Guoxiang Xie

List of Publications by Year in descending order

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Version: 2024-02-01

41344 40979 10,078 137 49 93 citations h-index g-index papers 146 146 146 14313 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Bile acid–microbiota crosstalk in gastrointestinal inflammation and carcinogenesis. Nature Reviews Gastroenterology and Hepatology, 2018, 15, 111-128.	17.8	1,100
2	Theabrownin from Pu-erh tea attenuates hypercholesterolemia via modulation of gut microbiota and bile acid metabolism. Nature Communications, 2019, 10, 4971.	12.8	418
3	Altered bile acid profile associates with cognitive impairment in Alzheimer's disease—An emerging role for gut microbiome. Alzheimer's and Dementia, 2019, 15, 76-92.	0.8	396
4	The Footprints of Gut Microbial–Mammalian Co-Metabolism. Journal of Proteome Research, 2011, 10, 5512-5522.	3.7	268
5	Serum and Urine Metabolite Profiling Reveals Potential Biomarkers of Human Hepatocellular Carcinoma. Molecular and Cellular Proteomics, 2011, 10, M110.004945.	3.8	267
6	Altered Bile Acid Metabolome in Patients with Nonalcoholic Steatohepatitis. Digestive Diseases and Sciences, 2015, 60, 3318-3328.	2.3	251
7	A targeted metabolomic protocol for short-chain fatty acids and branched-chain amino acids. Metabolomics, 2013, 9, 818-827.	3.0	212
8	Bile acid is a significant host factor shaping the gut microbiome of diet-induced obese mice. BMC Biology, 2017, 15, 120.	3.8	208
9	Altered bile acid profile in mild cognitive impairment and Alzheimer's disease: Relationship to neuroimaging and CSF biomarkers. Alzheimer's and Dementia, 2019, 15, 232-244.	0.8	198
10	Salivary metabolite signatures of oral cancer and leukoplakia. International Journal of Cancer, 2011, 129, 2207-2217.	5.1	185
11	Hyocholic acid species improve glucose homeostasis through a distinct TGR5 and FXR signaling mechanism. Cell Metabolism, 2021, 33, 791-803.e7.	16.2	185
12	Distinct Urinary Metabolic Profile of Human Colorectal Cancer. Journal of Proteome Research, 2012, 11, 1354-1363.	3.7	184
13	Distinctly altered gut microbiota in the progression of liver disease. Oncotarget, 2016, 7, 19355-19366.	1.8	180
14	Quercetin is equally or more effective than resveratrol in attenuating tumor necrosis factor-α–mediated inflammation and insulin resistance in primary human adipocytes. American Journal of Clinical Nutrition, 2010, 92, 1511-1521.	4.7	177
15	Enhanced Fructose Utilization Mediated by SLC2A5 Is a Unique Metabolic Feature of Acute Myeloid Leukemia with Therapeutic Potential. Cancer Cell, 2016, 30, 779-791.	16.8	176
16	Dysregulated hepatic bile acids collaboratively promote liver carcinogenesis. International Journal of Cancer, 2016, 139, 1764-1775.	5.1	169
17	Metabonomics Identifies Serum Metabolite Markers of Colorectal Cancer. Journal of Proteome Research, 2013, 12, 3000-3009.	3.7	163
18	Alteration of bile acid metabolism in the rat induced by chronic ethanol consumption. FASEB Journal, 2013, 27, 3583-3593.	0.5	162

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19	Ultraâ€performance LC/TOF MS analysis of medicinal <i>Panax</i> herbs for metabolomic research. Journal of Separation Science, 2008, 31, 1015-1026.	2.5	161
20	A Distinct Metabolic Signature of Human Colorectal Cancer with Prognostic Potential. Clinical Cancer Research, 2014, 20, 2136-2146.	7.0	141
21	Branched-chain and aromatic amino acid profiles and diabetes risk in Chinese populations. Scientific Reports, 2016, 6, 20594.	3.3	140
22	Melamine-Induced Renal Toxicity Is Mediated by the Gut Microbiota. Science Translational Medicine, 2013, 5, 172ra22.	12.4	129
23	Profiling of Serum Bile Acids in a Healthy Chinese Population Using UPLC–MS/MS. Journal of Proteome Research, 2015, 14, 850-859.	3.7	129
24	A dysregulated bile acid-gut microbiota axis contributes to obesity susceptibility. EBioMedicine, 2020, 55, 102766.	6.1	128
25	Metabonomics of Human Colorectal Cancer: New Approaches for Early Diagnosis and Biomarker Discovery. Journal of Proteome Research, 2014, 13, 3857-3870.	3.7	127
26	Novel personalized pathway-based metabolomics models reveal key metabolic pathways for breast cancer diagnosis. Genome Medicine, 2016, 8, 34.	8.2	122
27	High Throughput and Quantitative Measurement of Microbial Metabolome by Gas Chromatography/Mass Spectrometry Using Automated Alkyl Chloroformate Derivatization. Analytical Chemistry, 2017, 89, 5565-5577.	6.5	117
28	Chronic Ethanol Consumption Alters Mammalian Gastrointestinal Content Metabolites. Journal of Proteome Research, 2013, 12, 3297-3306.	3.7	116
29	A Metabolite Array Technology for Precision Medicine. Analytical Chemistry, 2021, 93, 5709-5717.	6.5	112
30	Characterization of Pu-erh Tea Using Chemical and Metabolic Profiling Approaches. Journal of Agricultural and Food Chemistry, 2009, 57, 3046-3054.	5.2	111
31	Circulating Unsaturated Fatty Acids Delineate the Metabolic Status of Obese Individuals. EBioMedicine, 2015, 2, 1513-1522.	6.1	110
32	Germline BAP1 mutations induce a Warburg effect. Cell Death and Differentiation, 2017, 24, 1694-1704.	11.2	105
33	Metabolite profiling of Panax notoginseng using UPLC–ESI-MS. Phytochemistry, 2008, 69, 2237-2244.	2.9	103
34	Insulin resistance and the metabolism of branched-chain amino acids. Frontiers of Medicine, 2013, 7, 53-59.	3.4	101
35	Urinary Metabolite Markers of Precocious Puberty. Molecular and Cellular Proteomics, 2012, 11, M111.011072.	3.8	91
36	Metabolic Substrates Exhibit Differential Effects on Functional Parameters of Mouse Sperm Capacitation1. Biology of Reproduction, 2012, 87, 75.	2.7	89

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37	Metabonomic Evaluation of Melamine-Induced Acute Renal Toxicity in Rats. Journal of Proteome Research, 2010, 9, 125-133.	3.7	87
38	Supplementation of Milled Chia Seeds Increases Plasma ALA and EPA in Postmenopausal Women. Plant Foods for Human Nutrition, 2012, 67, 105-110.	3.2	87
39	Serum Bile Acids Are Associated with Pathological Progression of Hepatitis B-Induced Cirrhosis. Journal of Proteome Research, 2016, 15, 1126-1134.	3.7	78
40	Sex-dependent effects on gut microbiota regulate hepatic carcinogenic outcomes. Scientific Reports, 2017, 7, 45232.	3.3	71
41	Gut microbiota remodeling reverses aging-associated inflammation and dysregulation of systemic bile acid homeostasis in mice sex-specifically. Gut Microbes, 2020, 11, 1450-1474.	9.8	71
42	Metabolomics in human type 2 diabetes research. Frontiers of Medicine, 2013, 7, 4-13.	3.4	70
43	Plasma Metabolite Biomarkers for the Detection of Pancreatic Cancer. Journal of Proteome Research, 2015, 14, 1195-1202.	3.7	70
44	High Fat Diet Feeding Exaggerates Perfluorooctanoic Acid-Induced Liver Injury in Mice via Modulating Multiple Metabolic Pathways. PLoS ONE, 2013, 8, e61409.	2.5	69
45	Hyocholic acid species as novel biomarkers for metabolic disorders. Nature Communications, 2021, 12, 1487.	12.8	66
46	Clinical prediction of HBV and HCV related hepatic fibrosis using machine learning. EBioMedicine, 2018, 35, 124-132.	6.1	65
47	Metabolic Fate of Tea Polyphenols in Humans. Journal of Proteome Research, 2012, 11, 3449-3457.	3.7	56
48	Transcriptomic and Metabonomic Profiling Reveal Synergistic Effects of Quercetin and Resveratrol Supplementation in High Fat Diet Fed Mice. Journal of Proteome Research, 2012, 11, 4961-4971.	3.7	54
49	Metabolomics approaches for characterizing metabolic interactions between host and its commensal microbes. Electrophoresis, 2013, 34, 2787-2798.	2.4	53
50	The Metabolite Profiles of the Obese Population Are Gender-Dependent. Journal of Proteome Research, 2014, 13, 4062-4073.	3.7	53
51	Distinct Plasma Bile Acid Profiles of Biliary Atresia and Neonatal Hepatitis Syndrome. Journal of Proteome Research, 2015, 14, 4844-4850.	3.7	52
52	Ursodeoxycholic acid accelerates bile acid enterohepatic circulation. British Journal of Pharmacology, 2019, 176, 2848-2863.	5.4	52
53	The Brain Metabolome of Male Rats across the Lifespan. Scientific Reports, 2016, 6, 24125.	3.3	51
54	Dysregulated bile acid signaling contributes to the neurological impairment in murine models of acute and chronic liver failure. EBioMedicine, 2018, 37, 294-306.	6.1	51

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55	Analysis of dencichine in Panax notoginseng by gas chromatography–mass spectrometry with ethyl chloroformate derivatization. Journal of Pharmaceutical and Biomedical Analysis, 2007, 43, 920-925.	2.8	49
56	Key Role for the 12-Hydroxy Group in the Negative Ion Fragmentation of Unconjugated C24 Bile Acids. Analytical Chemistry, 2016, 88, 7041-7048.	6.5	49
57	Comparison of Flavonoid Composition of Red Raspberries (Rubus idaeus L.) Grown in the Southern United States. Journal of Agricultural and Food Chemistry, 2012, 60, 5779-5786.	5.2	48
58	Herbal medicine Yinchenhaotang protects against \hat{l} ±-naphthylisothiocyanate-induced cholestasis in rats. Scientific Reports, 2017, 7, 4211.	3.3	48
59	Conjugated secondary 12α-hydroxylated bile acids promote liver fibrogenesis. EBioMedicine, 2021, 66, 103290.	6.1	47
60	An Investigation of an Acute Gastroenteritis Outbreak: Cronobacter sakazakii, a Potential Cause of Food-Borne Illness. Frontiers in Microbiology, 2018, 9, 2549.	3.5	45
61	Dietary fat sources differentially modulate intestinal barrier and hepatic inflammation in alcohol-induced liver injury in rats. American Journal of Physiology - Renal Physiology, 2013, 305, G919-G932.	3.4	44
62	Lowered circulating aspartate is a metabolic feature of human breast cancer. Oncotarget, 2015, 6, 33369-33381.	1.8	44
63	A panel of free fatty acid ratios to predict the development of metabolic abnormalities in healthy obese individuals. Scientific Reports, 2016, 6, 28418.	3.3	43
64	Metabolic Transformation of DMBA-Induced Carcinogenesis and Inhibitory Effect of Salvianolic Acid B and Breviscapine Treatment. Journal of Proteome Research, 2012, 11, 1302-1316.	3.7	41
65	Chenodeoxycholic Acid as a Potential Prognostic Marker for Roux-en-Y Gastric Bypass in Chinese Obese Patients. Journal of Clinical Endocrinology and Metabolism, 2015, 100, 4222-4230.	3.6	40
66	Urinary metabolic insights into host-gut microbial interactions in healthy and IBD children. World Journal of Gastroenterology, 2017, 23, 3643.	3.3	38
67	Waterborne manganese exposure alters plasma, brain, and liver metabolites accompanied by changes in stereotypic behaviors. Neurotoxicology and Teratology, 2012, 34, 27-36.	2.4	37
68	Towards Polypharmacokinetics: Pharmacokinetics of Multicomponent Drugs and Herbal Medicines Using a Metabolomics Approach. Evidence-based Complementary and Alternative Medicine, 2013, 2013, 1-12.	1.2	37
69	Factors affecting separation and detection of bile acids by liquid chromatography coupled with mass spectrometry in negative mode. Analytical and Bioanalytical Chemistry, 2017, 409, 5533-5545.	3.7	37
70	Metformin suppressed the proliferation of LoVo cells and induced a time-dependent metabolic and transcriptional alteration. Scientific Reports, 2015, 5, 17423.	3.3	36
71	IP4M: an integrated platform for mass spectrometry-based metabolomics data mining. BMC Bioinformatics, 2020, 21, 444.	2.6	35
72	Differential Effects of Grape Powder and Its Extract on Glucose Tolerance and Chronic Inflammation in High-Fat-Fed Obese Mice. Journal of Agricultural and Food Chemistry, 2012, 60, 12458-12468.	5.2	34

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73	A rapid ultraâ€performance liquid chromatography–electrospray Ionisation mass spectrometric method for the analysis of saponins in the adventitious roots of <i>Panax notoginseng</i> . Phytochemical Analysis, 2009, 20, 68-76.	2.4	33
74	Serum metabolite profiles are associated with the presence of advanced liver fibrosis in Chinese patients with chronic hepatitis B viral infection. BMC Medicine, 2020, 18, 144.	5.5	33
75	Very Low Carbohydrate Diet Significantly Alters the Serum Metabolic Profiles in Obese Subjects. Journal of Proteome Research, 2013, 12, 5801-5811.	3.7	32
76	Dietary Nicotinic Acid Supplementation Ameliorates Chronic Alcohol-Induced Fatty Liver in Rats. Alcoholism: Clinical and Experimental Research, 2014, 38, 1982-1992.	2.4	32
77	Serum Metabolite Signatures of Type 2 Diabetes Mellitus Complications. Journal of Proteome Research, 2015, 14, 447-456.	3.7	32
78	Strategy for an Association Study of the Intestinal Microbiome and Brain Metabolome Across the Lifespan of Rats. Analytical Chemistry, 2018, 90, 2475-2483.	6.5	32
79	Serum Metabolic Signatures of Fulminant Type 1 Diabetes. Journal of Proteome Research, 2012, 11, 4705-4711.	3.7	30
80	Metabonomic Variations Associated with AOM-Induced Precancerous Colorectal Lesions and Resveratrol Treatment. Journal of Proteome Research, 2012, 11, 3436-3448.	3.7	29
81	Serum stearic acid/palmitic acid ratio as a potential predictor of diabetes remission after Rouxâ€en‥ gastric bypass in obesity. FASEB Journal, 2017, 31, 1449-1460.	0.5	29
82	Polyâ€pharmacokinetic Study of a Multicomponent Herbal Medicine in Healthy Chinese Volunteers. Clinical Pharmacology and Therapeutics, 2018, 103, 692-702.	4.7	29
83	Association between Pre-Diagnostic Serum Bile Acids and Hepatocellular Carcinoma: The Singapore Chinese Health Study. Cancers, 2021, 13, 2648.	3.7	29
84	Simultaneous determination of saponins in flower buds of Panax notoginseng using high performance liquid chromatography. Biomedical Chromatography, 2008, 22, 244-249.	1.7	28
85	Urinary Metabolite Variation Is Associated with Pathological Progression of the Post-Hepatitis B Cirrhosis Patients. Journal of Proteome Research, 2012, 11, 3838-3847.	3.7	28
86	Analysis of human C24 bile acids metabolome in serum and urine based on enzyme digestion of conjugated bile acids and LC-MS determination of unconjugated bile acids. Analytical and Bioanalytical Chemistry, 2018, 410, 5287-5300.	3.7	28
87	Toward Personalized Nutrition: Comprehensive Phytoprofiling and Metabotyping. Journal of Proteome Research, 2013, 12, 1547-1559.	3.7	27
88	Long-term Proton Pump Inhibitor Administration Caused Physiological and Microbiota Changes in Rats. Scientific Reports, 2020, 10, 866.	3.3	27
89	Bile Acid Profiles Are Distinct among Patients with Different Etiologies of Chronic Liver Disease. Journal of Proteome Research, 2021, 20, 2340-2351.	3.7	27
90	Metabonomic Profiling Reveals Cancer Chemopreventive Effects of American Ginseng on Colon Carcinogenesis in <i>Apc</i> ^{<i>Min/+</i>} Mice. Journal of Proteome Research, 2015, 14, 3336-3347.	3.7	26

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91	Metabolic profiling reveals therapeutic effects of Herba Cistanches in an animal model of hydrocortisone-induced 'kidney-deficiency syndrome'. Chinese Medicine, 2008, 3, 3.	4.0	25
92	Metabolomics analysis reveals variation in <i>Schisandra chinensis</i> metabolites from different origins. Journal of Separation Science, 2014, 37, 731-737.	2.5	25
93	Metabonomic Profiling of Human Placentas Reveals Different Metabolic Patterns among Subtypes of Neural Tube Defects. Journal of Proteome Research, 2014, 13, 934-945.	3.7	25
94	Peripheral serum metabolomic profiles inform central cognitive impairment. Scientific Reports, 2020, 10, 14059.	3.3	25
95	The ratio of dihomoâ€Î³â€linolenic acid to deoxycholic acid species is a potential biomarker for the metabolic abnormalities in obesity. FASEB Journal, 2017, 31, 3904-3912.	0.5	24
96	Distinct Metabolic Signature of Human Bladder Cancer Cells Carrying an Impaired Fanconi Anemia Tumor-Suppressor Signaling Pathway. Journal of Proteome Research, 2016, 15, 1333-1341.	3.7	23
97	Fingerprint Analysis of Flos Carthami by Pressurized CEC and LC. Chromatographia, 2006, 64, 739-743.	1.3	22
98	Fingerprint analysis of Rhizoma chuanxiong by pressurized capillary electrochromatography and high-performance liquid chromatography. Biomedical Chromatography, 2007, 21, 867-875.	1.7	22
99	Global and Targeted Metabolomics Evidence of the Protective Effect of Chinese Patent Medicine <i>Jinkui Shenqi</i> Pill on Adrenal Insufficiency after Acute Glucocorticoid Withdrawal in Rats. Journal of Proteome Research, 2016, 15, 2327-2336.	3.7	22
100	Blocking glycine utilization inhibits multiple myeloma progression by disrupting glutathione balance. Nature Communications, 2022, 13, .	12.8	21
101	Activation of PPARα-catalase pathway reverses alcoholic liver injury via upregulating NAD synthesis and accelerating alcohol clearance. Free Radical Biology and Medicine, 2021, 174, 249-263.	2.9	17
102	A new silymarin preparation based on solid dispersion technique. Advances in Therapy, 2005, 22, 595-600.	2.9	16
103	Analysis of urinary metabolites for metabolomic study by pressurized CEC. Electrophoresis, 2007, 28, 4459-4468.	2.4	16
104	Asymmetric dimethylarginine attenuates serum starvation-induced apoptosis via suppression of the Fas (APO-1/CD95)/JNK (SAPK) pathway. Cell Death and Disease, 2013, 4, e830-e830.	6.3	16
105	Effects of ADMA on gene expression and metabolism in serum-starved LoVo cells. Scientific Reports, 2016, 6, 25892.	3.3	16
106	Metabonomics Reveals Metabolite Changes in Biliary Atresia Infants. Journal of Proteome Research, 2015, 14, 2569-2574.	3.7	15
107	Altered bile acid glycineÂ:Âtaurine ratio in the progression of chronic liver disease. Journal of Gastroenterology and Hepatology (Australia), 2022, 37, 208-215.	2.8	15
108	Outbreak of Haff Disease caused by consumption of crayfish (Procambarus clarkii), Nanjing, Jiangsu Province, China. Food Control, 2016, 59, 690-694.	5.5	14

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109	Histamine is correlated with liver fibrosis in biliary atresia. Digestive and Liver Disease, 2016, 48, 921-926.	0.9	14
110	Metabolomics analysis delineates the therapeutic effects of Huangqi decoction and astragalosides on \hat{l}_{\pm} -naphthylisothiocyanate (ANIT) -induced cholestasis in rats. Journal of Ethnopharmacology, 2021, 268, 113658.	4.1	14
111	Association of dietary sodium:potassium ratio with the metabolic syndrome in Chinese adults. British Journal of Nutrition, 2018, 120, 612-618.	2.3	13
112	A Metabolomics-Based Strategy for the Quality Control of Traditional Chinese Medicine: Shengmai Injection as a Case Study. Evidence-based Complementary and Alternative Medicine, 2013, 2013, 1-8.	1.2	12
113	Urinary Metabolite Profiling Offers Potential for Differentiation of Liver-Kidney Yin Deficiency and Dampness-Heat Internal Smoldering Syndromes in Posthepatitis B Cirrhosis Patients. Evidence-based Complementary and Alternative Medicine, 2015, 2015, 1-11.	1.2	12
114	polyPK: an R package for pharmacokinetic analysis of multi-component drugs using a metabolomics approach. Bioinformatics, 2018, 34, 1792-1794.	4.1	12
115	Outbreak of Haff disease caused by consumption of crayfish (Procambarus clarkii) in nanjing, China. Clinical Toxicology, 2019, 57, 331-337.	1.9	12
116	The Metabolic Responses to Aerial Diffusion of Essential Oils. PLoS ONE, 2012, 7, e44830.	2. 5	12
117	Theabrownin and Poria cocos Polysaccharide Improve Lipid Metabolism via Modulation of Bile Acid and Fatty Acid Metabolism. Frontiers in Pharmacology, 0, 13, .	3.5	12
118	Serum and Urine Metabolite Profiling Reveals Potential Biomarkers of Human Hepatocellular Carcinoma*. Molecular and Cellular Proteomics, 2011, 10, A110.004945.	3.8	11
119	Differentiation of <i>Schisandra chinensis </i> and <i>Schisandra sphenanthera </i> using metabolite profiles based on UPLC-MS and GC-MS. Natural Product Research, 2012, 26, 255-263.	1.8	10
120	Assessing the Metabolic Effects of Aromatherapy in Human Volunteers. Evidence-based Complementary and Alternative Medicine, 2013, 2013, 1-9.	1.2	10
121	Fanconi Anemia complementation group C protein in metabolic disorders. Aging, 2018, 10, 1506-1522.	3.1	10
122	Metabolomic profiling in colorectal cancer: opportunities for personalized medicine. Personalized Medicine, 2013, 10, 741-755.	1.5	8
123	Phospholipids are A Potentially Important Source of Tissue Biomarkers for Hepatocellular Carcinoma: Results of a Pilot Study Involving Targeted Metabolomics. Diagnostics, 2019, 9, 167.	2.6	8
124	Diagnosis of Fibrosis Using Blood Markers and Logistic Regression in Southeast Asian Patients With Non-alcoholic Fatty Liver Disease. Frontiers in Medicine, 2021, 8, 637652.	2.6	8
125	Molecular typing of Shigella sonnei isolates circulating in Nanjing, China, 2007–2011. Journal of Infection in Developing Countries, 2014, 8, 1525-1532.	1.2	7
126	Probiotics, bile acids and gastrointestinal carcinogenesis. Nature Reviews Gastroenterology and Hepatology, 2018, 15, 205-205.	17.8	7

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127	Next generation sequencing for the investigation of an outbreak of Salmonella Schwarzengrund in Nanjing, China. International Journal of Biological Macromolecules, 2018, 107, 393-396.	7.5	7
128	Prepregnant Obesity of Mothers in a Multiethnic Cohort Is Associated with Cord Blood Metabolomic Changes in Offspring. Journal of Proteome Research, 2020, 19, 1361-1374.	3.7	7
129	Distinct Urinary Metabolic Biomarkers of Human Colorectal Cancer. Disease Markers, 2022, 2022, 1-18.	1.3	7
130	Combined Omics Reveals That Disruption of the Selenocysteine Lyase Gene Affects Amino Acid Pathways in Mice. Nutrients, 2019, 11, 2584.	4.1	6
131	Management of Hepatic Encephalopathy by Traditional Chinese Medicine. Evidence-based Complementary and Alternative Medicine, 2012, 2012, 1-8.	1.2	4
132	Genetic characterization of emergent GII.17 norovirus variants from 2013 to 2015 in Nanjing, China. Journal of Medical Microbiology, 2016, 65, 1274-1280.	1.8	4
133	3MCor: an integrative web server for metabolome–microbiome-metadata correlation analysis. Bioinformatics, 2022, 38, 1378-1384.	4.1	3
134	Metabonomics in Translational Research for Personalized Medicine and Nutrition. Molecular and Integrative Toxicology, 2015, , 63-82.	0.5	1
135	Identification of recombinant coxsackievirus A6 variants in hand, foot and mouth disease in Nanjing, China, 2013. Journal of Medical Microbiology, 2018, 67, 1120-1129.	1.8	1
136	Determination of Endogenous Metabolites in Obesity Rat Urine by Pressurized Capillary Electrochromatography with Ethyl Chloroformate Derivatization. Chinese Journal of Analytical Chemistry, 2007, 35, 1111-1115.	1.7	0
137	Studies on the in Vitro Dissolution of Insoluble Volatile Drug from Su-Anxin Nasal Inhalant and Its Correlation on the Nose Steady Self-Controllable Expiration and Inspiration at Night. Pharmacology & Pharmacy, 2011, 02, 67-72.	0.7	0