## Huitao Zeng

List of Publications by Year in descending order

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Version: 2024-02-01

393982 360668 1,361 49 19 35 citations h-index g-index papers 50 50 50 1742 citing authors docs citations times ranked all docs

#	Article	IF	Citations
1	Deep-Learning-Based Drug–Target Interaction Prediction. Journal of Proteome Research, 2017, 16, 1401-1409.	1.8	381
2	Deep learning-based component identification for the Raman spectra of mixtures. Analyst, The, 2019, 144, 1789-1798.	1.7	130
3	DeepMirTar: a deep-learning approach for predicting human miRNA targets. Bioinformatics, 2018, 34, 3781-3787.	1.8	65
4	Prediction of Liquid Chromatographic Retention Time with Graph Neural Networks to Assist in Small Molecule Identification. Analytical Chemistry, 2021, 93, 2200-2206.	3.2	60
5	Predicting a Molecular Fingerprint from an Electron Ionization Mass Spectrum with Deep Neural Networks. Analytical Chemistry, 2020, 92, 8649-8653.	3.2	59
6	Deep MS/MS-Aided Structural-Similarity Scoring for Unknown Metabolite Identification. Analytical Chemistry, 2019, 91, 5629-5637.	3.2	47
7	Synthesis of Multi-Au-Nanoparticle-Embedded Mesoporous Silica Microspheres as Self-Filtering and Reusable Substrates for SERS Detection. ACS Applied Materials & Samp; Interfaces, 2017, 9, 42156-42166.	4.0	44
8	Comprehensive metabolic profiles of seminal plasma with different forms of male infertility and their correlation with sperm parameters. Journal of Pharmaceutical and Biomedical Analysis, 2020, 177, 112888.	1.4	39
9	Baseline correction of high resolution spectral profile data based on exponential smoothing. Chemometrics and Intelligent Laboratory Systems, 2014, 139, 97-108.	1.8	37
10	Classification of Green and Black Teas by PCA and SVM Analysis of Cyclic Voltammetric Signals from Metallic Oxide-Modified Electrode. Food Analytical Methods, 2014, 7, 472-480.	1.3	36
11	Selective iteratively reweighted quantile regression for baseline correction. Analytical and Bioanalytical Chemistry, 2014, 406, 1985-1998.	1.9	26
12	Absolute quantitative imaging of sphingolipids in brain tissue by exhaustive liquid microjunction surface sampling–liquid chromatography–mass spectrometry. Journal of Chromatography A, 2020, 1609, 460436.	1.8	26
13	In situ fabrication of label-free optical sensing paper strips for the rapid surface-enhanced Raman scattering (SERS) detection of brassinosteroids in plant tissues. Talanta, 2017, 165, 313-320.	2.9	25
14	KPIC2: An Effective Framework for Mass Spectrometry-Based Metabolomics Using Pure Ion Chromatograms. Analytical Chemistry, 2017, 89, 7631-7640.	3.2	25
15	UPLC-ESI-IT-TOF-MS metabolomic study of the therapeutic effect of Xuefu Zhuyu decoction on rats with traumatic brain injury. Journal of Ethnopharmacology, 2019, 245, 112149.	2.0	24
16	Qualitative analysis of major constituents from Xue Fu Zhu Yu Decoction using ultra high performance liquid chromatography with hybrid ion trap timeâ€ofâ€flight mass spectrometry. Journal of Separation Science, 2016, 39, 3457-3468.	1.3	23
17	Characterizing semen abnormality male infertility using non-targeted blood plasma metabolomics. PLoS ONE, 2019, 14, e0219179.	1.1	23
18	The rapid determination of total polyphenols content and antioxidant activity in Dendrobium officinale using near-infrared spectroscopy. Analytical Methods, 2016, 8, 4584-4589.	1.3	21

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19	Sensitive surface enhanced Raman spectroscopy (SERS) detection of methotrexate by core-shell-satellite magnetic microspheres. Talanta, 2017, 171, 152-158.	2.9	21
20	Direct calibration transfer to principal components via canonical correlation analysis. Chemometrics and Intelligent Laboratory Systems, 2018, 181, 21-28.	1.8	21
21	Retention time prediction in hydrophilic interaction liquid chromatography with graph neural network and transfer learning. Journal of Chromatography A, 2021, 1656, 462536.	1.8	17
22	Recursive Wavelet Peak Detection of Analytical Signals. Chromatographia, 2016, 79, 1247-1255.	0.7	15
23	Enhancing coverage in LC–MS-based untargeted metabolomics by a new sample preparation procedure using mixed-mode solid-phase extraction and two derivatizations. Analytical and Bioanalytical Chemistry, 2019, 411, 6189-6202.	1.9	15
24	Fast and Low-Cost Surface-Enhanced Raman Scattering (SERS) Method for On-Site Detection of Flumetsulam in Wheat. Molecules, 2020, 25, 4662.	1.7	15
25	Feature extraction from resolution perspective for gas chromatography-mass spectrometry datasets. RSC Advances, 2016, 6, 113997-114004.	1.7	14
26	Scalable calibration transfer without standards <i>via</i> dynamic time warping for near-infrared spectroscopy. Analytical Methods, 2019, 11, 4481-4493.	1.3	14
27	Fully automatic resolution of untargeted GC-MS data with deep learning assistance. Talanta, 2022, 244, 123415.	2.9	13
28	Application of sparse linear discriminant analysis for metabolomics data. Analytical Methods, 2014, 6, 9037-9044.	1.3	12
29	Deep-Learning-Assisted multivariate curve resolution. Journal of Chromatography A, 2021, 1635, 461713.	1.8	12
30	Evaluation and prediction of the antioxidant activity of Epimedium from multi-wavelength chromatographic fingerprints and chemometrics. Analytical Methods, 2014, 6, 1036.	1.3	11
31	Pure ion chromatogram extraction via optimal k-means clustering. RSC Advances, 2016, 6, 56977-56985.	1.7	10
32	Deep Learning-Based Method for Compound Identification in NMR Spectra of Mixtures. Molecules, 2022, 27, 3653.	1.7	10
33	Separation of Glycolipids/Sphingolipids from Glycerophospholipids on TiO <sub>2</sub> Coating in Aprotic Solvent for Rapid Comprehensive Lipidomic Analysis with Liquid Microjunction Surface Sampling-Mass Spectrometry. Analytical Chemistry, 2020, 92, 11250-11259.	3.2	9
34	Mixture analysis using nonâ€negative elastic net for Raman spectroscopy. Journal of Chemometrics, 2020, 34, e3293.	0.7	8
35	Unitary and binary chromatographic fingerprints analysis of Epimedium. Analytical Methods, 2013, 5, 5331.	1.3	7
36	IsoResolve: predicting splice isoform functions by integrating gene and isoform-level features with domain adaptation. Bioinformatics, 2021, 37, 522-530.	1.8	7

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37	Exploring asthenozoospermia seminal plasma amino acid disorder based on GC-SIM-MS combined with chemometrics methods. Analytical Methods, 2019, 11, 2895-2902.	1.3	6
38	Sample classification of GC-ToF-MS metabolomics data without the requirement for chromatographic deconvolution. Metabolomics, 2011, 7, 191-205.	1.4	5
39	Metabolic profiling putatively identifies plasma biomarkers of male infertility using UPLC-ESI-IT-TOFMS. RSC Advances, 2018, 8, 25974-25982.	1.7	5
40	TarMet: a reactive GUI tool for efficient and confident quantification of MS based targeted metabolic and stable isotope tracer analysis. Metabolomics, 2018, 14, 68.	1.4	5
41	Development of a sensitive and rapid UHPLC–MS/MS method for simultaneous quantification of nine compounds in rat plasma and application in a comparative pharmacokinetic study after oral administration of Xuefu Zhuyu Decoction and nimodipine. Biomedical Chromatography, 2020, 34, e4872.	0.8	5
42	Rapid and sensitive detection of neotame in instant grain beverages by paperâ€based silver nanoparticles substrates. Micro and Nano Letters, 2020, 15, 1099-1104.	0.6	5
43	Standardization of Raman spectra using variable penalty dynamic time warping. Analytical Methods, 2021, 13, 3414-3423.	1.3	2
44	Pure Ion Chromatograms Combined with Advanced Machine Learning Methods Improve Accuracy of Discriminant Models in LC–MS-Based Untargeted Metabolomics. Molecules, 2021, 26, 2715.	1.7	2
45	Chromatographic Profiling with Machine Learning Discriminates the Maturity Grades of Nicotiana tabacum L. Leaves. Separations, 2021, 8, 9.	1.1	2
46	Chemometrics in instrumental analysis of complex systemsâ€"in honor and memory of Yiâ€Zeng Liang. Journal of Chemometrics, 2018, 32, e3095.	0.7	1
47	Two-Way Data Analysis: Multivariate Curve Resolution: Noniterative Resolution Methods. , 2020, , 137-152.		1
48	A GC-MS study of the stability of rat serum metabolome during the sample preparation procedure. Analytical Methods, 2013, 5, 6807.	1.3	0
49	Detection of cimetidine in human plasma by surfaceâ€enhanced Raman scattering. Micro and Nano Letters, 2020, 15, 514-518.	0.6	0