

Fcx007

List of Publications by Year in descending order

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times ranked

744
citing authors

#	ARTICLE	IF	CITATIONS
1	Deciphering the Molecular Mechanisms of Chilling Tolerance in Lsi1-Overexpressing Rice. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4667.	4.1	4
2	Silicon Modulates Molecular and Physiological Activities in Lsi1 Transgenic and Wild Lemont Rice Seedlings under Arsenic Stress. <i>Agronomy</i> , 2021, 11, 1532.	3.0	4
3	Usage of Si, P, Se, and Ca Decrease Arsenic Concentration/Toxicity in Rice, a Review. <i>Applied Sciences (Switzerland)</i> , 2021, 11, 8090.	2.5	11
4	Comparison of Silicon-Evoked Responses on Arsenic Stress between Different Dular Rice Genotypes. <i>Plants</i> , 2021, 10, 2210.	3.5	4
5	MYB57 transcriptionally regulates MAPK11 to interact with PAL2;3 and modulate rice allelopathy. <i>Journal of Experimental Botany</i> , 2020, 71, 2127-2141.	4.8	14
6	Physiochemical mechanisms involved in the improvement of grain-filling, rice quality mediated by related enzyme activities in the ratoon cultivation system. <i>Field Crops Research</i> , 2020, 258, 107962.	5.1	27
7	Serine hydroxymethyltransferase localised in the endoplasmic reticulum plays a role in scavenging H ₂ O ₂ to enhance rice chilling tolerance. <i>BMC Plant Biology</i> , 2020, 20, 236.	3.6	15
8	Lsi1 plays an active role in enhancing the chilling tolerance of rice roots. <i>Plant Growth Regulation</i> , 2020, 90, 529-543.	3.4	7
9	Lsi1 modulates the antioxidant capacity of rice and protects against ultraviolet-B radiation. <i>Plant Science</i> , 2019, 278, 96-106.	3.6	14
10	Proteomic analysis of positive influence of alternate wetting and moderate soil drying on the process of rice grain filling. <i>Plant Growth Regulation</i> , 2018, 84, 533-548.	3.4	14
11	Increasing Rice Allelopathy by Induction of Barnyard Grass (<i>Echinochloa crus-galli</i>) Root Exudates. <i>Journal of Plant Growth Regulation</i> , 2018, 37, 745-754.	5.1	21
12	Protein Phosphatase (PP2C9) Induces Protein Expression Differentially to Mediate Nitrogen Utilization Efficiency in Rice under Nitrogen-Deficient Condition. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2827.	4.1	18
13	Lsi1-regulated Cd uptake and phytohormones accumulation in rice seedlings in presence of Si. <i>Plant Growth Regulation</i> , 2018, 86, 149-157.	3.4	7
14	Overexpression of Lsi1 in cold-sensitive rice mediates transcriptional regulatory networks and enhances resistance to chilling stress. <i>Plant Science</i> , 2017, 262, 115-126.	3.6	41
15	Cadmium-stress mitigation through gene expression of rice and silicon addition. <i>Plant Growth Regulation</i> , 2017, 81, 91-101.	3.4	32
16	Mixed Phenolic Acids Mediated Proliferation of Pathogens <i>Talaromyces helicus</i> and <i>Kosakonia sacchari</i> in Continuously Monocultured <i>Radix pseudostellariae</i> Rhizosphere Soil. <i>Frontiers in Microbiology</i> , 2016, 7, 335.	3.5	66
17	Effect of silicon on grain yield of rice under cadmium-stress. <i>Acta Physiologiae Plantarum</i> , 2016, 38, 1.	2.1	18
18	Methyl-CpG binding domain protein acts to regulate the repair of cyclobutane pyrimidine dimers on rice DNA. <i>Scientific Reports</i> , 2016, 6, 34569.	3.3	8

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19	Role of allene oxide cyclase in the regulation of rice phenolic acids synthesis and allelopathic inhibition on barnyardgrass. <i>Plant Growth Regulation</i> , 2016, 79, 265-273.	3.4	5
20	Soil Microbial Community Structure and Metabolic Activity of <i>Pinus elliottii</i> Plantations across Different Stand Ages in a Subtropical Area. <i>PLoS ONE</i> , 2015, 10, e0135354.	2.5	41
21	Interaction of <i>Pseudostellaria heterophylla</i> with <i>Fusarium oxysporum</i> f.sp. <i>heterophylla</i> mediated by its root exudates in a consecutive monoculture system. <i>Scientific Reports</i> , 2015, 5, 8197.	3.3	57
22	Identification and comparative analysis of microRNAs in barnyardgrass (<i>Chenopodium album</i>) in response to rice allelopathy. <i>Plant, Cell and Environment</i> , 2015, 38, 1368-1381.	5.7	30
23	Terminal Restriction Fragment Length Polymorphism Analysis of Soil Bacterial Communities under Different Vegetation Types in Subtropical Area. <i>PLoS ONE</i> , 2015, 10, e0129397.	2.5	18
24	Method for RNA extraction and cDNA library construction from microbes in crop rhizosphere soil. <i>World Journal of Microbiology and Biotechnology</i> , 2014, 30, 783-789.	3.6	7
25	Changes in Rice Allelopathy and Rhizosphere Microflora by Inhibiting Rice Phenylalanine Ammonia-lyase Gene Expression. <i>Journal of Chemical Ecology</i> , 2013, 39, 204-212.	1.8	77
26	Barnyard grass stress up regulates the biosynthesis of phenolic compounds in allelopathic rice. <i>Journal of Plant Physiology</i> , 2012, 169, 1747-1753.	3.5	46
27	Proteomic and phosphoproteomic determination of ABA's effects on grain-filling of <i>Oryza sativa</i> L. inferior spikelets. <i>Plant Science</i> , 2012, 185-186, 259-273.	3.6	38
28	Molecular physiological mechanism on consecutive monoculture problems of <i>Rehmannia glutinosa</i> . <i>Journal of Integrated OMICS</i> , 2011, 1, .	0.5	0
29	Rice allelopathy and its properties of molecular ecology. <i>Frontiers in Biology</i> , 2010, 5, 255-262.	0.7	7
30	Molecular Physiological Properties of Rice on the Enhanced Weed-Suppression Ability Induced by Lower Phosphorus Supplies*. <i>Ying Yong Yu Huan Jing Sheng Wu Xue Bao = Chinese Journal of Applied and Environmental Biology</i> , 2010, 2009, 289-294.	0.1	0
31	Allelopathic Enhancement and Differential Gene Expression in Rice under Low Nitrogen Treatment. <i>Journal of Chemical Ecology</i> , 2008, 34, 688-695.	1.8	41