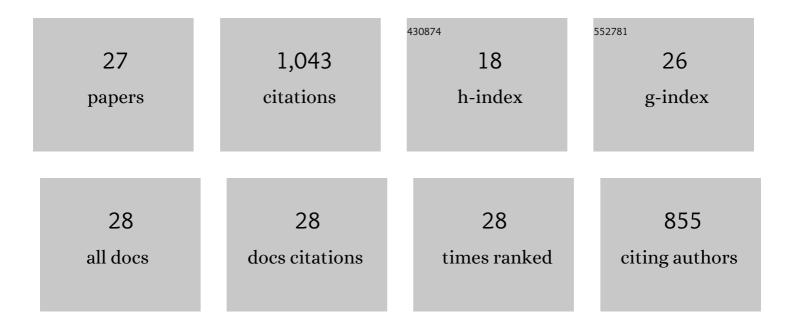
Prafull Salvi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcription factors as key molecular target to strengthen the drought stress tolerance in plants. Physiologia Plantarum, 2021, 172, 847-868.	5.2	131
2	Phytohormone signaling and crosstalk in regulating drought stress response in plants. Plant Cell Reports, 2021, 40, 1305-1329.	5.6	113
3	Differentially expressed seed aging responsive heat shock protein OsHSP18.2 implicates in seed vigor, longevity and improves germination and seedling establishment under abiotic stress. Frontiers in Plant Science, 2015, 6, 713.	3.6	103
4	Differentially expressed galactinol synthase(s) in chickpea are implicated in seed vigor and longevity by limiting the age induced ROS accumulation. Scientific Reports, 2016, 6, 35088.	3.3	76
5	Stress-Inducible Galactinol Synthase of Chickpea (CaGolS) is Implicated in Heat and Oxidative Stress Tolerance Through Reducing Stress-Induced Excessive Reactive Oxygen Species Accumulation. Plant and Cell Physiology, 2018, 59, 155-166.	3.1	76
6	Imperative role of sugar signaling and transport during drought stress responses in plants. Physiologia Plantarum, 2021, 171, 833-848.	5.2	73
7	Rice PROTEIN <scp>l</scp> â€ISOASPARTYL METHYLTRANSFERASE isoforms differentially accumulate during seed maturation to restrict deleterious isoAsp and reactive oxygen species accumulation and are implicated in seed vigor and longevity. New Phytologist, 2016, 211, 627-645.	7.3	63
8	Differentially expressed myo-inositol monophosphatase gene (CaIMP) in chickpea (Cicer arietinum L.) encodes a lithium-sensitive phosphatase enzyme with broad substrate specificity and improves seed germination and seedling growth under abiotic stresses. Journal of Experimental Botany, 2013, 64, 5623-5639.	4.8	56
9	Ectopic overexpression of cytosolic ascorbate peroxidase gene (Apx1) improves salinity stress tolerance in Brassica juncea by strengthening antioxidative defense mechanism. Acta Physiologiae Plantarum, 2020, 42, 1.	2.1	37
10	Exploration of silicate solubilizing bacteria for sustainable agriculture and silicon biogeochemical cycle. Plant Physiology and Biochemistry, 2021, 166, 827-838.	5.8	36
11	Ectopic over-expression of ABA-responsive Chickpea galactinol synthase (CaGolS) gene results in improved tolerance to dehydration stress by modulating ROS scavenging. Environmental and Experimental Botany, 2020, 171, 103957.	4.2	34
12	Arabidopsis SKP1-like protein13 (ASK13) positively regulates seed germination and seedling growth under abiotic stress. Journal of Experimental Botany, 2018, 69, 3899-3915.	4.8	33
13	Biological potential of bioactive metabolites derived from fungal endophytes associated with medicinal plants. Mycological Progress, 2021, 20, 577-594.	1.4	32
14	Sugar transporters and their molecular tradeoffs during abiotic stress responses in plants. Physiologia Plantarum, 2022, 174, e13652.	5.2	31
15	Insertional Mutagenesis Approaches and Their Use in Rice for Functional Genomics. Plants, 2019, 8, 310.	3.5	25
16	Assessment of Biological Activities of Fungal Endophytes Derived Bioactive Compounds Isolated from Amoora rohituka. Journal of Fungi (Basel, Switzerland), 2022, 8, 285.	3.5	24
17	Molecular cloning, in-silico characterization and functional validation of monodehydroascorbate reductase gene in Eleusine coracana. PLoS ONE, 2017, 12, e0187793.	2.5	21
18	Arabidopsis protein l-ISOASPARTYL METHYLTRANSFERASE repairs isoaspartyl damage to antioxidant enzymes and increases heat and oxidative stress tolerance. Journal of Biological Chemistry, 2020, 295, 783-799.	3.4	20

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19	<i>Arabidopsis</i> protein l-ISOASPARTYL METHYLTRANSFERASE repairs isoaspartyl damage to antioxidant enzymes and increases heat and oxidative stress tolerance. Journal of Biological Chemistry, 2020, 295, 783-799.	3.4	16
20	Antimicrobial Potential of Essential Oils from Aromatic Plant Ocimum sp.; A Comparative Biochemical Profiling and In-Silico Analysis. Agronomy, 2022, 12, 627.	3.0	13
21	Efficient Genetic Transformation of Rice for CRISPR/Cas9 Mediated Genome-Editing and Stable Overexpression Studies: A Case Study on Rice Lipase 1 and Galactinol Synthase Encoding Genes. Agronomy, 2022, 12, 179.	3.0	9
22	Deciphering the structural basis of the broad substrate specificity of myo-inositol monophosphatase (IMP) from Cicer arietinum. International Journal of Biological Macromolecules, 2020, 151, 967-975.	7.5	6
23	A rapid, efficient, and low-cost BiFC protocol and its application in studying in vivo interaction of seed-specific transcription factors, RISBZ and RPBF. Functional and Integrative Genomics, 2021, 21, 593-603.	3.5	6
24	A conserved NAG motif is critical to the catalytic activity of galactinol synthase, a key regulatory enzyme of RFO biosynthesis. Biochemical Journal, 2021, 478, 3939-3955.	3.7	5
25	Expression of ECMYB Transcription Factor Gene Under Different Abiotic Stress Conditions in Eleusine coracana. International Journal of Agriculture Environment and Biotechnology, 2018, 11, .	0.1	3
26	Gateway cloning and in-planta transformation of drought stress responsive Ecmyb1 gene isolated from Eleusine coracana var.PRM 6107. Environment Conservation Journal, 2021, 22, 205-211.	0.2	1
27	Stress ResponsiveÂOshyprp16ÂPromoter Driven Early Expression of Resistance GeneÂPi54ÂPotentiate the Resistance AgainstÂMagnaporthe OryzaeÂIn Transgenic Rice. SSRN Electronic Journal, 0, , .	0.4	Ο