Akshaya Kumar Biswal

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5101822/publications.pdf

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	933264	996849
906	10	15
citations	h-index	g-index
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1/	1/	1328
docs citations	times ranked	citing authors
	17	906 10 citations h-index 17 17

#	Article	IF	CITATIONS
1	Closer vein spacing by ectopic expression of nucleotide-binding and leucine-rich repeat proteins in rice leaves. Plant Cell Reports, 2022, 41, 319-335.	2.8	1
2	Globally Important Wheat Diseases: Status, Challenges, Breeding and Genomic Tools to Enhance Resistance Durability., 2021,, 59-128.		12
3	Novel Mutant Alleles Reveal a Role of the Extra-Large G Protein in Rice Grain Filling, Panicle Architecture, Plant Growth, and Disease Resistance. Frontiers in Plant Science, 2021, 12, 782960.	1.7	14
4	Rice with reduced stomatal density conserves water and has improved drought tolerance under future climate conditions. New Phytologist, 2019, 221, 371-384.	3.5	330
5	CRISPR mediated genome engineering to develop climate smart rice: Challenges and opportunities. Seminars in Cell and Developmental Biology, 2019, 96, 100-106.	2.3	32
6	The Nucleotideâ€Dependent Interactome of Rice Heterotrimeric Gâ€Protein α â€Subunit. Proteomics, 2019, 19, 1800385.	1.3	6
7	CRISPR-Cas9-Mediated Genome Editing of Rice Towards Better Grain Quality. Methods in Molecular Biology, 2019, 1892, 311-336.	0.4	7
8	Role of Biotechnology in Rice Production. , 2017, , 487-547.		7
9	CRISPR-Cas9 and CRISPR-Cpf1 mediated targeting of a stomatal developmental gene EPFL9 in rice. Plant Cell Reports, 2017, 36, 745-757.	2.8	170
10	Candidate regulators of Early Leaf Development in Maize Perturb Hormone Signalling and Secondary Cell Wall Formation When Constitutively Expressed in Rice. Scientific Reports, 2017, 7, 4535.	1.6	18
11	Action of multiple intra-QTL genes concerted around a co-localized transcription factor underpins a large effect QTL. Scientific Reports, 2015, 5, 15183.	1.6	58
12	Cereal flag leaf adaptations for grain yield under drought: knowledge status and gaps. Molecular Breeding, 2013, 31, 749-766.	1.0	70
13	An Absolute Index (Ab-index) to Measure a Researcher's Useful Contributions and Productivity. PLoS ONE, 2013, 8, e84334.	1.1	33
14	Development and validation of class I SSR markers targeting (GATA) n repeat motifs in rice. Euphytica, 2009, 169, 263-271.	0.6	7
15	In silico analysis of microsatellites in organellar genomes of major cereals for understanding their phylogenetic relationships. In Silico Biology, 2008, 8, 87-104.	0.4	18
16	Simple sequence repeats in organellar genomes of rice: frequency and distribution in genic and intergenic regions. Bioinformatics, 2007, 23, 1-4.	1.8	99
17	A Mitochondrial Repeat Specific Marker for Distinguishing Wild Abortive Type Cytoplasmic Male Sterile Rice Lines from their Cognate Isogenic Maintainer Lines. Crop Science, 2007, 47, 207-211.	0.8	24