

Ujjal Jyoti phukan

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

1,243
citations

933447

10
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

1834
citing authors

#	ARTICLE	IF	CITATIONS
1	WRKY Transcription Factors: Molecular Regulation and Stress Responses in Plants. <i>Frontiers in Plant Science</i> , 2016, 7, 760.	3.6	598
2	Regulation of Apetala2/Ethylene Response Factors in Plants. <i>Frontiers in Plant Science</i> , 2017, 8, 150.	3.6	128
3	Waterlogging and submergence stress: affects and acclimation. <i>Critical Reviews in Biotechnology</i> , 2016, 36, 956-966.	9.0	84
4	Wound Induced Transcriptional Regulation of Benzylisoquinoline Pathway and Characterization of Wound Inducible PsWRKY Transcription Factor from <i>Papaver somniferum</i> . <i>PLoS ONE</i> , 2013, 8, e52784.	2.5	80
5	MaRAP4, a waterlogging-responsive ERF from <i>Mentha</i> , regulates bidirectional sugar transporter <i>AtSWEET10</i> to modulate stress response in <i>Arabidopsis</i> . <i>Plant Biotechnology Journal</i> , 2018, 16, 221-233.	8.3	70
6	De novo leaf and root transcriptome analysis identified novel genes involved in Steroidal sapogenin biosynthesis in <i>Asparagus racemosus</i> . <i>BMC Genomics</i> , 2014, 15, 746.	2.8	68
7	Cell Death in Plant Immunity. <i>Cold Spring Harbor Perspectives in Biology</i> , 2020, 12, a036483.	5.5	67
8	PsAP2 an AP2/ERF family transcription factor from <i>Papaver somniferum</i> enhances abiotic and biotic stress tolerance in transgenic tobacco. <i>Plant Molecular Biology</i> , 2015, 89, 173-186.	3.9	66
9	<i>Mentha arvensis</i> exhibit better adaptive characters in contrast to <i>Mentha piperita</i> when subjugated to sustained waterlogging stress. <i>Protoplasma</i> , 2014, 251, 603-614.	2.1	30
10	Interaction of host and <i>Staphylococcus aureus</i> protease-system regulates virulence and pathogenicity. <i>Medical Microbiology and Immunology</i> , 2019, 208, 585-607.	4.8	30
11	Robust transcriptional indicators of immune cell death revealed by spatiotemporal transcriptome analyses. <i>Molecular Plant</i> , 2022, 15, 1059-1075.	8.3	17