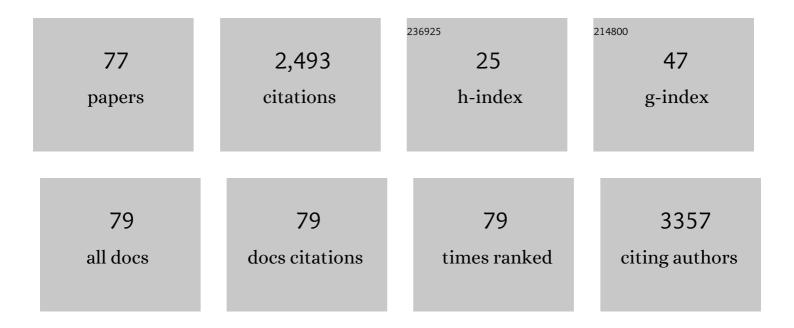
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular Pathological Diagnosis of Thyroid Tumors Using Spatially Resolved Metabolomics. Molecules, 2022, 27, 1390.	3.8	8
2	Contrast-enhanced CT-based radiomics model for differentiating risk subgroups of thymic epithelial tumors. BMC Medical Imaging, 2022, 22, 37.	2.7	4
3	Morus alba L. (Sangzhi) Alkaloids Promote Insulin Secretion, Restore Diabetic β-Cell Function by Preventing Dedifferentiation and Apoptosis. Frontiers in Pharmacology, 2022, 13, 841981.	3.5	12
4	Development and validation of a sensitive and reliable targeted metabolomics method for the quantification of cardiovascular diseaseâ€related biomarkers in plasma using ultrahighâ€performance liquid chromatography–tandem mass spectrometry. Rapid Communications in Mass Spectrometry, 2022, 36, e9292.	1.5	3
5	An Organ-Specific Metabolite Annotation Approach for Ambient Mass Spectrometry Imaging Reveals Spatial Metabolic Alterations of a Whole Mouse Body. Analytical Chemistry, 2022, 94, 7286-7294.	6.5	15
6	Norm ISWSVR: A Data Integration and Normalization Approach for Large-Scale Metabolomics. Analytical Chemistry, 2022, 94, 7500-7509.	6.5	4
7	Cryptotanshinone alleviates chemotherapy-induced colitis in mice with colon cancer via regulating fecal-bacteria-related lipid metabolism. Pharmacological Research, 2021, 163, 105232.	7.1	27
8	Rewiring of purine metabolism in response to acidosis stress in glioma stem cells. Cell Death and Disease, 2021, 12, 277.	6.3	18
9	Spatially resolved metabolomics combined with multicellular tumor spheroids to discover cancer tissue relevant metabolic signatures. Analytica Chimica Acta, 2021, 1155, 338342.	5.4	29
10	Mapping Metabolic Networks in the Brain by Ambient Mass Spectrometry Imaging and Metabolomics. Analytical Chemistry, 2021, 93, 6746-6754.	6.5	35
11	Hair growth predicts a depression-like phenotype in rats as a mirror of stress traceability. Neurochemistry International, 2021, 148, 105110.	3.8	1
12	Writing sequence identification of seals and signatures in documents using ambient mass spectrometry imaging with chemometric methods. Talanta, 2021, 235, 122804.	5.5	5
13	Enhanced On-Tissue Chemical Derivatization with Hydrogel Assistance for Mass Spectrometry Imaging. Analytical Chemistry, 2021, 93, 15373-15380.	6.5	17
14	Ratiometric Mass Spectrometry Imaging for Stain-Free Delineation of Ischemic Tissue and Spatial Profiling of Ischemia-Related Molecular Signatures. Frontiers in Chemistry, 2021, 9, 807868.	3.6	4
15	A targeted neurotransmitter quantification and nontargeted metabolic profiling method for pharmacometabolomics analysis of olanzapine by using UPLC-HRMS. RSC Advances, 2020, 10, 18305-18314.	3.6	6
16	Gegen Qinlian Decoction Coordinately Regulates PPARγ and PPARα to Improve Glucose and Lipid Homeostasis in Diabetic Rats and Insulin Resistance 3T3-L1 Adipocytes. Frontiers in Pharmacology, 2020, 11, 811.	3.5	18
17	Strategy for Global Profiling and Identification of 2- and 3-Hydroxy Fatty Acids in Plasma by UPLC–MS/MS. Analytical Chemistry, 2020, 92, 5143-5151.	6.5	19
18	Evaluation of the tumor-targeting efficiency and intratumor heterogeneity of anticancer drugs using quantitative mass spectrometry imaging. Theranostics, 2020, 10, 2621-2630.	10.0	37

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19	Development of a high-coverage metabolome relative quantitative method for large-scale sample analysis. Analytica Chimica Acta, 2020, 1109, 44-52.	5.4	8
20	A high-performance bio-tissue imaging method using air flow-assisted desorption electrospray ionization coupled with a high-resolution mass spectrometer. Chinese Chemical Letters, 2019, 30, 461-464.	9.0	12
21	Rational design of a multifunctional molecular dye for dual-modal NIR-II/photoacoustic imaging and photothermal therapy. Chemical Science, 2019, 10, 8348-8353.	7.4	137
22	A graphical data processing pipeline for mass spectrometry imaging-based spatially resolved metabolomics on tumor heterogeneity. Analytica Chimica Acta, 2019, 1077, 183-190.	5.4	26
23	Systematic optimization and evaluation of sample pretreatment methods for LC-MS-based metabolomics analysis of adherent mammalian cancer cells. Analytical Methods, 2019, 11, 3014-3022.	2.7	8
24	Development of methionine methylation profiling and relative quantification in human breast cancer cells based on metabolic stable isotope labeling. Analyst, The, 2019, 144, 3988-3998.	3.5	4
25	Whole-body spatially-resolved metabolomics method for profiling the metabolic differences of epimer drug candidates using ambient mass spectrometry imaging. Talanta, 2019, 202, 198-206.	5.5	14
26	Fabrication of homogenous threeâ€dimensional biomimetic tissue for mass spectrometry imaging. Journal of Mass Spectrometry, 2019, 54, 378-388.	1.6	5
27	Development of a metabolic pathway-based pseudo-targeted metabolomics method using liquid chromatography coupled with mass spectrometry. Talanta, 2019, 192, 160-168.	5.5	36
28	Spatially resolved metabolomics to discover tumor-associated metabolic alterations. Proceedings of the United States of America, 2019, 116, 52-57.	7.1	222
29	The impact of chronic environmental metal and benzene exposure on human urinary metabolome among Chinese children and the elderly population. Ecotoxicology and Environmental Safety, 2019, 169, 232-239.	6.0	20
30	Virtual Calibration Quantitative Mass Spectrometry Imaging for Accurately Mapping Analytes across Heterogenous Biotissue. Analytical Chemistry, 2019, 91, 2838-2846.	6.5	35
31	Derivatization reagent-assisted enantioseparation of 3-hydroxyaspartate with two chiral centers in rat cerebrospinal fluid by capillary electrophoresis-mass spectrometry. Analytica Chimica Acta, 2019, 1047, 257-266.	5.4	18
32	Rapid and sensitive liquid chromatography–tandem mass spectrometric method for the quantitative determination of potentially harmful substance 5,5′-oxydimethylenebis (2-furfural) in traditional Chinese medicine injections. Acta Pharmaceutica Sinica B, 2018, 8, 235-241.	12.0	8
33	A rapid and sensitive UPLC–MS/MS method for quantitative determination of arformoterol in rat plasma, lung and trachea tissues. Chinese Chemical Letters, 2018, 29, 1284-1286.	9.0	3
34	Biotransformation-based metabolomics profiling method for determining and quantitating cancer-related metabolites. Journal of Chromatography A, 2018, 1580, 80-89.	3.7	11
35	Development of simultaneous targeted metabolite quantification and untargeted metabolomics strategy using dual-column liquid chromatography coupled with tandem mass spectrometry. Analytica Chimica Acta, 2018, 1037, 369-379.	5.4	24
36	Combination of Droplet Extraction and Pico-ESI-MS Allows the Identification of Metabolites from Single Cancer Cells. Analytical Chemistry, 2018, 90, 9897-9903.	6.5	68

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37	Mass spectrometry imaging of intact cholesterol in a mouse esophagus tissue section and mouse zygotes using VUV laser desorption/ionization method. International Journal of Mass Spectrometry, 2018, 432, 9-13.	1.5	15
38	Systematic evaluation of serum and plasma collection on the endogenous metabolome. Bioanalysis, 2017, 9, 239-250.	1.5	24
39	Development of a Data-Independent Targeted Metabolomics Method for Relative Quantification Using Liquid Chromatography Coupled with Tandem Mass Spectrometry. Analytical Chemistry, 2017, 89, 6954-6962.	6.5	42
40	LC-MS-based metabolomics reveals metabolic signatures related to glioma stem-like cell self-renewal and differentiation. RSC Advances, 2017, 7, 24221-24232.	3.6	10
41	Methods used to increase the comprehensive coverage of urinary and plasma metabolomes by MS. Bioanalysis, 2016, 8, 981-997.	1.5	32
42	Global metabolomics reveals potential urinary biomarkers of esophageal squamous cell carcinoma for diagnosis and staging. Scientific Reports, 2016, 6, 35010.	3.3	32
43	Optimization and Evaluation Strategy of Esophageal Tissue Preparation Protocols for Metabolomics by LC–MS. Analytical Chemistry, 2016, 88, 3459-3464.	6.5	11
44	Ambient Mass Spectrometry Imaging Metabolomics Method Provides Novel Insights into the Action Mechanism of Drug Candidates. Analytical Chemistry, 2015, 87, 5372-5379.	6.5	68
45	Human Metabolic Responses to Chronic Environmental Polycyclic Aromatic Hydrocarbon Exposure by a Metabolomic Approach. Journal of Proteome Research, 2015, 14, 2583-2593.	3.7	69
46	Targeted Data-Independent Acquisition and Mining Strategy for Trace Drug Metabolite Identification Using Liquid Chromatography Coupled with Tandem Mass Spectrometry. Analytical Chemistry, 2015, 87, 7535-7539.	6.5	23
47	A sensitive and rapid HPLC–MS/MS method for the quantitative determination of trace amount of bromocriptine in small clinical prolactinoma tissue. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2015, 989, 91-97.	2.3	5
48	An integrated approach for detection and characterization of the trace impurities in levofloxacin using liquid chromatography-tandem mass spectrometry. Rapid Communications in Mass Spectrometry, 2014, 28, 1164-1174.	1.5	13
49	Combination of Injection Volume Calibration by Creatinine and MS Signals' Normalization to Overcome Urine Variability in LC-MS-Based Metabolomics Studies. Analytical Chemistry, 2013, 85, 7659-7665.	6.5	53
50	Plasma metabolome analysis by integrated ionization rapidâ€resolution liquid chromatography/tandem mass spectrometry. Rapid Communications in Mass Spectrometry, 2013, 27, 2071-2080.	1.5	21
51	Assessment of data pre-processing methods for LC-MS/MS-based metabolomics of uterine cervix cancer. Analyst, The, 2013, 138, 2669.	3.5	41
52	A rapid and sensitive liquid chromatography-tandem mass spectrometry method for the quantitation of S-phenylmercapturic acid in human urine. Analytical Methods, 2013, 5, 6081.	2.7	4
53	Air Flow-Assisted Ionization Imaging Mass Spectrometry Method for Easy Whole-Body Molecular Imaging under Ambient Conditions. Analytical Chemistry, 2013, 85, 2977-2982.	6.5	98
54	Liquid Chromatography–Tandem Mass Spectrometry-Based Plasma Metabonomics Delineate the Effect of Metabolites' Stability on Reliability of Potential Biomarkers. Analytical Chemistry, 2013, 85, 2606-2610.	6.5	63

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55	Global and Targeted Metabolomics of Esophageal Squamous Cell Carcinoma Discovers Potential Diagnostic and Therapeutic Biomarkers. Molecular and Cellular Proteomics, 2013, 12, 1306-1318.	3.8	113
56	Studies on the Interactions of Copper and Zinc Ions with β-Amyloid Peptides by a Surface Plasmon Resonance Biosensor. International Journal of Molecular Sciences, 2012, 13, 11832-11843.	4.1	12
57	Analysis of multiplex endogenous estrogen metabolites in human urine using ultra-fast liquid chromatography–tandem mass spectrometry: A case study for breast cancer. Analytica Chimica Acta, 2012, 711, 60-68.	5.4	42
58	Integrated rapid resolution liquid chromatography–tandem mass spectrometric approach for screening and identification of metabolites of the potential anticancer agent 3,6,7-trimethoxyphenanthroindolizidine in rat urine. Analytica Chimica Acta, 2012, 731, 60-67.	5.4	14
59	Hepatoprotective activities of a sesquiterpene-rich fraction from the aerial part of Cichorium glandulosum. Chinese Medicine, 2012, 7, 21.	4.0	13
60	Fast and Direct Analysis of Active Ingredient in Unknown Tablet Using Air Flow Assisted Ionization-Mass Spectrometry. Chinese Journal of Analytical Chemistry, 2012, 39, 1743-1747.	1.7	1
61	Time-Course Changes in Potential Biomarkers Detected Using a Metabonomic Approach in Walker 256 Tumor-Bearing Rats. Journal of Proteome Research, 2011, 10, 1953-1961.	3.7	26
62	Plasma Preparation Method for Metabolomic Analysis Based on Rapid Resolution Liquid Chromatography-Mass Spectrometry. Chinese Journal of Analytical Chemistry, 2011, 39, 1793-1797.	1.7	5
63	Air flow assisted ionization for remote sampling of ambient mass spectrometry and its application. Rapid Communications in Mass Spectrometry, 2011, 25, 843-850.	1.5	73
64	Simultaneous Structural Identification of Natural Products in Fractions of Crude Extract of the Rare Endangered Plant Anoectochilus roxburghii Using 1H NMR/RRLC-MS Parallel Dynamic Spectroscopy. International Journal of Molecular Sciences, 2011, 12, 2556-2571.	4.1	15
65	The characteristic fragmentation and rearrangement reaction of cationized glucopyranosyloxybenzyl tartrates by tandem mass spectrometry. Journal of Mass Spectrometry, 2010, 45, 824-828.	1.6	16
66	Integrated Ionization Approach for RRLCâ^'MS/MS-based Metabonomics: Finding Potential Biomarkers for Lung Cancer. Journal of Proteome Research, 2010, 9, 4071-4081.	3.7	97
67	Study of the characteristic fragmentation behavior of hydroquinone glycosides by electrospray ionization tandem mass spectrometry with optimization of collision energy. Journal of Mass Spectrometry, 2009, 44, 1182-1187.	1.6	19
68	Characterization of acid-induced protein conformational changes and noncovalent complexes in solution by using coldspray ionization mass spectrometry. Journal of the American Society for Mass Spectrometry, 2009, 20, 845-851.	2.8	14
69	An investigation of the fragmentation differences of isomeric flavonolâ€ <i>O</i> â€glycosides under different collisionâ€induced dissociation based mass spectrometry. Rapid Communications in Mass Spectrometry, 2009, 23, 1519-1524.	1.5	31
70	Nuclear magnetic resonance and liquid chromatography–mass spectrometry combined with an incompleted separation strategy for identifying the natural products in crude extract. Analytica Chimica Acta, 2009, 632, 221-228.	5.4	30
71	RRLC-MS/MS-based metabonomics combined with in-depth analysis of metabolic correlation network: finding potential biomarkers for breast cancer. Analyst, The, 2009, 134, 2003.	3.5	160
72	Development of a liquid chromatography/electrospray ionization tandem mass spectrometric method for the determination of hydroxyl radical. Rapid Communications in Mass Spectrometry, 2007, 21, 107-111.	1.5	10

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73	Fast profiling of the integral metabolism of flavonols in the active fraction ofGossypium herbaceam L. using liquid chromatography/multi-stage tandem mass spectrometry. Rapid Communications in Mass Spectrometry, 2007, 21, 1877-1888.	1.5	26
74	Direct On-Line Method To Monitor the Dynamic Structure of Noncovalent Titanium Complexes in Solution by Using Cold-Spray Ionization Time-of-Flight Mass Spectrometry. Analytical Chemistry, 2006, 78, 4737-4740.	6.5	22
75	Investigation of interconversion between aspacochiosides A and B by fast-atom bombardment mass spectrometry. Rapid Communications in Mass Spectrometry, 2006, 20, 328-330.	1.5	1
76	Structural characterization of flavonol 3,7-di-O-glycosides and determination of the glycosylation position by using negative ion electrospray ionization tandem mass spectrometry. Journal of Mass Spectrometry, 2006, 41, 352-360.	1.6	204
77	Characteristic elimination reactions of 1,2-disubstituted phenylbenzimidazoles and their isosteres 2,3-disubstituted phenylindoles in electron ionization mass spectrometry. Rapid Communications in Mass Spectrometry, 2004, 18, 584-587.	1.5	2