

Jun Huang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5311513/publications.pdf>

Version: 2024-02-01

8
papers

73
citations

1937685
4
h-index

1588992
8
g-index

8
all docs

8
docs citations

8
times ranked

52
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-Wide Identification and Characterization of Actin-Depolymerizing Factor (ADF) Family Genes and Expression Analysis of Responses to Various Stresses in Zea Mays L.. International Journal of Molecular Sciences, 2020, 21, 1751.	4.1	22
2	QTL mapping and transcriptome analysis identify candidate genes regulating pericarp thickness in sweet corn. BMC Plant Biology, 2020, 20, 117.	3.6	12
3	Genome-wide identification, phylogenetic and expression analysis of the maize HECT E3 ubiquitin ligase genes. Genetica, 2019, 147, 391-400.	1.1	10
4	Mitochondrion-encoded circular RNAs are widespread and translatable in plants. Plant Physiology, 2022, 189, 1482-1500.	4.8	9
5	Construction of High-Density Genetic Map and Identification of QTLs Associated with Seed Vigor after Exposure to Artificial Aging Conditions in Sweet Corn Using SLAF-seq. Genes, 2020, 11, 37.	2.4	7
6	Widely Targeted Metabolomics Analysis Reveals Key Quality-Related Metabolites in Kernels of Sweet Corn. International Journal of Genomics, 2021, 2021, 1-12.	1.6	7
7	Comparative Transcriptome Analysis Reveals Regulatory Networks during the Maize Ear Shank Elongation Process. International Journal of Molecular Sciences, 2021, 22, 7029.	4.1	3
8	Construction of maize-teosinte introgression line population and identification of major quantitative trait loci. Euphytica, 2021, 217, 1.	1.2	3