)

Imaging Processing Spatial and Temporal Imaging Dynamic Signal Processing Fourier Transform Phase

**Dynamic volumetric imaging and cilia beat mapping in the mouse male reproductive tract with optical coherence tomography** 2022

Kohei Umezu, [Tian Xia](#), Irina Larina

 [Biomedical Optics Express](#)

Imaging Analysis Volumetric 3D Imaging Dynamic Signal Processing Reproductive Biology

**Tracking spermatozoa movement toward the egg with functional optical coherence tomography** 2022

[Tian Xia](#), Kohei Umezu, Shang Wang, Irina Larina

 [Dynamics and Fluctuations in Biomedical Photonics XIX](#)

Object Detection Dynamic Signal Processing Denoising

**The inflammatory cytokine IL-6 induces FRA1 deacetylation promoting colorectal cancer stem-like properties** 2019

Tingyang Wang, Ping Song, Tingting Zhong, Xianjun Wang, Xueping Xiang, Qian Liu, Haiyi Chen, [Tian Xia](#), ..., Riccardo Fodde, Jimin Shao

 [Oncogene](#)

Cancer Immunology Pathway Immunofluorescence Imaging Imaging Analysis

## POSITION OF RESPONSIBILITY

## ACHIEVEMENTS & RECOGNITIONS

 PUBLIC MEDIA EXPOSURE3