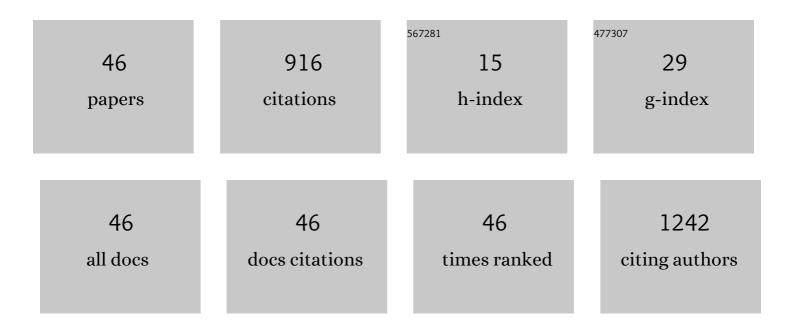
Dev Mani Pandey

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Biochemistry and biosynthesis of insect pigments. European Journal of Entomology, 2014, 111, 149-164.	1.2	130
2	Molecular marker survey and expression analyses of the rice submergence-tolerance gene SUB1A. Theoretical and Applied Genetics, 2010, 121, 1441-1453.	3.6	89
3	Identification of novel drought-responsive microRNAs and trans-acting siRNAs from Sorghum bicolor (L.) Moench by high-throughput sequencing analysis. Frontiers in Plant Science, 2015, 6, 506.	3.6	76
4	Transcriptional Regulatory Network Analysis of MYB Transcription Factor Family Genes in Rice. Frontiers in Plant Science, 2015, 6, 1157.	3.6	76
5	Effect of Growth Regulators on Photosynthesis, Transpiration and Related Parameters in Water Stressed Cotton. Biologia Plantarum, 2001, 44, 475-478.	1.9	49
6	ldentification of new stress-induced microRNA and their targets in wheat using computational approach. Plant Signaling and Behavior, 2013, 8, e23932.	2.4	44
7	Extraction and characterization of essential oil components based on geraniol and citronellol from Java citronella (Cymbopogon winterianus Jowitt). Plant Growth Regulation, 2014, 73, 133-145.	3.4	44
8	Identification of conserved drought stress responsive gene-network across tissues and developmental stages in rice. Bioinformation, 2013, 9, 72-78.	0.5	42
9	Identification of miRNAs in sorghum by using bioinformatics approach. Plant Signaling and Behavior, 2012, 7, 246-259.	2.4	38
10	Hormonal Regulation of Photosynthetic Enzymes in Cotton under Water Stress. Photosynthetica, 2000, 38, 403-407.	1.7	34
11	Effect of drought on physiological aspects of Crassulacean acid metabolism in Doritaenopsis. Plant Science, 2004, 167, 1219-1226.	3.6	29
12	Effect of carbon dioxide on cell growth and saponin production in suspension cultures of Panax ginseng. Biologia Plantarum, 2006, 50, 752-754.	1.9	22
13	ACC deaminase producing rhizobacterium Enterobacter cloacae ZNP-4 enhance abiotic stress tolerance in wheat plant. PLoS ONE, 2022, 17, e0267127.	2.5	19
14	Effect of Growth Regulators on Photosynthetic Metabolites in Cotton under Water Stress. Biologia Plantarum, 2002, 45, 445-448.	1.9	18
15	Gene network modules associated with abiotic stress response in tolerant rice genotypes identified by transcriptome meta-analysis. Functional and Integrative Genomics, 2020, 20, 29-49.	3.5	17
16	Anticancer activity of plant leaves extract collected from a tribal region of India. 3 Biotech, 2019, 9, 399.	2.2	16
17	Identification of miRNA, their targets and miPEPs in peanut (Arachis hypogaea L.). Computational Biology and Chemistry, 2019, 83, 107100.	2.3	16
18	Effects of excessive photon on the photosynthetic pigments and violaxanthin de-epoxidase activity in the xanthophyll cycle of spinach leaf. Plant Science, 2005, 168, 161-166.	3.6	13

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#	Article	IF	CITATIONS
19	Method development for optimised green synthesis of gold nanoparticles from <i>Millettia pinnata</i> and their activity in nonâ€small cell lung cancer cell lines. IET Nanobiotechnology, 2019, 13, 626-633.	3.8	13
20	Drought effect on electrophoretic protein pattern of Anoectochilus formosanus. Scientia Horticulturae, 2006, 107, 205-209.	3.6	12
21	Stress-induced degradation of D1 protein and its photoprotection by DCPIP in isolated thylakoid membranes of barley leaf. Biologia Plantarum, 2008, 52, 291-298.	1.9	12
22	ldentification and expression analysis of hypoxia stress inducible CCCH-type zinc finger protein genes in rice. Journal of Plant Biology, 2012, 55, 489-497.	2.1	12
23	Identification and annotation of abiotic stress responsive candidate genes in peanut ESTs. Bioinformation, 2012, 8, 1211-1219.	0.5	10
24	Study on cocoonase, sericin, and degumming of silk cocoon: computational and experimental. Journal of Genetic Engineering and Biotechnology, 2021, 19, 32.	3.3	10
25	The molecular docking and molecular dynamics study of flavonol synthase and flavonoid 3'-monooxygenase enzymes involved for the enrichment of kaempferol. Journal of Biomolecular Structure and Dynamics, 2023, 41, 2478-2491.	3.5	10
26	Lac dye as a potential anti-neoplastic agent. Journal of Cancer Research and Therapeutics, 2016, 12, 1033.	0.9	8
27	Molecular Dynamics Simulation of Rap1 Myb-type domain in Saccharomyces cerevisiae. Bioinformation, 2012, 8, 881-885.	0.5	8
28	Photosystem 2-activity and thylakoid membrane polypeptides of in vitro cultured chrysanthemum as affected by NaCl. Biologia Plantarum, 2009, 53, 329-333.	1.9	7
29	Identification of genes related to resin biosynthesis in the Indian lac insect, Kerria lacca (Hemiptera:) Tj ETQq1	1 0.784314 1.0	ŀrg₿T /Overl⊂
30	Effects of different irradiances on the photosynthetic process during ex-vitro acclimation of Anoectochilus plantlets. Photosynthetica, 2006, 44, 419-424.	1.7	5
31	In silico analysis of motifs in promoters of Differentially Expressed Genes in rice (Oryza sativa L.) under anoxia. International Journal of Bioinformatics Research and Applications, 2009, 5, 525.	0.2	5
32	Dynamic Changes of Photosynthetic Pigments in Soybean Callus under High Irradiance. Photosynthetica, 2003, 41, 311-314.	1.7	4
33	In Silico Characterization and Analysis of RTBP1 and NgTRF1 Protein Through MD Simulation and Molecular Docking: A Comparative Study. Interdisciplinary Sciences, Computational Life Sciences, 2015, 7, 275-286.	3.6	3
34	Protein–protein docking and molecular dynamics studies of sericin and cocoonase of silkworm: an insight for cocoon softening. Journal of Biomolecular Structure and Dynamics, 2021, , 1-13.	3.5	3
35	High Irradiance Effects on the Xanthophyll Cycle Pigments and the Activity of Violaxanthin De-Epoxidase in Soybean Callus. Photosynthetica, 2004, 42, 153-156.	1.7	2
36	Long-term effects of growth regulators on growth and turnover of symplastic and apoplastic sugars in the suspension subculture of kidney bean. Journal of Plant Biology, 2004, 47, 21-26.	2.1	2

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#	Article	IF	CITATIONS
37	Identification of GCC-box and TCC-box motifs in the promoters of differentially expressed genes in rice (Oryza sativa L.): Experimental and computational approaches. PLoS ONE, 2019, 14, e0214964.	2.5	2
38	Microstructural changes in rice (Oryza sativa L.) leaves under varying low pH levels: a swept-source optical coherence tomography approach. Laser Physics, 2021, 31, 085601.	1.2	2
39	SVM based model generation for binding site prediction on helix turn helix motif type of transcription factors in eukaryotes. Bioinformation, 2013, 9, 500-505.	0.5	2
40	In silico structural and functional characterization of Antheraea mylitta cocoonase. Journal of Genetic Engineering and Biotechnology, 2022, 20, 102.	3.3	2
41	Identification of single exon genes and their encoded proteins in rice (Oryza sativa L.) genome: an in silico approach. International Journal of Bioinformatics Research and Applications, 2011, 7, 376.	0.2	1
42	In silico characterization and analysis of RTBP1 and NgTRF1 protein through MD simulation and molecular docking — A comparative study. Interdisciplinary Sciences, Computational Life Sciences, 2015, 7, 275.	3.6	1
43	Annotation of Stress-Responsive Candidate Genes in Peanut ESTs. Interdisciplinary Sciences, Computational Life Sciences, 2015, 7, 143-151.	3.6	1
44	Changes in contents of photosynthetic pigments and ribulose-1,5-bisphosphate carboxylase activity during the development of globular somatic embryo into the plantlet of Siberian ginseng. Photosynthetica, 2006, 44, 221-226.	1.7	0
45	Molecular modeling and dynamics study of nonsynonymous SNP in bread wheat HSP16.9B gene. , 2016, , \cdot		0
46	Co-expression network analysis of acidic-responsive genes in Arabidopsis thaliana signifies hub genes expression and their key role assessment for acidity tolerance in Oryza sativa L. Biologia (Poland), 2021, 76, 3175-3190.	1.5	0