SHEN, XIAOTAO Ph.D.

Michael Snyder's Lab, Stanford School of Medicine

1 INFORMATION

Email: shenxt@stanford.edu Github: jaspershen

In LinkedIn: shenxt ↑ Homepage: http://shenxt.sxl.cn/

2 Address: M339, Always Building, Stanford School of Medicine, Stanford, CA 94305.

➡ EDUCATION & RESEARCH EXPERIENCE

> Postdoctoral Research Fellow (Advisor: <u>Dr. Michael Snyder</u>)

Apr 2019 – *present*, **Stanford University**.

> Research Scientist (Advisor: <u>Dr. Zheng-Jiang Zhu</u>)

Jan. 2019 – Mar. 2019, Chinese Academy of Sciences (CAS).

> **PhD** Student in Metabolomics and Bioinformatics (Advisor: <u>Dr. Zheng-Jiang Zhu</u>)

Aug. 2013 – Dec. 2018, Chinese Academy of Sciences (CAS).

> **BSs** in Biotechnology Aug. 2009 - Jun. 2013, **Inner Mongolia University**.

A RESEARCH

- > Metabolomics Data Processing Methods and Software Development.
- > Multi-omics Data Analysis for Biomarker Discovery for Diseases.

THONORS AND AWARDS

> Student Travel Award for Oral Presentation

The International Metabolomics Society (2018).

- > International Conference Travel Award
- The *Metabolites* Journal (2018).
- > China National Scholarship Ministry of Education of the People's Republic of China (2017).
- > Award for Outstanding Youth Report

The 3th China Mass Spectrometry Analysis Conference (2017).

- Merit Student University of Chinese Academy of Sciences (2016).
- > Award for Outstanding Youth Report

The 34th China Mass Spectrometry Society Conference (2016).

> Inner Mongolia Outstanding Graduate (2013).

Inner Mongolia Autonomous Region

> National Encouragement Scholarship

Inner Mongolia University (2011).

■ PUBLICATIONS

> Z. Wang, B. Cui, F. Zhang, Y. Yang, X. Shen, Z. Li, W. Zhao, Y. Zhang, K. Deng, Z. Rong, K. Yang, X. Yu, K. Li, P. Han, and Z.-J. Zhu, Development of A Correlative Strategy to Discover Colorectal Tumor Tissue Derived Metabolite Biomarkers in Plasma Using Untargeted Metabolomics, *Analytical Chemistry*, 2018, in press, 10.1021/acs.analchem.8b05177.

- > X. Shen, R. Wang, X. Xiong, Y. Yin, Y. Cai, J. Ma, N. Liu and Z.-J. Zhu, Large-scale Metabolite Identification for Untargeted Metabolomics Using Metabolic Reaction Network, *Nature Communications*, 2019, 10:1516.
- > **X. Shen** and Z.-J. Zhu, MetFlow: An interactive and integrated workflow for metabolomics data cleaning and differential metabolite discovery, *Bioinformatics*, 2019, in press, https://doi.org/10.1093/bioinformatics/bty1066.
- > Z. Zhou, X. Shen, X. Chen, J. Tu, X. Xiong, and Z.-J. Zhu, LipidIMMS Analyzer: Integrating Multi-dimensional Information to Support Lipid Identification in Ion Mobility–Mass Spectrometry based Lipidomics, *Bioinformatics*, 2018, in press, https://doi.org/10.1093/bioinformatics/bty661.
- > H. Jia, X. Shen (Co-first author), Y. Guan, M. Xu, M. Mo, J. Zhu and Z.-J. Zhu, Assessment of The Response to Neoadjuvant Chemo-Radiation in Rectal Cancer Patients based on A Metabolomics Approach, *Radiotherapy and Oncology*, 2018, 128, 548-556.
- > Z. Zhou, J. Tu, X. Xiong, X. Shen, and Z.-J. Zhu, LipidCCS: Prediction of Collision Cross-Section Values for Lipids with High Precision to Support Ion Mobility-Mass Spectrometry based Lipidomics, *Analytical Chemistry*, 2017, 89, 9559–9566.
- > Z. Zhou, X. Shen, J. Tu, and Z.-J. Zhu, Large-Scale Prediction of Collision Cross-Section Values for Metabolites in Ion Mobility Mass Spectrometry, *Analytical Chemistry*, 2016, 88, 11084-11091.
- > J. Wang, T. Zhang, X. Shen (Co-first author), J. Liu, D. Zhao, Y. Sun, L. Wang, Y. Liu, X. Gong, Y. Liu, Z.-J. Zhu, F. Xue,* Serum Metabolomics for Early Diagnosis of Esophageal Squamous Cell Carcinoma by UHPLC-QTOF/MS, *Metabolomics*, 2016, 12: 116.
- > X. Shen, X. Gong, Y. Cai, Y. Guo, J. Tu, H. Li, T. Zhang, J. Wang, F. Xue, and Z.-J. Zhu, Normalization and Integration of Large-Scale Metabolomics Data Using Support Vector Regression, *Metabolomics*, 2016, 12: 89.

□ ORAL PRESENTATIONS

Metabolic Reaction Network-based Recursive Metabolite Identification for Untargeted Metabolomics. The 14th International Conference of the Metabolomics Society, June, 2018, Seattle, USA.

- Assessment of the Response to Neoadjuvant Chemo-Radiation in Rectal Cancer Patients based on a Metabolomics Approach. The 3th China Mass Spectrometry Analysis Conference, December, 2017, Xiamen, China.
- > Normalization and Integration of Large-Scale Mass Spectrometry-based Metabolomics Data Using Support Vector Regression. **The 34**th **China Mass Spectrometry Society Conference**, September, 2016, Xining, China.
- Normalization and Integration of Large-Scale Mass Spectrometry-based Metabolomics Data Using Support Vector Regression. The 64th American Society for Mass Spectrometry Conference, June, 2016, San Antonio, USA.

POSTERS PRESENTATION

> Metabolic Reaction Network based Metabolite Annotation in Untargeted Metabolomics. **The** 13th International Conference of the Metabolomics Society, June, 2017, Brisbane, Austria.

1 TECHNICAL STRENGTH

- Languages: Mandarin (Very fluent), English (Fluent).
- Mass Spectrometry Analysis: Agilent QT0F 6550, Agilent QQQ 6495, 6460, Agilent IMMS 6560, AB sciex Triple TOF 5600, 6600.
- **Bioinformatic Tools:** Agilent MassHunter work station, AB sciex PeakView, XCMS, MS-DIAL.
- ♦ Other Skills: Markdown, Photoshop, Illustrator, Linux (Ubuntu and CentOS), GitHub ♠, Shiny ℯ.