Xin Xiong 1

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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

Xin Xiong 2

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

- 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 (project participant), TensorFlow (8 years), PyTorch (1 years), 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

• Prof. Liang Tian

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• Prof. Xuefei Li

Shenzhen Institutes of Advanced Technology (SIAT), CAS

Xin Xiong 3

 $Email: \ xuefei.li@siat.ac.cn$

• Prof. Hai Zhao

Shanghai Jiao Tong University Email: zhaohai@cs.sjtu.edu.cn