

Yue-Ie Caroline Hsing

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

25
papers

629
citations

11
h-index

25
g-index

27
ext. papers

818
ext. citations

6
avg, IF

3.75
L-index

#	Paper	IF	Citations
25	Closer vein spacing by ectopic expression of nucleotide-binding and leucine-rich repeat proteins in rice leaves. <i>Plant Cell Reports</i> , 2021 , 1	5.1	0
24	A drought-responsive rice amidohydrolase is the elusive plant guanine deaminase with the potential to modulate the epigenome. <i>Physiologia Plantarum</i> , 2021 , 172, 1853-1866	4.6	2
23	Genome Analysis Traces Regional Dispersal of Rice in Taiwan and Southeast Asia. <i>Molecular Biology and Evolution</i> , 2021 , 38, 4832-4846	8.3	4
22	Modulation of Rice Leaf Angle and Grain Size by Expressing and under the Control of Promoter. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	4
21	Rice Big Grain 1 promotes cell division to enhance organ development, stress tolerance and grain yield. <i>Plant Biotechnology Journal</i> , 2020 , 18, 1969-1983	11.6	8
20	Studies of rice Hd1 haplotypes worldwide reveal adaptation of flowering time to different environments. <i>PLoS ONE</i> , 2020 , 15, e0239028	3.7	3
19	Studies of rice Hd1 haplotypes worldwide reveal adaptation of flowering time to different environments 2020 , 15, e0239028		
18	Studies of rice Hd1 haplotypes worldwide reveal adaptation of flowering time to different environments 2020 , 15, e0239028		
17	Studies of rice Hd1 haplotypes worldwide reveal adaptation of flowering time to different environments 2020 , 15, e0239028		
16	Studies of rice Hd1 haplotypes worldwide reveal adaptation of flowering time to different environments 2020 , 15, e0239028		
15	Studies of rice Hd1 haplotypes worldwide reveal adaptation of flowering time to different environments 2020 , 15, e0239028		
14	Studies of rice Hd1 haplotypes worldwide reveal adaptation of flowering time to different environments 2020 , 15, e0239028		
13	OsDCL1a activation impairs phytoalexin biosynthesis and compromises disease resistance in rice. <i>Annals of Botany</i> , 2019 , 123, 79-93	4.1	12
12	Osa-miR7695 enhances transcriptional priming in defense responses against the rice blast fungus. <i>BMC Plant Biology</i> , 2019 , 19, 563	5.3	17
11	Retrotranspositional landscape of Asian rice revealed by 3000 genomes. <i>Nature Communications</i> , 2019 , 10, 24	17.4	55
10	The Polycistronic miR166k-166h Positively Regulates Rice Immunity via Post-transcriptional Control of. <i>Frontiers in Plant Science</i> , 2018 , 9, 337	6.2	44
9	A northern Chinese origin of Austronesian agriculture: new evidence on traditional Formosan cereals. <i>Rice</i> , 2018 , 11, 57	5.8	6

8	The Rice Paradox: Multiple Origins but Single Domestication in Asian Rice. <i>Molecular Biology and Evolution</i> , 2017 , 34, 969-979	8.3	124
7	Broomcorn and foxtail millet were cultivated in Taiwan about 5000 years ago. <i>Botanical Studies</i> , 2017 , 58, 3	2.3	20
6	Lack of Genotype and Phenotype Correlation in a Rice T-DNA Tagged Line Is Likely Caused by Introgression in the Seed Source. <i>PLoS ONE</i> , 2016 , 11, e0155768	3.7	5
5	Genetic resources offer efficient tools for rice functional genomics research. <i>Plant, Cell and Environment</i> , 2016 , 39, 998-1013	8.4	33
4	Genome-Wide Analysis of Polycistronic MicroRNAs in Cultivated and Wild Rice. <i>Genome Biology and Evolution</i> , 2016 , 8, 1104-14	3.9	23
3	RiTE database: a resource database for genus-wide rice genomics and evolutionary biology. <i>BMC Genomics</i> , 2015 , 16, 538	4.5	56
2	Genetic factors responsible for eating and cooking qualities of rice grains in a recombinant inbred population of an inter-subspecific cross. <i>Molecular Breeding</i> , 2014 , 34, 655-673	3.4	41
1	A rice gene activation/knockout mutant resource for high throughput functional genomics. <i>Plant Molecular Biology</i> , 2007 , 63, 351-64	4.6	172