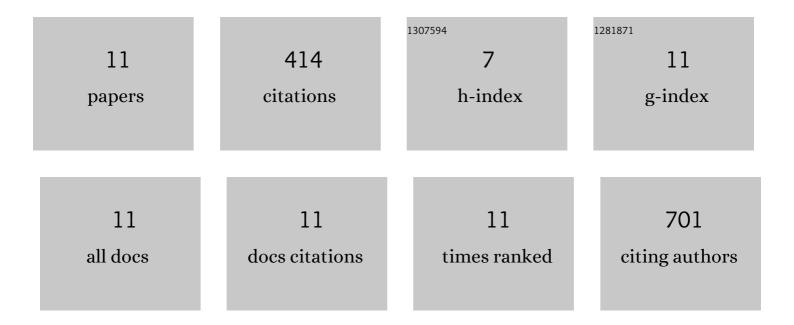
Bao-Jin Zhou

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6345873/publications.pdf Version: 2024-02-01



<u> Βλο-Ιινι Ζμου</u>

| # | 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. Journal of Agricultural and Food Chemistry, 2020, 68, 426-438. | 5.2 | 21 |
| 2 | Improving Gene Annotation of the Peanut Genome by Integrated Proteogenomics Workflow. Journal of Proteome Research, 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-CSCPB Study): A Prospective Study Protocol. Frontiers in Medicine, 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. Journal of Agricultural and Food Chemistry, 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. PLoS ONE, 2020, 15, e0243132. | 2.5 | 8 |
| 6 | Distinct gut metagenomics and metaproteomics signatures in prediabetics and treatment-naÃ ⁻ ve type 2 diabetics. EBioMedicine, 2019, 47, 373-383. | 6.1 | 101 |
| 7 | Identification and Characterization of Major Constituents in Different-Colored Rapeseed Petals by UPLC–HESI-MS/MS. Journal of Agricultural and Food Chemistry, 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. International Journal of Molecular Sciences, 2019, 20, 4542. | 4.1 | 8 |
| 9 | 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 |
| 10 | PPIP: Automated Software for Identification of Bioactive Endogenous Peptides. Journal of Proteome Research, 2019, 18, 721-727. | 3.7 | 2 |
| 11 | Deep functional analysis of synII, a 770-kilobase synthetic yeast chromosome. Science, 2017, 355, . | 12.6 | 163 |