

Shipei Xing

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Research Interests

Metabolomics, Mass spectrometry, Bioinformatics, Analytical chemistry

Employment

2023.8 – Present **University 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 (**Qiusi 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.](#)

- 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)

- 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))

Co-Authored Publications

- 2024 Abzer K Pakkir Shah, Axel Walter, Filip Ottosson, Francesco Russo, Marcelo Navarro-Díaz, Judith Boldt, Jarmo-Charles Kalinski, Eftychia E Kontou, James Elofson, Alexandros Polyzois, Carolina González-Marín, Shane Farrell, Marie R Aggerbeck, Thapanee Pruksatrakul, Nathan Chan, Yunshu Wang, Magdalena Pöchlacker, Corinna Brungs, Beatriz Cámara, Andrés M 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 Beemelmans, Hannes Link, Christoph Mayer, Justin JJ van der Hooft, Tito Damiani, Tomáš Pluskal, Pieter C Dorrestein, Jan Stanstrup, Robin Schmid, Mingxun Wang, Allegra T Aron, Madeleine Ernst, Daniel Petras. The Hitchhiker’s Guide to Statistical Analysis of Feature-based Molecular Networks from Non-Targeted Metabolomics Data. *Nature Protocols*. ([just accepted](#))
- 2024 Ipsita Mohanty, Helena Mannocho-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 Morais, 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 Aksenov, 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 Mannocho-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 Kirstin L Brown, Jessica MC Krekhno, Shipei Xing, Tao Huan, Lindsay D Eltis. Cholesterol-Mediated Coenzyme A Depletion in Catabolic Mutants of Mycobacteria Leads to Toxicity. *ACS Infectious Diseases*. ([10.1021/acsinfecdis.3c00237](https://doi.org/10.1021/acsinfecdis.3c00237))

- 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)
- 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)
- 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)
- 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)
- 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)
- 2022 Jian Guo, Huaxu Yu, Shipei Xing, Tao Huan. Addressing big data challenges in mass spectrometry-based metabolomics. *Chemical Communications*. (10.1039/D2CC03598G)
- 2022 Jian Guo, Sam Shen, Min Liu, Chenjingyi Wang, Brian Low, Ying Chen, Yaxi Hu, Shipei Xing, Huaxu Yu, Yu Gao, Mingliang Fang, Tao Huan. JPA: joint metabolic feature extraction increases the depth of chemical coverage for LC-MS-Based metabolomics and exposomics. *Metabolites*. (10.3390/metabo12030212)
- 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)
- 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)
- 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)
- 2021 Ying Chen, Jian Guo, Shipei Xing, Huaxu Yu, Tao Huan. Global-scale metabolomic profiling of human hair for simultaneous monitoring of endogenous metabolome, short-and long-term exposome. *Frontiers in Chemistry*. (10.3389/fchem.2021.674265)
- 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)

- 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)
- 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)

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.

Presentations

- 2024 **Poster presentation.** Discovering and annotating new molecules in untargeted metabolomics through structural coupling.
72nd ASMS Conference on Mass Spectrometry and Allied Topics. Anaheim, US.
- 2022 **Oral presentation.** Molecular formula discovery via bottom-up MS/MS interrogation.
34th 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.
2nd CASMS Virtual Conference.
- 2022 **Poster presentation.** Radical Fragment Ions in Collision-Induced Dissociation-based Tandem Mass Spectrometry.
70th ASMS Conference on Mass Spectrometry and Allied Topics. Minneapolis, US.
- 2021 **Lightning talk & poster presentation.** Radical Fragment Ions in Collision-Induced Dissociation Mass Spectrometry.
1st CASMS Virtual Conference.

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.

Service

Reviewer: Nature Microbiology, BMC Bioinformatics

Language Competency

English (fluent); Mandarin (native).