Ashutosh Pandey

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4798391/publications.pdf

Version: 2024-02-01

43 papers 2,209 citations

236833 25 h-index 289141 40 g-index

45 all docs

45 docs citations

45 times ranked

2497 citing authors

#	Article	IF	CITATIONS
1	Comparative transcriptome analysis of arsenate and arsenite stresses in rice seedlings. Chemosphere, 2009, 74, 688-702.	4.2	254
2	Modulation of Transcriptome and Metabolome of Tobacco by Arabidopsis Transcription Factor, <i>AtMYB12</i> , Leads to Insect Resistance. Plant Physiology, 2010, 152, 2258-2268.	2.3	216
3	MicroRNA858 Is a Potential Regulator of Phenylpropanoid Pathway and Plant Development. Plant Physiology, 2016, 171, 944-959.	2.3	163
4	CRISPR/Cas9 directed editing of lycopene epsilon-cyclase modulates metabolic flux for \hat{l}^2 -carotene biosynthesis in banana fruit. Metabolic Engineering, 2020, 59, 76-86.	3.6	144
5	Expression of Arabidopsis MYB transcription factor, AtMYB111, in tobacco requires light to modulate flavonol content. Scientific Reports, 2014, 4, 5018.	1.6	92
6	Isolation, screening and comprehensive characterization of candidate microalgae for biofuel feedstock production and dairy effluent treatment: A sustainable approach. Bioresource Technology, 2019, 293, 121998.	4.8	83
7	Coâ€expression of <i>Arabidopsis</i> transcription factor, <i>At<scp>MYB</scp>12</i> , and soybean isoflavone synthase, <i>Gm<scp>IFS</scp>1</i> , genes in tobacco leads to enhanced biosynthesis of isoflavones and flavonols resulting in osteoprotective activity. Plant Biotechnology Journal, 2014, 12, 69-80.	4.1	80
8	Constitutive expression of Arabidopsis MYB transcription factor, AtMYB11, in tobacco modulates flavonoid biosynthesis in favor of flavonol accumulation. Plant Cell Reports, 2015, 34, 1515-1528.	2.8	80
9	Genome-wide analysis of the AP2/ERF family in Musa species reveals divergence and neofunctionalisation during evolution. Scientific Reports, 2016, 6, 18878.	1.6	75
10	Molecular subtypes of colorectal cancer: AnÂemerging therapeutic opportunity for personalized medicine. Genes and Diseases, 2021, 8, 133-145.	1.5	71
11	Development of AtMYB12-expressing transgenic tobacco callus culture for production of rutin with biopesticidal potential. Plant Cell Reports, 2012, 31, 1867-1876.	2.8	66
12	AtMYB12 expression in tomato leads to large scale differential modulation in transcriptome and flavonoid content in leaf and fruit tissues. Scientific Reports, 2015, 5, 12412.	1.6	66
13	Genome-Wide Analysis of the Musa WRKY Gene Family: Evolution and Differential Expression during Development and Stress. Frontiers in Plant Science, 2016, 7, 299.	1.7	55
14	Low Temperature-Enhanced Flavonol Synthesis Requires Light-Associated Regulatory Components in Arabidopsis thaliana. Plant and Cell Physiology, 2018, 59, 2099-2112.	1.5	55
15	Genome-wide Expression Analysis and Metabolite Profiling Elucidate Transcriptional Regulation of Flavonoid Biosynthesis and Modulation under Abiotic Stresses in Banana. Scientific Reports, 2016, 6, 31361.	1.6	52
16	Nitric oxide alters nitrogen metabolism and PIN gene expressions by playing protective role in arsenic challenged Brassica juncea L Ecotoxicology and Environmental Safety, 2019, 176, 95-107.	2.9	48
17	Molecular components associated with the regulation of flavonoid biosynthesis. Plant Science, 2022, 317, 111196.	1.7	42
18	The R2R3-MYB gene family in banana (Musa acuminata): Genome-wide identification, classification and expression patterns. PLoS ONE, 2020, 15, e0239275.	1.1	39

#	Article	IF	CITATIONS
19	Development and cost-benefit analysis of a novel process for biofuel production from microalgae using pre-treated high-strength fresh cheese whey wastewater. Environmental Science and Pollution Research, 2020, 27, 23963-23980.	2.7	37
20	The R2R3-MYB transcription factor MtMYB134 orchestrates flavonol biosynthesis in Medicago truncatula. Plant Molecular Biology, 2021, 106, 157-172.	2.0	37
21	Multi-objective optimization of media nutrients for enhanced production of algae biomass and fatty acid biosynthesis from Chlorella pyrenoidosa NCIM 2738. Bioresource Technology, 2016, 200, 940-950.	4.8	36
22	Genome-Wide Identification and Expression Analysis of Homeodomain Leucine Zipper Subfamily IV (HDZ) Tj ETQc	10,00 rgB 1.7	T /Overlock
23	Characterization of isoflavone synthase gene from Psoralea corylifolia: a medicinal plant. Plant Cell Reports, 2010, 29, 747-755.	2.8	31
24	Regulation of Banana Phytoene Synthase (MaPSY) Expression, Characterization and Their Modulation under Various Abiotic Stress Conditions. Frontiers in Plant Science, 2017, 8, 462.	1.7	30
25	Ultrasound-intensified biodiesel production from algal biomass: a review. Environmental Chemistry Letters, 2021, 19, 209-229.	8.3	28
26	DEVELOPMENT AND OPTIMIZATION OF HPLC-PDA-MS-MS METHOD FOR SIMULTANEOUS QUANTIFICATION OF THREE CLASSES OF FLAVONOIDS IN LEGUME SEEDS, VEGETABLES, FRUITS, AND MEDICINAL PLANTS. Journal of Liquid Chromatography and Related Technologies, 2011, 34, 1729-1742.	0.5	27
27	Compendium of Plant-Specific CRISPR Vectors and Their Technical Advantages. Life, 2021, 11, 1021.	1.1	26
28	Genetically engineered flavonol enriched tomato fruit modulates chondrogenesis to increase bone length in growing animals. Scientific Reports, 2016, 6, 21668.	1.6	24
29	Protective role of nitric oxide on nitrogen-thiol metabolism and amino acids profiling during arsenic exposure in Oryza sativa L. Ecotoxicology, 2020, 29, 825-836.	1.1	24
30	COP1 mediates light-dependent regulation of flavonol biosynthesis through HY5 in Arabidopsis. Plant Science, 2021, 303, 110760.	1.7	23
31	Characterization and Expression Analysis of Phytoene Synthase from Bread Wheat (Triticum aestivum) Tj ETQq1	1 0.78431 1:1	4 rgBT /Ove
32	Molecular Characterization Revealed the Role of Thaumatin-Like Proteins of Bread Wheat in Stress Response. Frontiers in Plant Science, 2021, 12, 807448.	1.7	23
33	Emerging tools and paradigm shift of gene editing in cereals, fruits, and horticultural crops for enhancing nutritional value and food security. Food and Energy Security, 2021, 10, e258.	2.0	21
34	Harvesting of freshwater microalgae Scenedesmus sp. by electro–coagulation–flocculation for biofuel production: effects on spent medium recycling and lipid extraction. Environmental Science and Pollution Research, 2020, 27, 3497-3507.	2.7	20
35	MAPK cascade gene family in Camellia sinensis: In-silico identification, expression profiles and regulatory network analysis. BMC Genomics, 2020, 21, 613.	1.2	15
36	Novel microRNAs regulating ripening-associated processes in banana fruit. Plant Growth Regulation, 2020, 90, 223-235.	1.8	14

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#	Article	IF	CITATION
37	Molecular Characterization, Evolutionary Analysis, and Expression Profiling of BOR Genes in Important Cereals. Plants, 2022, 11, 911.	1.6	14
38	Interplay between <scp>R2R3 MYB</scp> â€type activators and repressors regulates proanthocyanidin biosynthesis in banana (<i>Musa acuminata</i>). New Phytologist, 2022, 236, 1108-1127.	3.5	14
39	Provitamin A Enrichment for Tackling Malnutrition. , 2016, , 277-299.		9
40	Scenedesmus sp. ASK22 cultivation using simulated dairy wastewater for nutrient sequestration and biofuel production: insight into fuel properties and their blends. Biomass Conversion and Biorefinery, 2024, 14, 3305-3317.	2.9	5
41	Two homeologous MATE transporter genes, <i>NtMATE21</i> and <i>NtMATE22</i> , are involved in the modulation of plant growth and flavonol transport in <i>Nicotiana tabacum</i> . Journal of Experimental Botany, 2022, 73, 6186-6206.	2.4	5
42	Synthetic Metabolism and Its Significance in Agriculture., 2019,, 365-391.		3
43	Targeted Metabolite Profiling of Five Cultivars of Vitis vinifera L. Fruits. ACS Food Science & Technology, 2021, 1, 653-659.	1.3	O