

Bin Zhou

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

2,740
citations

1040056

9
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

5431
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetics and Pathogenesis of Diffuse Large B-Cell Lymphoma. <i>New England Journal of Medicine</i> , 2018, 378, 1396-1407.	27.0	1,443
2	Ionizing Radiation Impairs T Cell Activation by Affecting Metabolic Reprogramming. <i>International Journal of Biological Sciences</i> , 2015, 11, 726-736.	6.4	35
3	Hepatitis C virus Genotype 1a core gene nucleotide patterns associated with hepatocellular carcinoma risk. <i>Journal of General Virology</i> , 2015, 96, 2928-2937.	2.9	7
4	Evaluation of Metabolite Biomarkers for Hepatocellular Carcinoma through Stratified Analysis by Gender, Race, and Alcoholic Cirrhosis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 64-72.	2.5	9
5	Prioritization of putative metabolite identifications in LC-MS/MS experiments using a computational pipeline. <i>Proteomics</i> , 2013, 13, 248-260.	2.2	8
6	Feasibility of Identifying the Tobacco-related Global Metabolome in Blood by UPLC-QTOF-MS. <i>Journal of Proteome Research</i> , 2013, 12, 679-691.	3.7	34
7	LC-MS-based metabolomics. <i>Molecular BioSystems</i> , 2012, 8, 470-481.	2.9	440
8	Utilization of metabolomics to identify serum biomarkers for hepatocellular carcinoma in patients with liver cirrhosis. <i>Analytica Chimica Acta</i> , 2012, 743, 90-100.	5.4	146
9	LC-MS Based Serum Metabolomics for Identification of Hepatocellular Carcinoma Biomarkers in Egyptian Cohort. <i>Journal of Proteome Research</i> , 2012, 11, 5914-5923.	3.7	116
10	Metabolite identification and quantitation in LC-MS/MS-based metabolomics. <i>TrAC - Trends in Analytical Chemistry</i> , 2012, 32, 1-14.	11.4	424
11	MetaboSearch: Tool for Mass-Based Metabolite Identification Using Multiple Databases. <i>PLoS ONE</i> , 2012, 7, e40096.	2.5	71
12	A Computational Pipeline for LC-MS/MS Based Metabolite Identification. , 2011, , .		3
13	Meta-analysis of LC-MS based metabolomic experiments. , 2011, , .		0
14	SVM-based spectral matching for metabolite identification. , 2010, 2010, 756-9.		4