

Jian-ping Cheng

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

594
citations

1162889

8
h-index

1058333

14
g-index

14
all docs

14
docs citations

14
times ranked

601
citing authors

#	ARTICLE	IF	CITATIONS
1	Bioactive Components and Health Functions of Oat. <i>Food Reviews International</i> , 2023, 39, 4545-4564.	4.3	13
2	Genome-wide identification, phylogenetic and expression pattern analysis of MADS-box family genes in foxtail millet (<i>Setaria italica</i>). <i>Scientific Reports</i> , 2022, 12, 4979.	1.6	12
3	Genome-wide identification and characterization of the SPL gene family and its expression in the various developmental stages and stress conditions in foxtail millet (<i>Setaria italica</i>). <i>BMC Genomics</i> , 2022, 23, .	1.2	9
4	Genome-wide identification and phylogenetic relationships of the Hsp70 gene family of <i>Aegilops tauschii</i> , wild emmer wheat (<i>Triticum dicoccoides</i>) and bread wheat (<i>Triticum aestivum</i>). <i>3 Biotech</i> , 2021, 11, 301.	1.1	6
5	Genome-wide identification and expression analysis of the bHLH transcription factor family and its response to abiotic stress in sorghum [<i>Sorghum bicolor</i> (L.) Moench]. <i>BMC Genomics</i> , 2021, 22, 415.	1.2	29
6	Genome-wide identification, expression analysis, and functional study of the GRAS transcription factor family and its response to abiotic stress in sorghum [<i>Sorghum bicolor</i> (L.) Moench]. <i>BMC Genomics</i> , 2021, 22, 509.	1.2	28
7	Genome-wide identification and expression analysis of the bHLH transcription factor family and its response to abiotic stress in foxtail millet (<i>Setaria italica</i> L.). <i>BMC Genomics</i> , 2021, 22, 778.	1.2	10
8	Genome-wide investigation of the GRAS transcription factor family in foxtail millet (<i>Setaria italica</i> L.). <i>BMC Plant Biology</i> , 2021, 21, 508.	1.6	19
9	JAZ8 Interacts With VirE3 Attenuating <i>Agrobacterium</i> Mediated Root Tumorigenesis. <i>Frontiers in Plant Science</i> , 2021, 12, 685533.	1.7	6
10	The Complete Chloroplast Genome Sequences of Eight <i>Fagopyrum</i> Species: Insights Into Genome Evolution and Phylogenetic Relationships. <i>Frontiers in Plant Science</i> , 2021, 12, 799904.	1.7	17
11	Genetic dissection of winter barley seedling response to salt and osmotic stress. <i>Molecular Breeding</i> , 2019, 39, 1.	1.0	11
12	Jasmonic Acid Signaling Pathway in Plants. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2479.	1.8	417
13	Molecular Cloning and Structureâ€“Function Analysis of a Trypsin Inhibitor from Tartary Buckwheat and Its Application in Combating Phytopathogenic Fungi. <i>Agronomy</i> , 2018, 8, 46.	1.3	1
14	A major QTL on chromosome 7HS controls the response of barley seedling to salt stress in the Nureâ€“Tremois population. <i>BMC Genetics</i> , 2017, 18, 79.	2.7	16