

# Shipei Xing

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## Research Interests

Mass spectrometry, Bioinformatics, Metabolomics, Analytical chemistry

## Employment

2023.8 – Present

**Univeristy of California, San Diego** – La Jolla, US

Postdoctoral scholar

Research: *Mass spectrometry informatics & Collaborative Microbial Metabolite Center*

Supervisor: Pieter C. Dorrestein

## Education

2019.1 – 2023.4

**University of British Columbia** – Vancouver, Canada

PhD in Chemistry

Research: *Towards accurate compound annotation in mass spectrometry-based global metabolomics*

Supervisor: Tao Huan

2014.9 – 2018.6

**Zhejiang University** – Hangzhou, China

BSc in Chemistry (**Qiushi Honors Program**, Chu Kochen Honors College)

Research: *Synthesis of Chiral Imidazoline Iminopyridine Ligands* (Outstanding thesis award)

Supervisor: Zhan Lu

2017.3 – 2017.9

**University of Utah** – Salt Lake City, US

Visiting scholar

Supervisor: Peter J. Stang

## Featured Publications

[See full publication list here.](#)

- 2024 [Shipei Xing, Vincent Charron-Lamoureux, Måns Ekelöf, Yasin El Abiead, Huaxu Yu, Oliver Fiehn, Theodore Alexandrov, Pieter C. Dorrestein. Structural annotation of full-scan MS data: A unified solution for LC-MS and MS imaging analyses. \*bioRxiv\*. \(10.1101/2024.10.14.618269\)](#)
- 2023 [Shipei Xing, Sam Shen, Banghua Xu, Xiaoxiao Li, Tao Huan. BUDDY: molecular formula discovery via bottom-up MS/MS interrogation. \*Nature Methods\*. \(10.1038/s41592-023-01850-x\)](#)
- 2021 [Shipei Xing, Yibo Jiao, Melody Salehzadeh, Kiran K Soma, Tao Huan. SteroidXtract: deep learning-based pattern recognition enables comprehensive and rapid extraction of steroid-like metabolic features for automated biology-driven metabolomics. \*Analytical Chemistry\*. \(10.1021/acs.analchem.0c04834\)](#)

- 2020 Shipei Xing, Yan Hu, Zixuan Yin, Min Liu, Xiaoyu Tang, Mingliang Fang, Tao Huan. Retrieving and utilizing hypothetical neutral losses from tandem mass spectra for spectral similarity analysis and unknown metabolite annotation. *Analytical Chemistry*. ([10.1021/acs.analchem.0c02521](https://doi.org/10.1021/acs.analchem.0c02521))
- 2022 Shipei Xing, Tao Huan. Radical fragment ions in collision-induced dissociation-based tandem mass spectrometry. *Analytica Chimica Acta*. ([10.1016/j.aca.2022.339613](https://doi.org/10.1016/j.aca.2022.339613))
- 2021 Shipei Xing, Huaxu Yu, Min Liu, Qingquan Jia, Zhi Sun, Mingliang Fang, Tao Huan. Recognizing contamination fragment ions in liquid chromatography–tandem mass spectrometry data. *Journal of the American Society for Mass Spectrometry*. (Invited, JASMS ‘Emerging Investigators’ focus section, [10.1021/jasms.0c00478](https://doi.org/10.1021/jasms.0c00478))

## Selected Co-authored Publications

- 2024 Abzer K Pakkir Shah, Axel Walter, Filip Ottosson, Francesco Russo, Marcelo Navarro-Diaz, Judith Boldt, Jarmo-Charles J Kalinski, Eftychia Eva Kontou, James Elofson, Alexandros Polyzois, Carolina González-Marín, Shane Farrell, Marie R Aggerbeck, Thapanee Prusatrakul, Nathan Chan, Yunshu Wang, Magdalena Pöchhacker, Corinna Brungs, Beatriz Cámara, Andrés Mauricio Caraballo-Rodríguez, Andres Cumsille, Fernanda de Oliveira, Kai Dührkop, Yasin El Abiead, Christian Geibel, Lana G Graves, Martin Hansen, Steffen Heuckeroth, Simon Knoblauch, Anastasiia Kostenko, Mirte CM Kuijpers, Kevin Mildau, Stilianos Papadopoulos Lambidis, Paulo Wender Portal Gomes, Tilman Schramm, Karoline Steuer-Lodd, Paolo Stincone, Sibgha Tayyab, Giovanni Andrea Vitale, Berenike C Wagner, Shipei Xing, Marquis T Yazzie, Simone Zuffa, Martinus de Kruijff, Christine Beemelmanns, Hannes Link, Christoph Mayer, Justin JJ van der Hooft, Tito Damiani, Tomáš Pluskal, Pieter Dorrestein, Jan Stanstrup, Robin Schmid, Mingxun Wang, Allegra Aron, Madeleine Ernst, Daniel Petras. Statistical analysis of feature-based molecular networking results from non-targeted metabolomics data. *Nature Protocols*. ([10.1038/s41596-024-01046-3](https://doi.org/10.1038/s41596-024-01046-3))
- 2024 Ipsita Mohanty, Helena Mannochio-Russo, Joshua V Schweer, Yasin El Abiead, Wout Bittremieux, Shipei Xing, Robin Schmid, Simone Zuffa, Felipe Vasquez, Valentina B Muti, Jasmine Zemlin, Omar E Tovar-Herrera, Sarah Moraïs, Dhimant Desai, Shantu Amin, Imhoi Koo, Christoph W Turck, Itzhak Mizrahi, Penny M Kris-Etherton, Kristina S Petersen, Jennifer A Fleming, Tao Huan, Andrew D Patterson, Dionicio Siegel, Lee R Hagey, Mingxun Wang, Allegra T Aron, Pieter C Dorrestein. The underappreciated diversity of bile acid modifications. *Cell*. ([10.1016/j.cell.2024.02.019](https://doi.org/10.1016/j.cell.2024.02.019))
- 2024 Tingting Zhao, Nicholas JP Wawryk, Shipei Xing, Brian Low, Gigi Li, Huaxu Yu, Yukai Wang, Qiming Shen, Xing-Fang Li, Tao Huan. ChloroDBPFinder: Machine Learning-Guided Recognition of Chlorinated Disinfection Byproducts from Nontargeted LC-HRMS Analysis. *Analytical Chemistry*. ([10.1021/acs.analchem.3c05124](https://doi.org/10.1021/acs.analchem.3c05124))
- 2023 Wout Bittremieux, Nicole E Avalon, Sydney P Thomas, Sarvar A Kakhkhorov, Alexander A Ak-senov, Paulo Wender P Gomes, Christine M Aceves, Andrés Mauricio Caraballo-Rodríguez, Julia M Gauglitz, William H Gerwick, Tao Huan, Alan K Jarmusch, Rima F Kaddurah-Daouk, Kyo Bin Kang, Hyun Woo Kim, Todor Kondić, Helena Mannochio-Russo, Michael J Meehan, Alexey V Melnik, Louis-Felix Nothias, Claire O’Donovan, Morgan Panitchpakdi, Daniel Petras, Robin Schmid, Emma L Schymanski, Justin JJ van der Hooft, Kelly C Weldon, Heejung Yang, Shipei Xing, Jasmine Zemlin, Mingxun Wang, Pieter C Dorrestein. Open access repository-scale propagated nearest neighbor suspect spectral library for untargeted metabolomics. *Nature Communications*. ([10.1038/s41467-023-44035-y](https://doi.org/10.1038/s41467-023-44035-y))

- 2023 Tingting Zhao, Shipei Xing, Huaxu Yu, Tao Huan. De novo cleaning of chimeric MS/MS spectra for LC-MS/MS-based metabolomics. *Analytical Chemistry*. ([10.1021/acs.analchem.3c00736](https://doi.org/10.1021/acs.analchem.3c00736))
- 2023 Ting Fu, Tao Huan, Gibraan Rahman, Hui Zhi, Zhenjiang Xu, Tae Gyu Oh, Jian Guo, Sally Coulter, Anupriya Tripathi, Cameron Martino, Justin L McCarville, Qiyun Zhu, Fritz Cayabyab, Brian Low, Mingxiao He, Shipei Xing, Fernando Vargas, T Yu Ruth, Annette Atkins, Christopher Liddle, Janelle Ayres, Manuela Raffatellu, Pieter C Dorrestein, Michael Downes, Rob Knight, Ronald M Evans. Paired microbiome and metabolome analyses associate bile acid changes with colorectal cancer progression. *Cell Reports*. ([10.1016/j.celrep.2023.112997](https://doi.org/10.1016/j.celrep.2023.112997))
- 2023 Fanrong Zhao, Li Li, Penghui Lin, Yue Chen, Shipei Xing, Huili Du, Zheng Wang, Junjie Yang, Tao Huan, Cheng Long, Limao Zhang, Bin Wang, Mingliang Fang. HExpPredict: in vivo exposure prediction of human blood exposome using a random forest model and its application in chemical risk prioritization. *Environmental Health Perspectives*. ([10.1289/EHP11305](https://doi.org/10.1289/EHP11305))
- 2023 Tingting Zhao, Kristin Carroll, Caley Craven, Nicholas JP Wawryk, Shipei Xing, Jian Guo, Xing-Fang Li, Tao Huan. HDPairFinder: A data processing platform for hydrogen/deuterium isotopic labeling-based nontargeted analysis of trace-level amino-containing chemicals in environmental water. *Journal of Environmental Sciences*. ([10.1016/j.jes.2023.02.033](https://doi.org/10.1016/j.jes.2023.02.033))
- 2022 Daniel GC Treen, Mingxun Wang, Shipei Xing, Katherine B Louie, Tao Huan, Pieter C Dorrestein, Trent R Northen, Benjamin P Bowen. SIMILE enables alignment of tandem mass spectra with statistical significance. *Nature Communications*. ([10.1038/s41467-022-30118-9](https://doi.org/10.1038/s41467-022-30118-9))
- 2022 Jian Guo, Huaxu Yu, Shipei Xing, Tao Huan. Addressing big data challenges in mass spectrometry-based metabolomics. *Chemical Communications*. ([10.1039/D2CC03598G](https://doi.org/10.1039/D2CC03598G))
- 2021 Fanrong Zhao, Li Li, Yue Chen, Yichao Huang, Tharushi Prabha Keerthisinghe, Agnes Chow, Ting Dong, Shenglan Jia, Shipei Xing, Benedikt Warth, Tao Huan, Mingliang Fang. Risk-based chemical ranking and generating a prioritized human exposome database. *Environmental Health Perspectives*. ([10.1289/EHP7722](https://doi.org/10.1289/EHP7722))
- 2021 Jian Guo, Sam Shen, Shipei Xing, Ying Chen, Frank Chen, Elizabeth M Porter, Huaxu Yu, Tao Huan. EVA: evaluation of metabolic feature fidelity using a deep learning model trained with over 25000 extracted ion chromatograms. *Analytical Chemistry*. ([10.1021/acs.analchem.1c01309](https://doi.org/10.1021/acs.analchem.1c01309))
- 2021 Jian Guo, Sam Shen, Shipei Xing, Huaxu Yu, Tao Huan. ISFrag: de novo recognition of in-source fragments for liquid chromatography–mass spectrometry data. *Analytical Chemistry*. ([10.1021/acs.analchem.1c01644](https://doi.org/10.1021/acs.analchem.1c01644))
- 2021 Jian Guo, Sam Shen, Shipei Xing, Tao Huan. DaDIA: hybridizing data-dependent and data-independent acquisition modes for generating high-quality metabolomic data. *Analytical Chemistry*. ([10.1021/acs.analchem.0c05022](https://doi.org/10.1021/acs.analchem.0c05022))
- 2021 Polina Beskrovnaya, Doaa Fakih, Isabelle Morneau, Ameena Hashimi, Dainelys Guadarrama Bello, Shipei Xing, Antonio Nanci, Tao Huan, Elitza I Tocheva. No Endospore Formation Confirmed in Members of the Phylum Proteobacteria. *Applied and Environmental Microbiology*. ([10.1128/AEM.02312-20](https://doi.org/10.1128/AEM.02312-20))
- 2020 Huaxu Yu, Shipei Xing, Lorenz Nierves, Philipp F Lange, Tao Huan. Fold-change compression: an unexplored but correctable quantitative bias caused by nonlinear electrospray ionization responses in untargeted metabolomics. *Analytical Chemistry*. ([10.1021/acs.analchem.0c00246](https://doi.org/10.1021/acs.analchem.0c00246))

## Patents

- 2022 "Bottom-Up" Annotation of Molecular Formula Through Tandem Mass Spectra  
Shipei Xing, Tao Huan.  
*Invention ID: 2022-106. Technology ID: 23-062.*
- 2020 Core structure-based tandem MS spectral similarity algorithm  
Shipei Xing, Tao Huan.  
*Invention ID: 2020-050. Technology ID: 20-178.*

## Honors and Awards

- 2022 Canadian Society of Mass Spectrometry (CSMS) Lake Louise Travel Award (sponsored by Thermo).
- 2022 – 2023 Affiliated Fellowship, UBC.
- 2022 Pei-Huang Tung and Tan-Wen Tung Graduate Fellowship, UBC.
- 2022 Gladys Estella Laird Research Fellowship, UBC.
- 2022 C L Wang Memorial Scholarship, UBC.
- 2022 Dr. Arnold By Travel Fellowship, UBC Chemistry.
- 2022 Graduate Student Travel Award, UBC.
- 2021 Sandra Morris and Richard Tillyer Scholarship in Chemistry, UBC Chemistry.
- 2021 Best Poster Award. 1st Chinese American Society for Mass Spectrometry (CASMS) Virtual Conference.
- 2020 – 2022 President's Academic Excellence Initiative PhD Award, UBC.
- 2019 – 2022 Faculty of Science PhD Tuition Award, UBC.
- 2019 – 2022 International Tuition Award, UBC.
- 2018 Outstanding Thesis Award, ZJU.

## Conference Presentations

- 2024 **Poster presentation.** Discovering and annotating new molecules in untargeted metabolomics through structural coupling.  
*72<sup>nd</sup> ASMS Conference on Mass Spectrometry and Allied Topics. Anaheim, US.*
- 2022 **Oral presentation.** Molecular formula discovery via bottom-up MS/MS interrogation.  
*34<sup>th</sup> Lake Louise Tandem MS Workshop. Lake Louise, Canada.*
- 2022 **Lightning talk & poster presentation.** BUDDY: Bottom-up MS/MS interrogation enables large-scale discovery of unreported molecular formulae with significance control.  
*2<sup>nd</sup> CASMS Virtual Conference.*
- 2022 **Poster presentation.** Radical Fragment Ions in Collision-Induced Dissociation-based Tandem Mass Spectrometry.  
*70<sup>th</sup> ASMS Conference on Mass Spectrometry and Allied Topics. Minneapolis, US.*

- 2021    **Lightning talk & poster presentation.** Radical Fragment Ions in Collision-Induced Dissociation Mass Spectrometry.  
*1<sup>st</sup> CASMS Virtual Conference.*

## Service

### Reviewer

- Co-reviewer: Nature, Cell, Nature Microbiology
- Independent reviewer: Nature Communications, BMC Bioinformatics, Journal of Cheminformatics, Chemosphere, Metabolites, etc.

### Editorial Board Member

- BMC Chemistry

### Guest Editor

- Metabolites – Special Issue on "Metabolomics and Bioinformatics Approaches to Studying Human Gut Microbiota-Derived Metabolites"

## Teaching Experience

- 2022    Teaching assistant (lecture), CHEM 211: Analytical Chemistry. **UBC**.
- 2020    Teaching assistant (lab), CHEM 154: Chemistry for Engineering. **UBC**.
- 2019 & 2021    Teaching assistant (lab), CHEM 123: Thermodynamics, Kinetics and Organic Chemistry. **UBC**.

## Professional Affiliations

- 2022 – Present    Canadian Society for Mass Spectrometry (CSMS), Member.
- 2021 – Present    Chinese American Society for Mass Spectrometry (CASMS), Member.
- 2020 – Present    American Society for Mass Spectrometry (ASMS), Member.