

Pradeep K Agarwal

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

3,279
citations

218677

26
h-index

161849

54
g-index

58
all docs

58
docs citations

58
times ranked

3721
citing authors

#	ARTICLE	IF	CITATIONS
1	Role of DREB transcription factors in abiotic and biotic stress tolerance in plants. <i>Plant Cell Reports</i> , 2006, 25, 1263-1274.	5.6	864
2	Improved Salinity Tolerance of <i>Arachis hypogaea</i> (L.) by the Interaction of Halotolerant Plant-Growth-Promoting Rhizobacteria. <i>Journal of Plant Growth Regulation</i> , 2012, 31, 195-206.	5.1	256
3	Proteomics, metabolomics, and ionomics perspectives of salinity tolerance in halophytes. <i>Frontiers in Plant Science</i> , 2015, 6, 537.	3.6	226
4	Bioengineering for Salinity Tolerance in Plants: State of the Art. <i>Molecular Biotechnology</i> , 2013, 54, 102-123.	2.4	220
5	Overexpression of PgDREB2A transcription factor enhances abiotic stress tolerance and activates downstream stress-responsive genes. <i>Molecular Biology Reports</i> , 2010, 37, 1125-1135.	2.3	153
6	The <i>SbSOS1</i> gene from the extreme halophyte <i>Salicornia brachiata</i> enhances Na ⁺ loading in xylem and confers salt tolerance in transgenic tobacco. <i>BMC Plant Biology</i> , 2012, 12, 188.	3.6	147
7	Dehydration responsive element binding transcription factors and their applications for the engineering of stress tolerance. <i>Journal of Experimental Botany</i> , 2017, 68, 2135-2148.	4.8	144
8	Stress-inducible DREB2A transcription factor from <i>Pennisetum glaucum</i> is a phosphoprotein and its phosphorylation negatively regulates its DNA-binding activity. <i>Molecular Genetics and Genomics</i> , 2007, 277, 189-198.	2.1	106
9	Accumulation of heavy metals and its biochemical responses in <i>Salicornia brachiata</i> , an extreme halophyte. <i>Marine Biology Research</i> , 2010, 6, 511-518.	0.7	78
10	<i>SbDREB2A</i> , an A-2 type DREB transcription factor from extreme halophyte <i>Salicornia brachiata</i> confers abiotic stress tolerance in <i>Escherichia coli</i> . <i>Plant Cell Reports</i> , 2010, 29, 1131-1137.	5.6	72
11	Constitutive overexpression of a stress-inducible small GTP-binding protein PgRab7 from <i>Pennisetum glaucum</i> enhances abiotic stress tolerance in transgenic tobacco. <i>Plant Cell Reports</i> , 2007, 27, 105-115.	5.6	65
12	Identification of salt-induced genes from <i>Salicornia brachiata</i> , an extreme halophyte through expressed sequence tags analysis. <i>Genes and Genetic Systems</i> , 2009, 84, 111-120.	0.7	61
13	Cloning and characterization of the <i>Salicornia brachiata</i> Na ⁺ /H ⁺ antiporter gene <i>SbNHX1</i> and its expression by abiotic stress. <i>Molecular Biology Reports</i> , 2011, 38, 1965-1973.	2.3	54
14	The <i>SbASR-1</i> Gene Cloned from an Extreme Halophyte <i>Salicornia brachiata</i> Enhances Salt Tolerance in Transgenic Tobacco. <i>Marine Biotechnology</i> , 2012, 14, 782-792.	2.4	50
15	Ion homeostasis in a salt-secreting halophytic grass. <i>AoB PLANTS</i> , 2015, 7, plv055.	2.3	50
16	A Dehydration-Responsive Element Binding (DREB) Transcription Factor from the Succulent Halophyte <i>Salicornia brachiata</i> Enhances Abiotic Stress Tolerance in Transgenic Tobacco. <i>Marine Biotechnology</i> , 2014, 16, 657-673.	2.4	44
17	Molecular characterization of the <i>Salicornia brachiata</i> <i>SbMAPKK</i> gene and its expression by abiotic stress. <i>Molecular Biology Reports</i> , 2010, 37, 981-986.	2.3	43
18	PCIB an Antiauxin Enhances Microspore Embryogenesis in Microspore Culture of <i>Brassica juncea</i> . <i>Plant Cell, Tissue and Organ Culture</i> , 2006, 86, 201-210.	2.3	42

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19	Overexpression of a novel SbMYB15 from <i>Salicornia brachiata</i> confers salinity and dehydration tolerance by reduced oxidative damage and improved photosynthesis in transgenic tobacco. <i>Planta</i> , 2015, 242, 1291-1308.	3.2	41
20	<i>Kappaphycus alvarezii</i> sap mitigates abiotic-induced stress in <i>Triticum durum</i> by modulating metabolic coordination and improves growth and yield. <i>Journal of Applied Phycology</i> , 2018, 30, 2659-2673.	2.8	40
21	Insights into the role of seaweed <i>Kappaphycus alvarezii</i> sap towards phytohormone signalling and regulating defence responsive genes in <i>Lycopersicon esculentum</i> . <i>Journal of Applied Phycology</i> , 2016, 28, 2529-2537.	2.8	38
22	Molecular characterization of an MYB transcription factor from a succulent halophyte involved in stress tolerance. <i>AoB PLANTS</i> , 2015, 7, plv054.	2.3	35
23	Plant Rabs: Characterization, Functional Diversity, and Role in Stress Tolerance. <i>Plant Molecular Biology Reporter</i> , 2009, 27, 417-430.	1.8	32
24	A SNARE-Like Superfamily Protein SbSLSP from the Halophyte <i>Salicornia brachiata</i> Confers Salt and Drought Tolerance by Maintaining Membrane Stability, K ⁺ /Na ⁺ Ratio, and Antioxidant Machinery. <i>Frontiers in Plant Science</i> , 2016, 7, 737.	3.6	30
25	Ectopic Expression of JcWRKY Transcription Factor Confers Salinity Tolerance via Salicylic Acid Signaling. <i>Frontiers in Plant Science</i> , 2016, 7, 1541.	3.6	29
26	Somatic embryogenesis and in vitro plantlet regeneration in <i>Salicornia brachiata</i> Roxb.. <i>Plant Cell, Tissue and Organ Culture</i> , 2015, 120, 355-360.	2.3	26
27	SbMYB15 transcription factor mitigates cadmium and nickel stress in transgenic tobacco by limiting uptake and modulating antioxidative defence system. <i>Functional Plant Biology</i> , 2019, 46, 702.	2.1	26
28	Molecular Cloning and Characterization of a Group II WRKY Transcription Factor from <i>Jatropha curcas</i> , an Important Biofuel Crop. <i>DNA and Cell Biology</i> , 2014, 33, 503-513.	1.9	25
29	A Low-Affinity K ⁺ Transporter ALHKT2;1 from Recretohalophyte <i>Aeluropus lagopoides</i> Confers Salt Tolerance in Yeast. <i>Molecular Biotechnology</i> , 2015, 57, 489-498.	2.4	24
30	Deciphering hydrogen peroxide-induced signalling towards stress tolerance in plants. <i>3 Biotech</i> , 2019, 9, 395.	2.2	23
31	Seaweed extracts: Potential biodegradable, environmentally friendly resources for regulating plant defence. <i>Algal Research</i> , 2021, 58, 102363.	4.6	23
32	Improved Shoot Regeneration, Salinity Tolerance and Reduced Fungal Susceptibility in Transgenic Tobacco Constitutively Expressing PR-10a Gene. <i>Frontiers in Plant Science</i> , 2016, 7, 217.	3.6	22
33	Chemical Derivatization of Metabolite Mass Profiling of the Recretohalophyte <i>Aeluropus lagopoides</i> Revealing Salt Stress Tolerance Mechanism. <i>Marine Biotechnology</i> , 2017, 19, 207-218.	2.4	19
34	The JcWRKY tobacco transgenics showed improved photosynthetic efficiency and wax accumulation during salinity. <i>Scientific Reports</i> , 2019, 9, 19617.	3.3	18
35	AINAC4 Transcription Factor From Halophyte <i>Aeluropus lagopoides</i> Mitigates Oxidative Stress by Maintaining ROS Homeostasis in Transgenic Tobacco. <i>Frontiers in Plant Science</i> , 2018, 9, 1522.	3.6	17
36	A novel salt-inducible gene SbSI-1 from <i>Salicornia brachiata</i> confers salt and desiccation tolerance in <i>E. coli</i> . <i>Molecular Biology Reports</i> , 2012, 39, 1943-1948.	2.3	15

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37	High-frequency in vitro shoot regeneration in <i>Cucumis sativus</i> by inhibition of endogenous auxin. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 2014, 50, 729-737.	2.1	14
38	Sargassum seaweed extract enhances <i>Macrophomina phaseolina</i> resistance in tomato by regulating phytohormones and antioxidative activity. <i>Journal of Applied Phycology</i> , 2020, 32, 4373-4384.	2.8	14
39	Transcriptional regulation of salinity stress: role and spatio-temporal expressions of ion-transporter gene promoters. <i>Biologia Plantarum</i> , 2018, 62, 641-646.	1.9	10
40	Overexpression of <i>JcWRKY2</i> confers increased resistance towards <i>Macrophomina phaseolina</i> in transgenic tobacco. <i>3 Biotech</i> , 2020, 10, 490.	2.2	8
41	MSAP marker based DNA methylation study in <i>Salicornia brachiata</i> DREB2A transgenic tobacco. <i>Plant Gene</i> , 2016, 6, 77-81.	2.3	7
42	Ectopic Expression of <i>JcWRKY</i> Confers Enhanced Resistance in Transgenic Tobacco Against <i>Macrophomina phaseolina</i> . <i>DNA and Cell Biology</i> , 2018, 37, 298-307.	1.9	7
43	Functional Validation of <i>JcWRKY2</i> , a Group III Transcription Factor Toward Mitigating Salinity Stress in Transgenic Tobacco. <i>DNA and Cell Biology</i> , 2019, 38, 1278-1291.	1.9	7
44	Molecular cloning and characterization of high-affinity potassium transporter (<i>AlHKT2;1</i>) gene promoter from halophyte <i>Aeluropus lagopoides</i> . <i>International Journal of Biological Macromolecules</i> , 2021, 181, 1254-1264.	7.5	7
45	Mechanism of high affinity potassium transporter (HKT) towards improved crop productivity in saline agricultural lands. <i>3 Biotech</i> , 2022, 12, 51.	2.2	7
46	Artificial miRNA mediated resistance in tobacco against <i>Jatropha</i> leaf curl Gujarat virus by targeting RNA silencing suppressors. <i>Scientific Reports</i> , 2021, 11, 890.	3.3	5
47	Special Adaptive Features of Plant Species in Response to Salinity. <i>Signaling and Communication in Plants</i> , 2020, , 53-76.	0.7	5
48	An economical and efficient protocol for total RNA isolation from <i>Jatropha curcas</i> . <i>International Journal of Environmental Studies</i> , 2015, 72, 624-630.	1.6	4
49	Geminiviruses: Molecular biodiversity and global distribution in <i>Jatropha</i> . <i>Physiological and Molecular Plant Pathology</i> , 2019, 108, 101439.	2.5	4
50	Overexpression of <i>AINAC1</i> from recretohalophyte <i>Aeluropus lagopoides</i> alleviates drought stress in transgenic tobacco. <i>Environmental and Experimental Botany</i> , 2021, 181, 104277.	4.2	4
51	Development of a high-frequency adventitious shoot regeneration using cotyledon explants of an important oilseed crop <i>Sesamum indicum</i> L.. <i>In Vitro Cellular and Developmental Biology - Plant</i> , 2022, 58, 470-478.	2.1	4
52	Expression of <i>B. subtilis</i> Phytase gene driven by fruit specific E8 promoter for enhanced minerals, metabolites and phytonutrient in cucumber fruit. <i>Food Research International</i> , 2022, 156, 111138.	6.2	4
53	Transcription Factor-Based Genetic Engineering for Salinity Tolerance in Crops. , 2018, , 185-211.		3
54	<i>AlRab7</i> from <i>Aeluropus lagopoides</i> ameliorates ion toxicity in transgenic tobacco by regulating hormone signaling and reactive oxygen species homeostasis. <i>Physiologia Plantarum</i> , 2021, 173, 1448-1462.	5.2	3

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55	The AlRab7 E3-Ub-ligase mediates AlRab7 ubiquitination and improves ionic and oxidative stress tolerance in <i>Saccharomyces cerevisiae</i> . <i>Plant Physiology and Biochemistry</i> , 2020, 151, 689-704.	5.8	1
56	Soil microbial diversity shift as affected by conversion of shallow and rocky wastelands to <i>Jatropha curcas</i> L. plantation. <i>International Journal of Environmental Studies</i> , 2015, 72, 631-649.	1.6	0