

# Xinjie Zhao

## List of Publications by Year in descending order

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54  
papers

4,002  
citations

172207

29  
h-index

161609

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all docs

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docs citations

57  
times ranked

7093  
citing authors

#	ARTICLE	IF	CITATIONS
1	Gut microbiome and serum metabolome alterations in obesity and after weight-loss intervention. <i>Nature Medicine</i> , 2017, 23, 859-868.	15.2	1,074
2	Analyses of gut microbiota and plasma bile acids enable stratification of patients for antidiabetic treatment. <i>Nature Communications</i> , 2017, 8, 1785.	5.8	312
3	Relationship of Serum Trimethylamine N-Oxide (TMAO) Levels with early Atherosclerosis in Humans. <i>Scientific Reports</i> , 2016, 6, 26745.	1.6	224
4	Gut microbiome-related effects of berberine and probiotics on type 2 diabetes (the PREMOTÉ study). <i>Nature Communications</i> , 2020, 11, 5015.	5.8	184
5	Discovery and Validation of Plasma Biomarkers for Major Depressive Disorder Classification Based on Liquid Chromatography–Mass Spectrometry. <i>Journal of Proteome Research</i> , 2015, 14, 2322-2330.	1.8	152
6	Integration of Metabolomics and Transcriptomics Reveals Major Metabolic Pathways and Potential Biomarker Involved in Prostate Cancer. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 154-163.	2.5	149
7	Changes of the plasma metabolome during an oral glucose tolerance test: is there more than glucose to look at?. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2009, 296, E384-E393.	1.8	143
8	Development of a High Coverage Pseudotargeted Lipidomics Method Based on Ultra-High Performance Liquid Chromatography–Mass Spectrometry. <i>Analytical Chemistry</i> , 2018, 90, 7608-7616.	3.2	138
9	Development of a plasma pseudotargeted metabolomics method based on ultra-high-performance liquid chromatography–mass spectrometry. <i>Nature Protocols</i> , 2020, 15, 2519-2537.	5.5	127
10	Integration of lipidomics and transcriptomics unravels aberrant lipid metabolism and defines cholesteryl oleate as potential biomarker of prostate cancer. <i>Scientific Reports</i> , 2016, 6, 20984.	1.6	103
11	Effect of a traditional Chinese medicine preparation Xindi soft capsule on rat model of acute blood stasis: A urinary metabolomics study based on liquid chromatography–mass spectrometry. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2008, 873, 151-158.	1.2	98
12	Plasma lipidomics reveals potential lipid markers of major depressive disorder. <i>Analytical and Bioanalytical Chemistry</i> , 2016, 408, 6497-6507.	1.9	95
13	Comprehensive Strategy to Construct In-House Database for Accurate and Batch Identification of Small Molecular Metabolites. <i>Analytical Chemistry</i> , 2018, 90, 7635-7643.	3.2	90
14	Serum Metabolomics Study of Polycystic Ovary Syndrome Based on Liquid Chromatography–Mass Spectrometry. <i>Journal of Proteome Research</i> , 2014, 13, 1101-1111.	1.8	78
15	A data preprocessing strategy for metabolomics to reduce the mask effect in data analysis. <i>Frontiers in Molecular Biosciences</i> , 2015, 2, 4.	1.6	78
16	Serum metabolic profiling study of lung cancer using ultra high performance liquid chromatography/quadrupole time-of-flight mass spectrometry. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2014, 966, 147-153.	1.2	70
17	Metabolomics and transcriptomics profiles reveal the dysregulation of the tricarboxylic acid cycle and related mechanisms in prostate cancer. <i>International Journal of Cancer</i> , 2018, 143, 396-407.	2.3	57
18	Deep Annotation of Hydroxycinnamic Acid Amides in Plants Based on Ultra-High-Performance Liquid Chromatography–High-Resolution Mass Spectrometry and Its In Silico Database. <i>Analytical Chemistry</i> , 2018, 90, 14321-14330.	3.2	54

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19	Urinary profiling investigation of metabolites with cis-diol structure from cancer patients based on UPLC-MS and HPLC-MS as well as multivariate statistical analysis. <i>Journal of Separation Science</i> , 2006, 29, 2444-2451.	1.3	50
20	Metabolomics Study of Roux-en-Y Gastric Bypass Surgery (RYGB) to Treat Type 2 Diabetes Patients Based on Ultraperformance Liquid Chromatography-Mass Spectrometry. <i>Journal of Proteome Research</i> , 2016, 15, 1288-1299.	1.8	48
21	Strategy for Comprehensive Identification of Acylcarnitines Based on Liquid Chromatography-High-Resolution Mass Spectrometry. <i>Analytical Chemistry</i> , 2018, 90, 5712-5718.	3.2	42
22	Serum lipid profiling of patients with chronic hepatitis B, cirrhosis, and hepatocellular carcinoma by ultra fast LC/IT-TOF MS. <i>Electrophoresis</i> , 2013, 34, 2848-2856.	1.3	41
23	Nontargeted Screening Method for Illegal Additives Based on Ultrahigh-Performance Liquid Chromatography-High-Resolution Mass Spectrometry. <i>Analytical Chemistry</i> , 2016, 88, 8870-8877.	3.2	41
24	Association of Serum Bile Acids Profile and Pathway Dysregulation With the Risk of Developing Diabetes Among Normoglycemic Chinese Adults: Findings From the 4C Study. <i>Diabetes Care</i> , 2021, 44, 499-510.	4.3	40
25	Lysophosphatidylcholines activate PPAR $\gamma$ and protect human skeletal muscle cells from lipotoxicity. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2016, 1861, 1980-1992.	1.2	38
26	Human Prostate Cancer Is Characterized by an Increase in Urea Cycle Metabolites. <i>Cancers</i> , 2020, 12, 1814.	1.7	37
27	A multi-omics investigation of the molecular characteristics and classification of six metabolic syndrome relevant diseases. <i>Theranostics</i> , 2020, 10, 2029-2046.	4.6	35
28	Pseudotargeted Method Based on Parallel Column Two-Dimensional Liquid Chromatography-Mass Spectrometry for Broad Coverage of Metabolome and Lipidome. <i>Analytical Chemistry</i> , 2020, 92, 6043-6050.	3.2	34
29	Ion-Pair Selection Method for Pseudotargeted Metabolomics Based on SWATH MS Acquisition and Its Application in Differential Metabolite Discovery of Type 2 Diabetes. <i>Analytical Chemistry</i> , 2018, 90, 11401-11408.	3.2	33
30	Metabolomics profiling of metformin-mediated metabolic reprogramming bypassing AMPK. <i>Metabolism: Clinical and Experimental</i> , 2019, 91, 18-29.	1.5	30
31	Muscle-Liver Substrate Fluxes in Exercising Humans and Potential Effects on Hepatic Metabolism. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, 1196-1209.	1.8	27
32	Rapid lipidomic profiling based on ultra-high performance liquid chromatography-mass spectrometry and its application in diabetic retinopathy. <i>Analytical and Bioanalytical Chemistry</i> , 2020, 412, 3585-3594.	1.9	27
33	Combined berberine and probiotic treatment as an effective regimen for improving postprandial hyperlipidemia in type 2 diabetes patients: a double blinded placebo controlled randomized study. <i>Gut Microbes</i> , 2022, 14, 2003176.	4.3	27
34	Serum Metabolomics Study of the Acute Graft Rejection in Human Renal Transplantation Based on Liquid Chromatography-Mass Spectrometry. <i>Journal of Proteome Research</i> , 2014, 13, 2659-2667.	1.8	25
35	GC/MS-based metabolomic studies reveal key roles of glycine in regulating silk synthesis in silkworm, <i>Bombyx mori</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2015, 57, 41-50.	1.2	24
36	Serum metabolomics study of Traditional Chinese medicine formula intervention to polycystic ovary syndrome. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2016, 120, 127-133.	1.4	24

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37	Metabolic Alterations Related to Glioma Grading Based on Metabolomics and Lipidomics Analyses. <i>Metabolites</i> , 2020, 10, 478.	1.3	18
38	Muscle and liver-specific alterations in lipid and acylcarnitine metabolism after a single bout of exercise in mice. <i>Scientific Reports</i> , 2016, 6, 22218.	1.6	17
39	An alignment algorithm for LC-MS-based metabolomics dataset assisted by MS/MS information. <i>Analytica Chimica Acta</i> , 2017, 990, 96-102.	2.6	17
40	Serum Metabolomics Study of Gliclazide-Modified-Release-Treated Type 2 Diabetes Mellitus Patients Using a Gas Chromatography–Mass Spectrometry Method. <i>Journal of Proteome Research</i> , 2018, 17, 1575-1585.	1.8	17
41	The delayed effects of antibiotics in type 2 diabetes, friend or foe?. <i>Journal of Endocrinology</i> , 2018, 238, 137-149.	1.2	15
42	Removal of false positive features to generate authentic peak table for high-resolution mass spectrometry-based metabolomics study. <i>Analytica Chimica Acta</i> , 2019, 1067, 79-87.	2.6	12
43	How to Screen and Prevent Metabolic Syndrome in Patients of PCOS Early: Implications From Metabolomics. <i>Frontiers in Endocrinology</i> , 2021, 12, 659268.	1.5	10
44	MetEx: A Targeted Extraction Strategy for Improving the Coverage and Accuracy of Metabolite Annotation in Liquid Chromatography–High-Resolution Mass Spectrometry Data. <i>Analytical Chemistry</i> , 2022, 94, 8561-8569.	3.2	10
45	Systematic, Modifying Group-Assisted Strategy Expanding Coverage of Metabolite Annotation in Liquid Chromatography–Mass Spectrometry-Based Nontargeted Metabolomics Studies. <i>Analytical Chemistry</i> , 2021, 93, 10916-10924.	3.2	8
46	A graph density-based strategy for features fusion from different peak extract software to achieve more metabolites in metabolic profiling from high-resolution mass spectrometry. <i>Analytica Chimica Acta</i> , 2020, 1139, 8-14.	2.6	5
47	Metabolite Triplet in Serum Improves the Diagnostic Accuracy of Prediabetes and Diabetes Screening. <i>Journal of Proteome Research</i> , 2021, 20, 1005-1014.	1.8	5
48	Association of plasma branched-chain amino acids with overweight: A Mendelian randomization analysis. <i>Obesity</i> , 2021, 29, 1708-1718.	1.5	4
49	Metabolomics biomarker analysis of threatened abortion in polycystic ovary syndrome: a clinical discovery study. <i>RSC Advances</i> , 2017, 7, 52923-52929.	1.7	3
50	A high throughput lipidomics method and its application in atrial fibrillation based on 96-well plate pretreatment and liquid chromatography-mass spectrometry. <i>Journal of Chromatography A</i> , 2021, 1651, 462271.	1.8	3
51	Comparison of the metabolome in urine prior and eight weeks after radical prostatectomy uncovers pathologic and molecular features of prostate cancer. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2021, 205, 114288.	1.4	3
52	Identification and regulation of the xenometabolite derivatives cis- and trans-3,4-methylene-heptanoylcarnitine in plasma and skeletal muscle of exercising humans. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2020, 318, E701-E709.	1.8	2
53	Serum Metabonomics Reveals Risk Factors in Different Periods of Cerebral Infarction in Humans. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 784288.	1.6	2
54	Liquid Chromatography-Mass Spectrometry of Biofluids and Extracts. <i>Methods in Molecular Biology</i> , 2015, 1277, 61-73.	0.4	1