Wen Zou

List of Publications by Year in descending order

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38	1,266	16	33
papers	citations	h-index	g-index
39	39	39	1808
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Software-Assisted Pattern Recognition of Persistent Organic Pollutants in Contaminated Human and Animal Food. Molecules, 2021, 26, 685.	3.8	2
2	Text Fingerprinting and Topic Mining in the Prescription Opioid Use Literature. , 2021, , .		1
3	An end-to-end framework for biomedical event trigger identification with hierarchical attention and adaptive cost learning. International Journal of Data Mining and Bioinformatics, 2020, 23, 189.	0.1	O
4	QUICK: Quality and Usability Investigation and Control Kit for Mass Spectrometric Data from Detection of Persistent Organic Pollutants. International Journal of Environmental Research and Public Health, 2019, 16, 4203.	2.6	5
5	Persistent Organic Pollutants in Food: Contamination Sources, Health Effects and Detection Methods. International Journal of Environmental Research and Public Health, 2019, 16, 4361.	2.6	214
6	An Improved Biomedical Event Trigger Identification Framework via Modeling Document with Hierarchical Attention. , $2019, \ldots$		1
7	Endocrine Disrupting Chemicals Mediated through Binding Androgen Receptor Are Associated with Diabetes Mellitus. International Journal of Environmental Research and Public Health, 2018, 15, 25.	2.6	18
8	Best Setting of Model Parameters in Applying Topic Modeling on Textual Documents , 2017, , .		0
9	Mechanistic roles of microRNAs in hepatocarcinogenesis: A study of thioacetamide with multiple doses and time-points of rats. Scientific Reports, 2017, 7, 3054.	3.3	14
10	A novel procedure on next generation sequencing data analysis using text mining algorithm. BMC Bioinformatics, 2016, 17, 213.	2.6	16
11	Text mining for identifying topics in the literatures about adolescent substance use and depression. BMC Public Health, 2016, 16, 279.	2.9	67
12	A heuristic approach to determine an appropriate number of topics in topic modeling. BMC Bioinformatics, 2015, 16, S8.	2.6	202
13	Molecular regulation of miRNAs and potential biomarkers in the progression of hepatic steatosis to NASH. Biomarkers in Medicine, 2015, 9, 1189-1200.	1.4	12
14	Biomarker identification from next-generation sequencing data for pathogen bacteria characterization and surveillance. Biomarkers in Medicine, 2015, 9, 1253-1264.	1.4	5
15	Differential gene expression in Staphylococcus aureus exposed to Orange II and Sudan III azo dyes. Journal of Industrial Microbiology and Biotechnology, 2015, 42, 745-757.	3.0	9
16	Asymmetric author-topic model for knowledge discovering of big data in toxicogenomics. Frontiers in Pharmacology, 2015, 6, 81.	3.5	12
17	A Composite Model for Subgroup Identification and Prediction via Bicluster Analysis. PLoS ONE, 2014, 9, e111318.	2.5	3
18	Topic modeling for cluster analysis of large biological and medical datasets. BMC Bioinformatics, 2014, 15, S11.	2.6	52

#	Article	IF	Citations
19	Data mining tools for Salmonella characterization: application to gel-based fingerprinting analysis. BMC Bioinformatics, 2013, 14, S15.	2.6	22
20	Meta-Analysis of Pulsed-Field Gel Electrophoresis Fingerprints Based on a Constructed Salmonella Database. PLoS ONE, 2013, 8, e59224.	2.5	18
21	Identification of Bicluster Regions in a Binary Matrix and Its Applications. PLoS ONE, 2013, 8, e71680.	2.5	12
22	Prediction System for Rapid Identification of Salmonella Serotypes Based on Pulsed-Field Gel Electrophoresis Fingerprints. Journal of Clinical Microbiology, 2012, 50, 1524-1532.	3.9	26
23	Microarray analysis of virulence gene profiles in Salmonella serovars from food/food animal environment. Journal of Infection in Developing Countries, 2011, 5, 94-105.	1.2	45
24	An FDA bioinformatics tool for microbial genomics research on molecular characterization of bacterial foodborne pathogens using microarrays. BMC Bioinformatics, 2010, 11, S4.	2.6	53
25	Evaluation of Pulsed-Field Gel Electrophoresis Profiles for Identification of <i>Salmonella</i> Serotypes. Journal of Clinical Microbiology, 2010, 48, 3122-3126.	3.9	46
26	Metabolism of azo dyes by human skin microbiota. Journal of Medical Microbiology, 2010, 59, 108-114.	1.8	79
27	Microarray analysis of antimicrobial resistance genes in <i>Salmonella enterica</i> from preharvest poultry environment. Journal of Applied Microbiology, 2009, 107, 906-914.	3.1	30
28	Azoreductase inStaphylococcus aureus. , 2009, Chapter 4, Unit4.28.		2
29	Detection of protein–protein interactions by a combination of a novel cytoplasmic membrane targeting system of recombinant proteins and fluorescence resonance energy transfer. Applied Microbiology and Biotechnology, 2006, 70, 451-457.	3.6	17
30	Activation of TBK1 and IKK $\hat{l}\mu$ Kinases by Vesicular Stomatitis Virus Infection and the Role of Viral Ribonucleoprotein in the Development of Interferon Antiviral Immunity. Journal of Virology, 2004, 78, 10636-10649.	3.4	164
31	Structure-function study of the amino-terminal stretch of the catalase subunit molecule in oligomerization, heme binding, and activity expression. Applied Microbiology and Biotechnology, 2003, 61, 488-494.	3.6	15
32	Construction of a Combinatorial Protein Library Displayed on Yeast Cell Surface and Its Application to Molecular Breeding. ACS Symposium Series, 2002, , 220-233.	0.5	0
33	Screening of a molecule endowing Saccharomyces cerevisiae with n -nonane-tolerance from a combinatorial random protein library. Applied Microbiology and Biotechnology, 2002, 58, 806-812.	3.6	31
34	Construction of a combinatorial protein library displayed on yeast cell surface using DNA random priming method. Journal of Bioscience and Bioengineering, 2001, 92, 393-396.	2.2	14
35	Genetically Controlled Self-Aggregation of Cell-Surface-Engineered Yeast Responding to Glucose Concentration. Applied and Environmental Microbiology, 2001, 67, 2083-2087.	3.1	7
36	Establishment of a simple system to analyse the molecular interaction in the agglutination of Saccharomyces cerevisiae. Yeast, 2000, 16, 995-1000.	1.7	3

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37	Cytochemical evaluation of localization and secretion of a heterologous enzyme displayed on yeast cell surface. FEMS Microbiology Letters, 2000, 192, 243-248.	1.8	12
38	Screening of Genes Involved in Isooctane Tolerance in Saccharomyces cerevisiae by Using mRNA Differential Display. Applied and Environmental Microbiology, 2000, 66, 4883-4889.	3.1	36