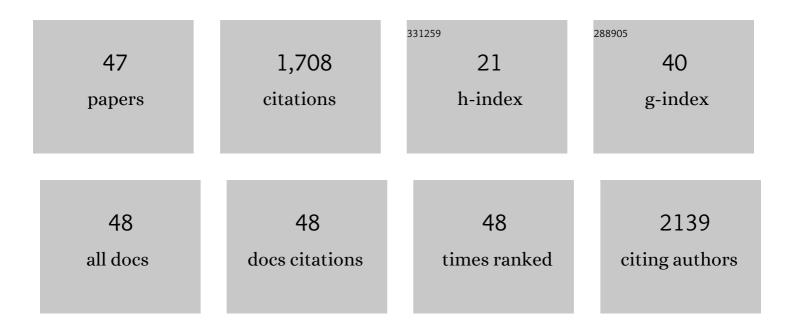
Nan Wang

List of Publications by Year in descending order

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NAN WANC

#	Article	IF	CITATIONS
1	A Sub-Nanostructural Transformable Nanozyme for Tumor Photocatalytic Therapy. Nano-Micro Letters, 2022, 14, 101.	14.4	24
2	Radiomics Analysis Based on Multiparametric <scp>MRI</scp> for Predicting Early Recurrence in Hepatocellular Carcinoma After Partial Hepatectomy. Journal of Magnetic Resonance Imaging, 2021, 53, 1066-1079.	1.9	49
3	High overage Metabolome Analysis Reveals Significant Diet Effects of Autoclaved and Irradiated Feed on Mouse Fecal and Urine Metabolomics. Molecular Nutrition and Food Research, 2021, 65, 2100110.	1.5	4
4	Association Between Antihypertensive Medication Use and Breast Cancer: A Systematic Review and Meta-Analysis. Frontiers in Pharmacology, 2021, 12, 609901.	1.6	12
5	Dynamic nanoassemblies for imaging and therapy of neurological disorders. Advanced Drug Delivery Reviews, 2021, 175, 113832.	6.6	15
6	Impact of Target Turnover on the Translation of Drug-Target Residence Time to Time-Dependent Antibacterial Activity. ACS Infectious Diseases, 2021, 7, 2755-2763.	1.8	5
7	Metabolomic profile overlap in prototypical autoimmune humoral disease: a comparison of myasthenia gravis and rheumatoid arthritis. Metabolomics, 2020, 16, 10.	1.4	25
8	Molecular Design of Conjugated Small Molecule Nanoparticles for Synergistically Enhanced PTT/PDT. Nano-Micro Letters, 2020, 12, 147.	14.4	82
9	Efficacy and Safety of Thalidomide for Chemotherapy-induced Nausea and Vomiting. Journal of Cancer, 2020, 11, 4560-4570.	1.2	6
10	Association of vitamin C intake with breast cancer risk and mortality: a meta-analysis of observational studies. Aging, 2020, 12, 18415-18435.	1.4	15
11	Beyond the antibodies: serum metabolomic profiling of myasthenia gravis. Metabolomics, 2019, 15, 109.	1.4	11
12	First-trimester metabolomic prediction of stillbirth. Journal of Maternal-Fetal and Neonatal Medicine, 2019, 32, 3435-3441.	0.7	5
13	Chemical Isotope Labeling LC-MS for Monitoring Disease Progression and Treatment in Animal Models: Plasma Metabolomics Study of Osteoarthritis Rat Model. Scientific Reports, 2017, 7, 40543.	1.6	18
14	Metabolomic analysis of oxidative stress: Superoxide dismutase mutation and paraquat induced stress in Drosophila melanogaster. Free Radical Biology and Medicine, 2017, 113, 323-334.	1.3	21
15	Elevated acetylâ€CoA by amino acid recycling fuels microalgal neutral lipid accumulation in exponential growth phase for biofuel production. Plant Biotechnology Journal, 2017, 15, 497-509.	4.1	36
16	The Impact of GFP Reporter Gene Transduction and Expression on Metabolomics of Placental Mesenchymal Stem Cells Determined by UHPLC-Q/TOF-MS. Stem Cells International, 2017, 2017, 1-12.	1.2	7
17	Cognitive Enhancement in Infants Associated with Increased Maternal Fruit Intake During Pregnancy: Results from a Birth Cohort Study with Validation in an Animal Model. EBioMedicine, 2016, 8, 331-340.	2.7	19
18	Dansylation isotope labeling liquid chromatography mass spectrometry for parallel profiling of human urinary and fecal submetabolomes. Analytica Chimica Acta, 2016, 903, 100-109.	2.6	29

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19	Development of High-Performance Chemical Isotope Labeling LC–MS for Profiling the Human Fecal Metabolome. Analytical Chemistry, 2015, 87, 829-836.	3.2	66
20	Effects of thyroxin and donepezil on hippocampal acetylcholine content and syntaxin-1 and munc-18 expression in adult rats with hypothyroidism. Experimental and Therapeutic Medicine, 2014, 7, 529-536.	0.8	7
21	Microbore liquid chromatography ultraviolet detection for quantification of total peptide amount and its application for assessing sample quality in shotgun proteome analysis of hundreds of cells. Journal of Chromatography A, 2014, 1338, 51-57.	1.8	1
22	Microwave-assisted acid hydrolysis of proteins combined with peptide fractionation and mass spectrometry analysis for characterizing protein terminal sequences. Journal of Proteomics, 2014, 100, 68-78.	1.2	6
23	In-Gel Microwave-Assisted Acid Hydrolysis of Proteins Combined with Liquid Chromatography Tandem Mass Spectrometry for Mapping Protein Sequences. Analytical Chemistry, 2014, 86, 600-607.	3.2	5
24	Development of microwave-assisted acid hydrolysis of proteins using a commercial microwave reactor and its combination with LC–MS for protein full-sequence analysis. Talanta, 2014, 129, 290-295.	2.9	9
25	Indigenous microbes survive in situ ozonation improving biodegradation of dissolved organic matter in aged oil sands process-affected waters. Chemosphere, 2013, 93, 2748-2755.	4.2	18
26	Impact of Ozonation on Naphthenic Acids Speciation and Toxicity of Oil Sands Process-Affected Water to <i>Vibrio fischeri</i> and Mammalian Immune System. Environmental Science & Technology, 2013, 47, 6518-6526.	4.6	111
27	Granule matrix property and rapid "kiss-and-run―exocytosis contribute to the different kinetics of catecholamine release from carotid glomus and adrenal chromaffin cells at matched quantal size. Canadian Journal of Physiology and Pharmacology, 2012, 90, 791-801.	0.7	4
28	Toxicity of untreated and ozone-treated oil sands process-affected water (OSPW) to early life stages of the fathead minnow (Pimephales promelas). Water Research, 2012, 46, 6359-6368.	5.3	128
29	Transcriptional Responses of the Brain–Gonad–Liver Axis of Fathead Minnows Exposed to Untreated and Ozone-Treated Oil Sands Process-Affected Water. Environmental Science & Technology, 2012, 46, 9701-9708.	4.6	68
30	Integrated SDS Removal and Peptide Separation by Strong-Cation Exchange Liquid Chromatography for SDS-Assisted Shotgun Proteome Analysis. Journal of Proteome Research, 2012, 11, 818-828.	1.8	34
31	Effect of Ozonation on the Estrogenicity and Androgenicity of Oil Sands Process-Affected Water. Environmental Science & Technology, 2011, 45, 6268-6274.	4.6	77
32	Naphthenic acids speciation and removal during petroleum-coke adsorption and ozonation of oil sands process-affected water. Science of the Total Environment, 2011, 409, 5119-5125.	3.9	143
33	Comparison of surfactant-assisted shotgun methods using acid-labile surfactants and sodium dodecyl sulfate for membrane proteome analysis. Analytica Chimica Acta, 2011, 698, 36-43.	2.6	41
34	Reproducible microwave-assisted acid hydrolysis of proteins using a household microwave oven and its combination with LC-ESI MS/MS for mapping protein sequences and modifications. Journal of the American Society for Mass Spectrometry, 2010, 21, 1573-1587.	1.2	30
35	Influence of Cholesterol on Catecholamine Release from the Fusion Pore of Large Dense Core Chromaffin Granules. Journal of Neuroscience, 2010, 30, 3904-3911.	1.7	45
36	Development of Mass Spectrometry-Based Shotgun Method for Proteome Analysis of 500 to 5000 Cancer Cells. Analytical Chemistry, 2010, 82, 2262-2271.	3.2	76

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37	Large-Scale Proteome Profile of the Zebrafish (<i>Danio rerio</i>) Gill for Physiological and Biomarker Discovery Studies. Zebrafish, 2009, 6, 229-238.	0.5	45
38	Off-Line Two-Dimensional Liquid Chromatography with Maximized Sample Loading to Reversed-Phase Liquid Chromatography-Electrospray Ionization Tandem Mass Spectrometry for Shotgun Proteome Analysis. Analytical Chemistry, 2009, 81, 1049-1060.	3.2	54
39	Identification of Arsenic-Binding Proteins in Human Cells by Affinity Chromatography and Mass Spectrometry. Analytical Chemistry, 2009, 81, 4144-4152.	3.2	53
40	Physicoâ€Chemical Processes. Water Environment Research, 2009, 81, 1056-1126.	1.3	2
41	Liquid Chromatography MALDI MS/MS for Membrane Proteome Analysis. Methods in Molecular Biology, 2009, 528, 295-310.	0.4	1
42	Proteome Profile of Human Breast Cancer Tissue Generated by LCâ^'ESIâ^'MS/MS Combined with Sequential Protein Precipitation and Solubilization. Journal of Proteome Research, 2008, 7, 3583-3590.	1.8	13
43	Absolute Threshold. , 2008, , 3-3.		0
44	Exploring the Precursor Ion Exclusion Feature of Liquid Chromatographyâ^'Electrospray Ionization Quadrupole Time-of-Flight Mass Spectrometry for Improving Protein Identification in Shotgun Proteome Analysis. Analytical Chemistry, 2008, 80, 4696-4710.	3.2	75
45	Proteome Profile of Cytosolic Component of Zebrafish Liver Generated by LCâ [~] ESI MS/MS Combined with Trypsin Digestion and Microwave-Assisted Acid Hydrolysis. Journal of Proteome Research, 2007, 6, 263-272.	1.8	69
46	Influence of Quantal Size and cAMP on the Kinetics of Quantal Catecholamine Release from Rat Chromaffin Cells. Biophysical Journal, 2007, 92, 2735-2746.	0.2	15
47	Characterization of Human Tear Proteome Using Multiple Proteomic Analysis Techniques. Journal of Proteome Research, 2005, 4, 2052-2061	1.8	129