## Ram Singh Purty

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Physiological responses among Brassica species under salinity stress show strong correlation with transcript abundance for SOS pathway-related genes. Journal of Plant Physiology, 2009, 166, 507-520.	3.5	120
2	Towards salinity tolerance in Brassica: an overview. Physiology and Molecular Biology of Plants, 2008, 14, 39-49.	3.1	81
3	Transformation of tomato cultivar â€ <sup>~</sup> Pusa Ruby' with bspA gene from Populus tremula for drought tolerance. Plant Cell, Tissue and Organ Culture, 2006, 84, 56-68.	2.3	37
4	Optimization of De Novo Short Read Assembly of Seabuckthorn (Hippophae rhamnoides L.) Transcriptome. PLoS ONE, 2013, 8, e72516.	2.5	36
5	In silico identification and validation of miRNA and their DIR specific targets in Oryza sativa Indica under abiotic stress. Non-coding RNA Research, 2020, 5, 167-177.	4.6	21
6	Maintenance of stress related transcripts in tolerant cultivar at a level higher than sensitive one appears to be a conserved salinity response among plants. Plant Signaling and Behavior, 2009, 4, 431-434.	2.4	15
7	Assessment of Bioremediation Potential of Cellulosimicrobium sp. for Treatment of Multiple Heavy Metals. Microbiology and Biotechnology Letters, 2019, 47, 269-277.	0.4	14
8	Downregulation of Candidate Gene Expression and Neuroprotection by Piperine in Streptozotocin-Induced Hyperglycemia and Memory Impairment in Rats. Frontiers in Pharmacology, 2020, 11, 595471.	3.5	12
9	Induction of a novel boiling stable protein in response to desiccation and ABA treatments in Sesbania sesban var. bicolor leaves. Biologia Plantarum, 2005, 49, 137-140.	1.9	7
10	Structural and Expression Analysis of Salinity Stress Responsive Phosphoserine Phosphatase from Brassica juncea (L.). Journal of Proteomics and Bioinformatics, 2017, 10, .	0.4	7
11	Stress mitigation strategies of plant growth-promoting rhizobacteria: Plant growth-promoting rhizobacteria mechanisms. Plant Science Today, 2021, 8, 25-32.	0.7	5
12	Promoterless gus gene shows leaky β-glucuronidase activity during transformation of tomato with bspA gene for drought tolerance. Biologia Plantarum, 2006, 50, 352-358.	1.9	4
13	Process Design For Removal of Heavy Metals By A Bio-sorbent Trickle Bed System: A Proof of Concept. Journal of Physics: Conference Series, 2020, 1531, 012119.	0.4	4
14	Emperical Modeling of Growth Parameters in Cellulosimicrobium cellulans during Heavy Metal Tolerance. Journal of Physics: Conference Series, 2020, 1531, 012120.	0.4	4
15	Development of male sterile transgenic lines in rice by tapetum specific expression of <i>barnase</i> gene. Journal of Plant Biotechnology, 2017, 44, 364-371.	0.4	4
16	COVID‑19 pandemic: Understanding the emergence, pathogenesis and containment (Review). World Academy of Sciences Journal, 0, , .	0.6	3
17	Isolation, Characterization, and In Silico Interaction Studies of Bioactive Compounds from Caesalpinia bonducella with Target Proteins Involved in Alzheimer's Disease. Applied Biochemistry and Biotechnology, 2023, 195, 2216-2234.	2.9	3
18	Comparative analysis of the Agrobacterium mediated transformation using primary and secondary callus of indica rice (Oryza sativa L.) using phosphinothricin as selecting medium. Australian Journal of Crop Science, 2018, 12, 1660-1667.	0.3	2

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#	Article	IF	CITATIONS
19	Genome-wide analysis of PHD finger gene family and identification of potential miRNA and their PHD finger gene specific targets in Oryza sativa indica. Non-coding RNA Research, 2020, 5, 191-200.	4.6	2
20	Nature of the tapetum-specific promoter is crucial for generation of rice transgenics possessing a lethal barnase gene. Journal of Crop Science and Biotechnology, 2021, 24, 579-587.	1.5	2
21	Isolation of Cellulose-Degrading Bacteria and to Use Their Cellulolytic Potential for Production of Bioethanol from Paper Waste. Lecture Notes in Bioengineering, 2021, , 3-11.	0.4	0
22	Synthesis and Characterization of TiO2 Nanoparticle and Checking Its Antimicrobial Activity Against Escherichia coli and Staphylococcus aureus. Lecture Notes in Bioengineering, 2021, , 317-325.	0.4	0