Bao-Jin Zhou

List of Publications by Year
in descending order

11	414	1307594	1281871
papers	citations	7	11
		h-index	g-index

11	11	11	701
all docs	docs citations	times ranked	citing authors

#	ARTICLE	IF	CITATIONS
1	Deep functional analysis of synII, a 770-kilobase synthetic yeast chromosome. Science, 2017, 355, .	12.6	163
2	Distinct gut metagenomics and metaproteomics signatures in prediabetics and treatment-naÃ¯ve type 2 diabetics. EBioMedicine, 2019, 47, 373-383.	6.1	101
3	Identification and Characterization of Major Constituents in Different-Colored Rapeseed Petals by UPLCâ€"HESI-MS/MS. Journal of Agricultural and Food Chemistry, 2019, 67, 11053-11065.	5.2	51
4	Comparative Analysis of the Metabolic Profiles of Yellow- versus Black-Seeded Rapeseed Using UPLCâ€"HESIâ€"MS/MS and Transcriptome Analysis. Journal of Agricultural and Food Chemistry, 2020, 68, 3033-3049.	5.2	23
5	iTRAQ-based pharmacoproteomics reveals potential targets of berberine, a promising therapy for ulcerative colitis. European Journal of Pharmacology, 2019, 850, 167-179.	3.5	22
6	Integrated Analysis of Comparative Lipidomics and Proteomics Reveals the Dynamic Changes of Lipid Molecular Species in High-Oleic Acid Peanut Seed. Journal of Agricultural and Food Chemistry, 2020, 68, 426-438.	5.2	21
7	Comparative Analysis of Proteomics and Transcriptomics during Fertility Transition in a Two-Line Hybrid Rice Line Wuxiang S. International Journal of Molecular Sciences, 2019, 20, 4542.	4.1	8
8	Clinical Multi-Omics Study on the Gut Microbiota in Critically Ill Patients After Cardiovascular Surgery Combined With Cardiopulmonary Bypass With or Without Sepsis (MUL-GM-CSCPB Study): A Prospective Study Protocol. Frontiers in Medicine, 2020, 7, 269.	2.6	8
9	A proteomic analysis of peanut seed at different stages of underground development to understand the changes of seed proteins. PLoS ONE, 2020, 15, e0243132.	2.5	8
10	Improving Gene Annotation of the Peanut Genome by Integrated Proteogenomics Workflow. Journal of Proteome Research, 2020, 19, 2226-2235.	3.7	7
11	PPIP: Automated Software for Identification of Bioactive Endogenous Peptides. Journal of Proteome Research, 2019, 18, 721-727.	3.7	2