N Manikanda Boopathi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8911697/publications.pdf

Version: 2024-02-01

40 papers 489

7 h-index 18 g-index

46 all docs 46 docs citations

46 times ranked

601 citing authors

#	Article	IF	CITATIONS
1	Phenotypic and genotypic analysis of drought-resistance traits for development of rice cultivars adapted to rainfed environments. Field Crops Research, 2008, 109, 1-23.	5.1	265
2	Molecular mapping and location of QTLs for drought-resistance traits in indica rice (Oryza sativa L.) lines adapted to target environments. Acta Physiologiae Plantarum, 2010, 32, 355-364.	2.1	52
3	Farmers' perceptions and management of plant viruses in vegetables and legumes in tropical and subtropical Asia. Crop Protection, 2015, 75, 115-123.	2.1	43
4	Genetic Mapping and Marker Assisted Selection. , 2020, , .		17
5	Marker-Assisted Breeding as Next-Generation Strategy for Genetic Improvement of Productivity and Quality: Can It Be Realized in Cotton?. International Journal of Plant Genomics, 2011, 2011, 1-16.	2.2	14
6	Evaluation and Bulked Segregant Analysis of Major Yield QTL qtl12.1 Introgressed into Indigenous Elite Line for Low Water Availability under Water Stress. Rice Science, 2013, 20, 25-30.	3.9	12
7	Genetic purity analysis of cotton (Gossypium spp.) hybrids using SSR markers. Seed Science and Technology, 2010, 38, 358-366.	1.4	11
8	GBS-SNP and SSR based genetic mapping and QTL analysis for drought tolerance in upland cotton. Physiology and Molecular Biology of Plants, 2021, 27, 1731-1745.	3.1	11
9	Plant miRNomics: Novel Insights in Gene Expression and Regulation. , 2015, , 181-211.		7
10	Association analysis of yield and fibre quality characters in interspecific population of cotton (Gossypium spp.). Journal of Crop Science and Biotechnology, 2012, 15, 239-243.	1.5	6
11	Genetic Diversity, Erosion, and Population Structure in Cotton Genetic Resources. Sustainable Development and Biodiversity, 2016, , 409-438.	1.7	6
12	Genetic improvement of a neglected and underutilised oilseed crop: sesame (Sesamum indicum L.) through mutation breeding. Nucleus (India), 2020, 63, 293-302.	2.2	6
13	Molecular Breeding for Genetic Improvement of Cotton (Gossypium spp.)., 2015,, 613-645.		5
14	Success Stories in MAS. , 2013, , 187-192.		3
15	Genotyping of Mapping Population. , 2013, , 39-80.		3
16	Usefulness and Utilization of Indian Cotton Germplasm. , 0, , .		3
17	Botanical Descriptions of Moringa spp.,. Compendium of Plant Genomes, 2021, , 11-20.	0.5	3
18	Linkage Map Construction. , 2013, , 81-108.		2

#	Article	IF	CITATIONS
19	Leapfrogging the Interventions of ICT Tools in Vegetable Intensification Pathways of Tamil Nadu: Farm Level Perceptions. Agricultural Economics Research Review, 2014, 27, 93.	0.2	2
20	Morphological characterization of certain Jasminum sambac genotypes using principal component analysis. The Pharma Innovation, 2021, 10, 118-123.	0.3	2
21	Molecular Markers and DNA Barcoding in Moringa. Compendium of Plant Genomes, 2021, , 85-100.	0.5	1
22	Tissue Culture and Genetic Engineering in Moringa. Compendium of Plant Genomes, 2021, , 67-83.	0.5	1
23	Profitability of begomovirus management strategies among chilli farmers in Tamil Nadu: A gross margin impact analysis. Indian Journal of Agricultural Research, 2015, , .	0.1	1
24	Correlation and path-coefficient analysis in Ramnad Mundu chilli (Capsicum annuum L.) for yield and quality traits. International Journal Plant Sciences, 2021, 16, 1-6.	0.0	1
25	Development of novel SSR markers derived from genomic and transcriptomic data of <i>Moringa oleifera</i> L. var. PKM1 and their applicability. Journal of Horticultural Science and Biotechnology, 2022, 97, 487-495.	1.9	1
26	Identification of Stable and Multiple Environment Interaction QTLs and Candidate Genes for Fiber Productive Traits Under Irrigated and Water Stress Conditions Using Intraspecific RILs of Gossypium hirsutum var. MCU5 X TCH1218. Frontiers in Plant Science, 2022, 13, 851504.	3.6	1
27	QTL Identification. , 2013, , 117-163.		O
28	Germplasm Characterisation: Utilising the Underexploited Resources. , 2013, , 1-21.		0
29	Curtain Raiser to Novel MAS Platforms. , 2013, , 193-244.		O
30	Recent Advances in MAS in Major Crops. , 2013, , 245-280.		О
31	Germplasm Characterization: Utilizing the Underexploited Resources. , 2020, , 21-68.		O
32	Cytogenetical Analysis of Moringa Genome. Compendium of Plant Genomes, 2021, , 51-55.	0.5	0
33	Genome Sequencing, Organellar Genomes and Comparative Genomics in Moringa. Compendium of Plant Genomes, 2021, , 101-132.	0.5	O
34	Medicinal and Therapeutic Properties of Moringa. Compendium of Plant Genomes, 2021, , 31-39.	0.5	0
35	Classical Genetics and Traditional Breeding in Moringa. Compendium of Plant Genomes, 2021, , 41-49.	0.5	O
36	Moringa and Its Genome: Future Prospects. Compendium of Plant Genomes, 2021, , 181-185.	0.5	0

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-	37	Moringa Functional Genomics: Implications of Long Read Sequencing Technologies. Compendium of Plant Genomes, 2021, , 133-143.	0.5	0
ć	38	In vitro Root Architectural Screening for Early Drought Adaptive Traits in Sorghum (Sorghum) Tj ETQq0 0 0 rgBT	Oyerlock	10 ₀ Tf 50 702
;	39	Transient albino and revertible to green (TARGreen) rice mutant: simple, affordable and beneficial novel tag for rapid genetic purity testing in rice (Oryza sativa L.). Genetic Resources and Crop Evolution, 2020, 67, 1949-1955.	1.6	0
4	40	Effect of Chemical Manipulation on Growth, Yield and Fiber Traits of Compact Cotton. Current Journal of Applied Science and Technology, 0, , 61-69.	0.3	0