Sajeevan Radha Sivarajan

List of Publications by Year in descending order

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18 papers	382 citations	933447 10 h-index	17 g-index
19	19	19	486
all docs	docs citations	times ranked	citing authors

#	Article	lF	CITATIONS
1	Multiple hybrid de novo genome assembly of finger millet, an orphan allotetraploid crop. DNA Research, 2018, 25, 39-47.	3.4	85
2	Expression of Arabidopsis SHN1 in Indian Mulberry (Morus indica L.) Increases Leaf Surface Wax Content and Reduces Post-harvest Water Loss. Frontiers in Plant Science, 2017, 8, 418.	3.6	41
3	Development and Characterization of Genic SSR Markers from Indian Mulberry Transcriptome and Their Transferability to Related Species of Moraceae. PLoS ONE, 2016, 11, e0162909.	2.5	41
4	An Approach to Function Annotation for Proteins of Unknown Function (PUFs) in the Transcriptome of Indian Mulberry. PLoS ONE, 2016, 11, e0151323.	2.5	40
5	Identification and Characterization of Genes Responsible for Drought Tolerance in Rice Mediated by Pseudomonas fluorescens. Rice Science, 2017, 24, 291-298.	3.9	35
6	The transcriptome enables the identification of candidate genes behind medicinal value of Drumstick tree (Moringa oleifera). Genomics, 2020, 112, 621-628.	2.9	22
7	An Efficient Protocol for Total RNA Isolation from Healthy and Stressed Tissues of Mulberry (<i>Morus</i> sp.) and Other Species. American Journal of Plant Sciences, 2014, 05, 2057-2065.	0.8	20
8	Full-Length Cloning and Characterization of Abiotic Stress Responsive & lt; i> CIPK31-Like< li> Gene from Finger Millet, a Drought-Tolerant Crop. Current Science, 2016, 111, 890.	0.8	19
9	Distinct Evolutionary Origins of Intron Retention Splicing Events in NHX1 Antiporter Transcripts Relate to Sequence Specific Distinctions in Oryza Species. Frontiers in Plant Science, 2020, 11, 267.	3.6	16
10	<i>In Vitro</i> Plant Regeneration of <i>Morus indica</i> L. cv. V1 Using Leaf Explant. American Journal of Plant Sciences, 2013, 04, 2001-2005.	0.8	16
11	Identification of splice variant of OsGBF1 in Oryza sativa ssp. indica genotypes under salinity stress. 3 Biotech, 2018, 8, 345.	2.2	11
12	Molecular cloning and characterization of a novel basic helix–loop–helix-144 (bHLH144)-like transcription factor from Morus alba (L.). Plant Gene, 2016, 5, 109-117.	2.3	10
13	Leaf wax trait in crops for drought and biotic stress tolerance: regulators of epicuticular wax synthesis and role of small RNAs. Indian Journal of Plant Physiology, 2017, 22, 434-447.	0.8	9
14	Identification and Characterization of OsWRKY72 Variant in Indica Genotypes. Rice Science, 2016, 23, 297-305.	3.9	6
15	A knowledge-driven protocol for prediction of proteins of interest with an emphasis on biosynthetic pathways. MethodsX, 2020, 7, 101053.	1.6	4
16	Dataset for the combined transcriptome assembly of M. oleifera and functional annotation. Data in Brief, 2020, 30, 105416.	1.0	4
17	Molecular Identification and Genetic Diversity of Lactobacillus Species Isolated from Different Edible Sources. Journal of Pure and Applied Microbiology, 2016, 10, 3155-3162.	0.9	2
18	Inhibition of ABA-mediated Responses by Dithiothreitol in Plants. Journal of Plant Growth Regulation, 0, , 1.	5.1	0