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# Education

## • PhD in Computational Biology

Hong Kong Baptist University, Expected 2025

Advisor: Professor Liang Tian

# • Master of Engineering in Computer Technology

Shanghai Jiao Tong University, 2021

Advisor: Professor Hai Zhao

#### • Bachelor of Engineering in Bioengineering

Xi'an Polytechnic University, 2010

# Work Experience

#### • Research Assistant

Shenzhen Institutes of Advanced Technology (SIAT), CAS, 2020–2021

- Focused on the deconvolution of bulk cell RNA-seq data from primary solid tumors.
- Investigated the relationship between specific gene expression programs (e.g., EMT) and the abundance of infiltrated immune cells in the tumor microenvironment.

#### • Research Assistant

Interdisciplinary Research Center of Biology and Chemistry (IRCBC), CAS, 2016–2019

 Developed machine-learning-based prediction algorithms for CCS values and contributed to web server development (MetCCS and LipidCCS).

#### • R&D Department Member

Genminix Informatics Ltd. Co., 2011–2016

 Developed a method for quick search and visualization of feed-forward loops, leading to a Chinese patent (201410112193X).

# Teaching Experience

- Teaching Assistant, Thermal and Statistical Physics (PHYS3047), 2023–2024
- Teaching Assistant, Thermal and Statistical Physics (PHYS3047), 2022–2023

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## Presentations

• DeSide: A Unified Deep Learning Approach for Cellular Decomposition of Bulk Tumors Based on Limited scRNA-seq Data

Poster Presentation, 28th IUPAP International Conference on Statistical Physics, August 7-11, 2023, University of Tokyo, Tokyo, Japan

# **Publications**

\$ denotes equal contributions

- \* denotes corresponding author(s)
- Xiong, X.\$, Liu, Y.\$, Pu, D., Yang, Z., Bi, Z., Tian, L.\*, and Li, X.\* (2023). DeSide: A unified deep learning approach for cellular decomposition of bulk tumors based on limited scRNA-seq data. bioRxiv, 2023.05.11.540466. 10.1101/2023.05.11.540466.
- Zhou, Z., Luo, M., Chen, X., Yin, Y., Xiong, X., Wang, R., and Zhu, Z.-J.\* (2020). Ion mobility collision cross-section atlas for known and unknown metabolite annotation in untargeted metabolomics. *Nat. Commun.* 11, 4334.
- Shen, X., Wang, R., Xiong, X., Yin, Y., Cai, Y., Ma, Z., Liu, N., and Zhu, Z.-J.\* (2019). Metabolic reaction network-based recursive metabolite annotation for untargeted metabolomics. *Nat. Commun.* 10, 1516.
- Zhou, Z., Shen, X., Chen, X., Tu, J., **Xiong, X.**, and Zhu, Z.J.\* (2019). LipidIMMS Analyzer: integrating multi-dimensional information to support lipid identification in ion mobility—mass spectrometry based lipidomics. *Bioinformatics* 35.
- Zhou, Z., Tu, J., Xiong, X., Shen, X., and Zhu, Z.J.\* (2017). LipidCCS: prediction of collision cross-section values for lipids with high precision to support ion mobility–mass spectrometry-based lipidomics. *Anal. Chem.* 89, 9559–9566.
- Zhou, Z., Xiong, X., and Zhu, Z.-J.\* (2017). MetCCS predictor: a web server for predicting collision cross-section values of metabolites in ion mobility-mass spectrometry based metabolomics. *Bioinformatics* 33, 2235–2237.

## Skills

- Software: Python (9 years), R (participated a big R project), TensorFlow (8 years), PyTorch (1 year), MySQL, Docker
- Languages: Chinese (native), English (fluent)
- Data Analysis: Bulk RNA-seq and Single-cell RNA-seq, LC-MS, Machine Learning Methods, Statistical Methods

## References

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Shenzhen Institutes of Advanced Technology (SIAT), CAS

Email: xuefei.li@siat.ac.cn

# • Prof. Hai Zhao

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