

Zhigang Li

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

1,251
citations

567281

15
h-index

888059

17
g-index

19
all docs

19
docs citations

19
times ranked

1490
citing authors

#	ARTICLE	IF	CITATIONS
1	Constitutive Expression of a <i>miR319</i> Gene Alters Plant Development and Enhances Salt and Drought Tolerance in Transgenic Creeping Bentgrass. <i>Plant Physiology</i> , 2013, 161, 1375-1391.	4.8	378
2	Heterologous expression of <i>Arabidopsis</i> H ⁺ pyrophosphatase enhances salt tolerance in transgenic creeping bentgrass (<i>Agrostis stolonifera</i> L.). <i>Plant, Cell and Environment</i> , 2010, 33, 272-289.	5.7	158
3	Constitutive Expression of Rice <i>MicroRNA528</i> Alters Plant Development and Enhances Tolerance to Salinity Stress and Nitrogen Starvation in Creeping Bentgrass. <i>Plant Physiology</i> , 2015, 169, 576-593.	4.8	136
4	AsHSP17, a creeping bentgrass small heat shock protein modulates plant photosynthesis and ABA-dependent and independent signalling to attenuate plant response to abiotic stress. <i>Plant, Cell and Environment</i> , 2016, 39, 1320-1337.	5.7	82
5	Heterologous expression of <i>OsSIZ1</i> , a rice <i>SUMO E3</i> ligase, enhances broad abiotic stress tolerance in transgenic creeping bentgrass. <i>Plant Biotechnology Journal</i> , 2013, 11, 432-445.	8.3	79
6	Transgenic creeping bentgrass overexpressing <i>Osa-miR393a</i> exhibits altered plant development and improved multiple stress tolerance. <i>Plant Biotechnology Journal</i> , 2019, 17, 233-251.	8.3	75
7	Ectopic expression of a cyanobacterial flavodoxin in creeping bentgrass impacts plant development and confers broad abiotic stress tolerance. <i>Plant Biotechnology Journal</i> , 2017, 15, 433-446.	8.3	66
8	MicroRNA396-mediated alteration in plant development and salinity stress response in creeping bentgrass. <i>Horticulture Research</i> , 2019, 6, 48.	6.3	64
9	STRESS INDUCED FACTOR 2, a Leucine-Rich Repeat Kinase Regulates Basal Plant Pathogen Defense. <i>Plant Physiology</i> , 2018, 176, 3062-3080.	4.8	49
10	Heterologous expression of a rice <i>miR395</i> gene in <i>Nicotiana tabacum</i> impairs sulfate homeostasis. <i>Scientific Reports</i> , 2016, 6, 28791.	3.3	29
11	AsHSP26.8a, a creeping bentgrass small heat shock protein integrates different signaling pathways to modulate plant abiotic stress response. <i>BMC Plant Biology</i> , 2020, 20, 184.	3.6	27
12	Comparative transcriptome profiling provides insights into plant salt tolerance in seashore paspalum (<i>Paspalum vaginatum</i>). <i>BMC Genomics</i> , 2020, 21, 131.	2.8	26
13	MiR396 is involved in plant response to vernalization and flower development in <i>Agrostis stolonifera</i> . <i>Horticulture Research</i> , 2020, 7, 173.	6.3	21
14	Transcriptomic profiles of non-embryogenic and embryogenic callus cells in a highly regenerative upland cotton line (<i>Gossypium hirsutum</i> L.). <i>BMC Developmental Biology</i> , 2020, 20, 25.	2.1	19
15	Adventitious shoot regeneration from in vitro cultured leaf explants of peach rootstock Guardian® is significantly enhanced by silver thiosulfate. <i>Plant Cell, Tissue and Organ Culture</i> , 2015, 120, 757-765.	2.3	17
16	DRMY1, a Myb-Like Protein, Regulates Cell Expansion and Seed Production in <i>Arabidopsis thaliana</i> . <i>Plant and Cell Physiology</i> , 2019, 60, 285-302.	3.1	15
17	Comparative Transcriptomics of Non-Embryogenic and Embryogenic Callus in Semi-Recalcitrant and Non-Recalcitrant Upland Cotton Lines. <i>Plants</i> , 2021, 10, 1775.	3.5	10
18	Manipulating Expression of Tonoplast Transporters. , 2012, 913, 359-369.		0

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19	Biolytic DNA Delivery in Turfgrass Embryonic Callus Initiated from Mature Seeds. <i>Methods in Molecular Biology</i> , 2020, 2124, 251-261.	0.9	0