

Jun Huang

List of Publications by Year in descending order

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

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