

Yiwei Jiang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7502703/publications.pdf>

Version: 2024-02-01

67
papers

2,328
citations

236925

25
h-index

223800

46
g-index

68
all docs

68
docs citations

68
times ranked

2538
citing authors

#	ARTICLE	IF	CITATIONS
1	Drought and Heat Stress Injury to Two Cool-Season Turfgrasses in Relation to Antioxidant Metabolism and Lipid Peroxidation. <i>Crop Science</i> , 2001, 41, 436-442.	1.8	379
2	Reactive oxygen species, antioxidant enzyme activities and gene expression patterns in leaves and roots of Kentucky bluegrass in response to drought stress and recovery. <i>Scientia Horticulturae</i> , 2009, 120, 264-270.	3.6	247
3	Research Advances in Mechanisms of Turfgrass Tolerance to Abiotic Stresses: From Physiology to Molecular Biology. <i>Critical Reviews in Plant Sciences</i> , 2014, 33, 141-189.	5.7	162
4	Protein Alterations in Tall Fescue in Response to Drought Stress and Abscisic Acid. <i>Crop Science</i> , 2002, 42, 202-207.	1.8	109
5	Effects of Drought or Heat Stress Alone and in Combination on Kentucky Bluegrass. <i>Crop Science</i> , 2000, 40, 1358-1362.	1.8	94
6	Association of candidate genes with drought tolerance traits in diverse perennial ryegrass accessions. <i>Journal of Experimental Botany</i> , 2013, 64, 1537-1551.	4.8	83
7	Osmotic Adjustment and Root Growth Associated with Drought Preconditioning-Enhanced Heat Tolerance in Kentucky Bluegrass. <i>Crop Science</i> , 2001, 41, 1168-1173.	1.8	79
8	Natural variation of drought response in <i>Brachypodium distachyon</i> . <i>Physiologia Plantarum</i> , 2011, 141, 19-29.	5.2	67
9	Genome-wide association analysis and QTL mapping reveal the genetic control of cadmium accumulation in maize leaf. <i>BMC Genomics</i> , 2018, 19, 91.	2.8	60
10	Genome-wide associations with flowering time in switchgrass using exome-capture sequencing data. <i>New Phytologist</i> , 2017, 213, 154-169.	7.3	56
11	Antioxidant Responses of Creeping Bentgrass Roots to Waterlogging. <i>Crop Science</i> , 2007, 47, 232-238.	1.8	51
12	Broadband Spectral Reflectance Models of Turfgrass Species and Cultivars to Drought Stress. <i>Crop Science</i> , 2007, 47, 1611-1618.	1.8	44
13	Physiological and transcriptomic responses of reproductive stage soybean to drought stress. <i>Plant Cell Reports</i> , 2018, 37, 1611-1624.	5.6	44
14	Identification of differentially expressed genes under drought stress in perennial ryegrass. <i>Physiologia Plantarum</i> , 2010, 139, no-no.	5.2	40
15	Genotypic variation in growth and metabolic responses of perennial ryegrass exposed to short-term waterlogging and submergence stress. <i>Plant Physiology and Biochemistry</i> , 2015, 95, 57-64.	5.8	39
16	Growth, Physiological, and Anatomical Responses of Creeping Bentgrass Cultivars to Different Depths of Waterlogging. <i>Crop Science</i> , 2006, 46, 2420-2426.	1.8	37
17	Candidate gene association mapping for winter survival and spring regrowth in perennial ryegrass. <i>Plant Science</i> , 2015, 235, 37-45.	3.6	37
18	Association of simple sequence repeat (SSR) markers with submergence tolerance in diverse populations of perennial ryegrass. <i>Plant Science</i> , 2011, 180, 391-398.	3.6	36

#	ARTICLE	IF	CITATIONS
19	Waterlogging Tolerance of Kentucky Bluegrass Cultivars. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2007, 42, 386-390.	1.0	36
20	Quantitative Trait Locus Mapping for Flowering Time in a Lowland × Upland Switchgrass Pseudo-F ₂ Population. <i>Plant Genome</i> , 2018, 11, 170093.	2.8	35
21	Correlations of Leaf Relative Water Content, Canopy Temperature, and Spectral Reflectance in Perennial Ryegrass Under Water Deficit Conditions. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2009, 44, 459-462.	1.0	35
22	Growth response, carbohydrate and ion accumulation of diverse perennial ryegrass accessions to increasing salinity. <i>Scientia Horticulturae</i> , 2013, 154, 73-81.	3.6	32
23	Differential growth response and carbohydrate metabolism of global collection of perennial ryegrass accessions to submergence and recovery following de-submergence. <i>Journal of Plant Physiology</i> , 2012, 169, 1040-1049.	3.5	30
24	Physiological Response, Cell Wall Components, and Gene Expression of Switchgrass under Short-Term Drought Stress and Recovery. <i>Crop Science</i> , 2012, 52, 2718-2727.	1.8	28
25	Quantitative Trait Loci Associated with Drought Tolerance in <i>Brachypodium distachyon</i> . <i>Frontiers in Plant Science</i> , 2017, 8, 811.	3.6	27
26	Natural variation of salinity response, population structure and candidate genes associated with salinity tolerance in perennial ryegrass accessions. <i>Plant, Cell and Environment</i> , 2013, 36, 2021-2033.	5.7	24
27	Ploidy Level and DNA Content of Perennial Ryegrass Germplasm as Determined by Flow Cytometry. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2009, 44, 2049-2052.	1.0	24
28	Genome-Wide Association Study in Pseudo-F ₂ Populations of Switchgrass Identifies Genetic Loci Affecting Heading and Anthesis Dates. <i>Frontiers in Plant Science</i> , 2018, 9, 1250.	3.6	22
29	Anaerobic metabolism in roots of Kentucky bluegrass in response to short-term waterlogging alone and in combination with high temperatures. <i>Plant and Soil</i> , 2009, 314, 221-229.	3.7	20
30	Antioxidative Responses and Candidate Gene Expression in Prairie Junegrass under Drought Stress. <i>Journal of the American Society for Horticultural Science</i> , 2010, 135, 303-309.	1.0	20
31	Genome-wide identification of <i>ZmHMAs</i> and association of natural variation in <i>ZmHMA2</i> and <i>ZmHMA3</i> with leaf cadmium accumulation in maize. <i>PeerJ</i> , 2019, 7, e7877.	2.0	20
32	Turfgrass Selection and Grass Clippings Management Influence Soil Carbon and Nitrogen Dynamics. <i>Agronomy Journal</i> , 2017, 109, 1719-1725.	1.8	19
33	Chlorophyll Metabolism and Gene Expression in Response to Submergence Stress and Subsequent Recovery in Perennial Ryegrass Accessions Differing in Growth Habits. <i>Journal of Plant Physiology</i> , 2020, 251, 153195.	3.5	19
34	Marker-Trait Association for Biomass Yield of Potential Bio-fuel Feedstock <i>Miscanthus sinensis</i> from Southwest China. <i>Frontiers in Plant Science</i> , 2016, 7, 802.	3.6	18
35	Natural variation of physiological traits, molecular markers, and chlorophyll catabolic genes associated with heat tolerance in perennial ryegrass accessions. <i>BMC Plant Biology</i> , 2020, 20, 520.	3.6	17
36	Differential antioxidant enzyme activity in rapid response glyphosate-resistant <i>Ambrosia trifida</i> . <i>Pest Management Science</i> , 2018, 74, 2125-2132.	3.4	14

#	ARTICLE	IF	CITATIONS
37	Growth and Hormone Alterations in Response to Heat Stress in Perennial Ryegrass Accessions Differing in Heat Tolerance. <i>Journal of Plant Growth Regulation</i> , 2020, 39, 1022-1029.	5.1	14
38	Effects of 6-Benzyladenine, $\hat{1}^3$ -Aminobutyric Acid, and Nitric Oxide on Plant Growth, Photochemical Efficiency, and Ion Accumulation of Perennial Ryegrass Cultivars to Salinity Stress. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2019, 54, 1418-1422.	1.0	13
39	Phenotypic and Genotypic Diversity for Drought Tolerance among and within Perennial Ryegrass Accessions. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2015, 50, 1148-1154.	1.0	11
40	Leaf and Root Growth, Carbon and Nitrogen Contents, and Gene Expression of Perennial Ryegrass to Different Nitrogen Supplies. <i>Journal of the American Society for Horticultural Science</i> , 2016, 141, 555-562.	1.0	10
41	Genotypic Variations in Plant Growth and Nutritional Elements of Perennial Ryegrass Accessions under Salinity Stress. <i>Journal of the American Society for Horticultural Science</i> , 2017, 142, 476-483.	1.0	10
42	Genome-Wide Association Study for Plant Architecture and Bioenergy Traits in Diverse Sorghum and Sudangrass Germplasm. <i>Agronomy</i> , 2020, 10, 1602.	3.0	10
43	Growth and Enzymatic Activity of Four Warm-season Turfgrass Species Exposed to Waterlogging. <i>Journal of the American Society for Horticultural Science</i> , 2015, 140, 151-162.	1.0	10
44	Transcriptome characterization of candidate genes for heat tolerance in perennial ryegrass after exogenous methyl Jasmonate application. <i>BMC Plant Biology</i> , 2022, 22, 68.	3.6	10
45	Interactive Short-term Effects of Waterlogging and Salinity Stress on Growth and Carbohydrate, Lipid Peroxidation, and Nutrients in Two Perennial Ryegrass Cultivars. <i>Journal of the American Society for Horticultural Science</i> , 2017, 142, 110-118.	1.0	9
46	Natural variation and genomic prediction of growth, physiological traits, and nitrogen-use efficiency in perennial ryegrass under low-nitrogen stress. <i>Journal of Experimental Botany</i> , 2020, 71, 6670-6683.	4.8	9
47	Diverse genotypic variations of photosynthetic capacity, transpiration and antioxidant enzymes of lily hybrids to increasing salinity stress. <i>Scientia Horticulturae</i> , 2021, 280, 109939.	3.6	9
48	Nucleotide diversity and linkage disequilibrium in antioxidant genes of <i>Brachypodium distachyon</i> . <i>Plant Science</i> , 2012, 197, 122-129.	3.6	8
49	The BRANCHING ENZYME1 gene, encoding a glycoside hydrolase family 13 protein, is required for in vitro plant regeneration in <i>Arabidopsis</i> . <i>Plant Cell, Tissue and Organ Culture</i> , 2014, 117, 279-291.	2.3	8
50	Specific peroxidases differentiate <i>Brachypodium distachyon</i> accessions and are associated with drought tolerance traits. <i>Annals of Botany</i> , 2016, 118, 259-270.	2.9	8
51	Association of Candidate Genes With Submergence Response in Perennial Ryegrass. <i>Frontiers in Plant Science</i> , 2017, 8, 791.	3.6	8
52	Growth, ionic response, and gene expression of shoots and roots of perennial ryegrass under salinity stress. <i>Acta Physiologiae Plantarum</i> , 2018, 40, 1.	2.1	8
53	Submergence stress alters fructan and hormone metabolism and gene expression in perennial ryegrass with contrasting growth habits. <i>Environmental and Experimental Botany</i> , 2020, 179, 104202.	4.2	8
54	Plant growth, ion accumulation, and antioxidant enzymes of endophyte-infected and endophyte-free tall fescue to salinity stress. <i>Acta Physiologiae Plantarum</i> , 2021, 43, 1.	2.1	7

#	ARTICLE	IF	CITATIONS
55	Association of candidate genes with drought tolerance traits in zoysiagrass germplasm. <i>Journal of Plant Physiology</i> , 2019, 237, 61-71.	3.5	6
56	Effects of Gamma-Aminobutyric Acid on Seed Germination, Ion Balance, and Metabolic Activity in Perennial Ryegrass Under Salinity Stress. <i>Journal of Plant Growth Regulation</i> , 2022, 41, 1835-1844.	5.1	6
57	Identification of Quantitative Trait Loci for Plant Height, Crown Diameter, and Plant Biomass in a Pseudo-F2 Population of Switchgrass. <i>Bioenergy Research</i> , 2019, 12, 267-274.	3.9	5
58	Nitrogen and carbon contents, nitrogen use efficiency, and antioxidant responses of perennial ryegrass accessions to nitrogen deficiency. <i>Journal of Plant Nutrition</i> , 2019, 42, 2092-2101.	1.9	5
59	The signal peptide of Cry1Ia can improve the expression of eGFP or mCherry in <i>Escherichia coli</i> and <i>Bacillus thuringiensis</i> and enhance the host's fluorescent intensity. <i>Microbial Cell Factories</i> , 2020, 19, 112.	4.0	5
60	Differential Metabolomic Responses of Kentucky Bluegrass Cultivars to Low Nitrogen Stress. <i>Frontiers in Plant Science</i> , 2021, 12, 808772.	3.6	5
61	Exogenous Application of Nitrogen and Cytokinin on Growth, Carbohydrate, and Antioxidant Metabolism of Creeping Bentgrass after De-submergence. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2016, 51, 1602-1606.	1.0	4
62	Minimal irrigation requirements of Kentucky bluegrass and tall fescue blends in the northern transition zone. <i>Crop Science</i> , 2020, 61, 2939.	1.8	4
63	Physiological and Molecular Responses of <i>Zoysia japonica</i> to Rust Infection. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4185.	4.1	3
64	Editorial: Genetics and Genomics of Polyploid Plants. <i>Frontiers in Plant Science</i> , 2019, 10, 934.	3.6	2
65	Transcriptome profiling reveals differentially expressed genes associated with flowering time in contrasting switchgrass genotypes. <i>Crop Science</i> , 2020, 60, 1472-1487.	1.8	2
66	Responses of Turfgrass to Low-Oxygen Stress. <i>Books in Soils, Plants, and the Environment</i> , 2007, , 531-545.	0.1	2
67	Turfgrass Water Use and Physiology. , 0, , 319-345.		0