

Jennifer A Kimball

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2974693/publications.pdf>

Version: 2024-02-01

22
papers

737
citations

933447

10
h-index

713466

21
g-index

27
all docs

27
docs citations

27
times ranked

1059
citing authors

#	ARTICLE	IF	CITATIONS
1	Recessive male floret color for tracking gene flow in cultivated northern wild rice (<i>Zizania) Tj ETQq1 1 0.784314.rgBT /Overlock 10	1.8	1
2	Dormancy breaking treatments in Northern Wild Rice (<i>Zizania palustris</i> L.) seed suggest a physiological source of Dormancy. <i>Plant Growth Regulation</i> , 2022, 98, 235-247.	3.4	1
3	Whole-genome assembly and annotation of northern wild rice, <i>Zizania palustris</i> L., supports a whole-genome duplication in the <i>Zizania</i> genus. <i>Plant Journal</i> , 2021, 107, 1802-1818.	5.7	7
4	Identification of single nucleotide polymorphism markers for population genetic studies in <i>Zizania palustris</i> L.. <i>Conservation Genetics Resources</i> , 2020, 12, 451-455.	0.8	4
5	<i>Herbaspirillum rubrisubalbicans</i> as a Phytopathogenic Model to Study the Immune System of <i>Sorghum bicolor</i>. <i>Molecular Plant-Microbe Interactions</i> , 2020, 33, 235-246.	2.6	15
6	Genome-wide association analysis of the strength of the MAMP-elicited defense response and resistance to target leaf spot in sorghum. <i>Scientific Reports</i> , 2020, 10, 20817.	3.3	12
7	Phenological stages of cultivated northern wild rice according to the BBCH scale. <i>Annals of Applied Biology</i> , 2020, 176, 350-356.	2.5	10
8	Investigation of variable storage conditions for cultivated northern wild rice and their effects on seed viability and dormancy. <i>Seed Science Research</i> , 2020, 30, 21-28.	1.7	10
9	Identification of QTL for Target Leaf Spot resistance in <i>Sorghum bicolor</i> and investigation of relationships between disease resistance and variation in the MAMP response. <i>Scientific Reports</i> , 2019, 9, 18285.	3.3	13
10	Quantifying MAMP-induced Production of Reactive Oxygen Species in Sorghum and Maize. <i>Bio-protocol</i> , 2019, 9, .	0.4	10
11	Linkage analysis and identification of quantitative trait loci associated with freeze tolerance and turf quality traits in St. Augustinegrass. <i>Molecular Breeding</i> , 2018, 38, 1.	2.1	7
12	Population Dynamics Among six Major Groups of the <i>Oryza rufipogon</i> Species Complex, Wild Relative of Cultivated Asian Rice. <i>Rice</i> , 2016, 9, 56.	4.0	80
13	Combining Ability for Winter Survival and Turf Quality Traits in St. Augustinegrass. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2016, 51, 810-815.	1.0	6
14	Validation of yield-enhancing quantitative trait loci from a low-yielding wild ancestor of rice. <i>Molecular Breeding</i> , 2013, 32, 101-120.	2.1	34
15	Genetic Relationships in <i>Zoysia</i> Species and the Identification of Putative Interspecific Hybrids Using Simple Sequence Repeat Markers and Inflorescence Traits. <i>Crop Science</i> , 2013, 53, 285-295.	1.8	24
16	Assessment of Genetic Diversity in <i>Zoysia</i> Species using Amplified Fragment Length Polymorphism Markers. <i>Crop Science</i> , 2012, 52, 360-370.	1.8	14
17	Use of sequence-related amplified polymorphism (SRAP) markers for comparing levels of genetic diversity in centipedegrass (<i>Eremochloa ophiuroides</i> (Munro) Hack.) germplasm. <i>Genetic Resources and Crop Evolution</i> , 2012, 59, 1517-1526.	1.6	12
18	Assessment of Molecular Variation within â€˜Raleighâ€™ St. Augustinegrass using Amplified Fragment Length Polymorphism Markers. <i>Hortscience: A Publication of the American Society for Horticultural Science</i> , 2012, 47, 839-844.	1.0	8

#	ARTICLE	IF	CITATIONS
19	A Rice Diversity Panel Evaluated for Genetic and Agro-Morphological Diversity between Subpopulations and its Geographic Distribution. <i>Crop Science</i> , 2011, 51, 2021-2035.	1.8	83
20	A universal core genetic map for rice. <i>Theoretical and Applied Genetics</i> , 2010, 120, 563-572.	3.6	60
21	Development of a Research Platform for Dissecting Phenotype-Genotype Associations in Rice (<i>Oryza</i>) Tj ETQq1 1 0.784314 rgBT / Q	4.0	75
22	Genomic Diversity and Introgression in <i>O. sativa</i> Reveal the Impact of Domestication and Breeding on the Rice Genome. <i>PLoS ONE</i> , 2010, 5, e10780.	2.5	250