

Bao-Jin Zhou

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

414
citations

1307594

7
h-index

1281871

11
g-index

11
all docs

11
docs citations

11
times ranked

701
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrated Analysis of Comparative Lipidomics and Proteomics Reveals the Dynamic Changes of Lipid Molecular Species in High-Oleic Acid Peanut Seed. <i>Journal of Agricultural and Food Chemistry</i> , 2020, 68, 426-438.	5.2	21
2	Improving Gene Annotation of the Peanut Genome by Integrated Proteogenomics Workflow. <i>Journal of Proteome Research</i> , 2020, 19, 2226-2235.	3.7	7
3	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-CSCP B Study): A Prospective Study Protocol. <i>Frontiers in Medicine</i> , 2020, 7, 269.	2.6	8
4	Comparative Analysis of the Metabolic Profiles of Yellow- versus Black-Seeded Rapeseed Using UPLC-HESI-MS/MS and Transcriptome Analysis. <i>Journal of Agricultural and Food Chemistry</i> , 2020, 68, 3033-3049.	5.2	23
5	A proteomic analysis of peanut seed at different stages of underground development to understand the changes of seed proteins. <i>PLoS ONE</i> , 2020, 15, e0243132.	2.5	8
6	Distinct gut metagenomics and metaproteomics signatures in prediabetics and treatment-naïve type 2 diabetics. <i>EBioMedicine</i> , 2019, 47, 373-383.	6.1	101
7	Identification and Characterization of Major Constituents in Different-Colored Rapeseed Petals by UPLC-HESI-MS/MS. <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 11053-11065.	5.2	51
8	Comparative Analysis of Proteomics and Transcriptomics during Fertility Transition in a Two-Line Hybrid Rice Line Wuxiang S. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4542.	4.1	8
9	iTRAQ-based pharmacoproteomics reveals potential targets of berberine, a promising therapy for ulcerative colitis. <i>European Journal of Pharmacology</i> , 2019, 850, 167-179.	3.5	22
10	PPIP: Automated Software for Identification of Bioactive Endogenous Peptides. <i>Journal of Proteome Research</i> , 2019, 18, 721-727.	3.7	2
11	Deep functional analysis of synIII, a 770-kilobase synthetic yeast chromosome. <i>Science</i> , 2017, 355, .	12.6	163