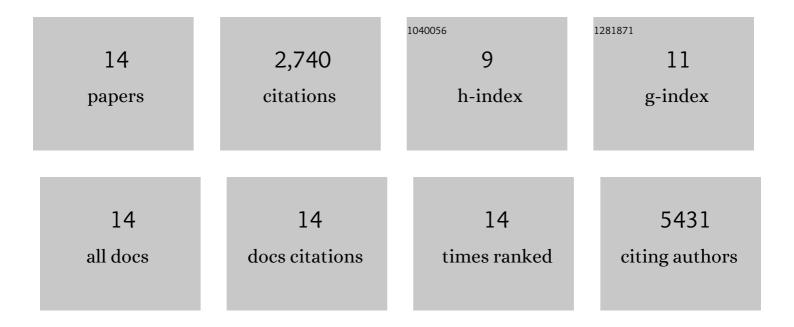
Bin Zhou

List of Publications by Year in descending order

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Βιν Ζησι

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

14 Meta-analysis of LC-MS based metabolomic experiments. , 2011, , .