Lihui Zeng

List of Publications by Year
in descending order

18	461	840776 11	940533 16
papers	citations	h-index	g-index

18	18	18	380
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	StMYB44 negatively regulates anthocyanin biosynthesis at high temperatures in tuber flesh of potato. Journal of Experimental Botany, 2019, 70, 3809-3824.	4.8	95
2	The Critical Role of miRNAs in Regulation of Flowering Time and Flower Development. Genes, 2020, 11, 319.	2.4	59
3	NtMYB3, an R2R3-MYB from Narcissus, Regulates Flavonoid Biosynthesis. International Journal of Molecular Sciences, 2019, 20, 5456.	4.1	56
4	Ectopic Overexpression of a Novel R2R3-MYB, NtMYB2 from Chinese Narcissus Represses Anthocyanin Biosynthesis in Tobacco. Molecules, 2018, 23, 781.	3.8	50
5	The red flesh of kiwifruit is differentially controlled by specific activationâ€“repression systems. New Phytologist, 2022, 235, 630-645.	7.3	37
6	Identification of Regulatory Genes Implicated in Continuous Flowering of Longan (Dimocarpus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54	2.5	26
7	Recent Advanced Metabolic and Genetic Engineering of Phenylpropanoid Biosynthetic Pathways. International Journal of Molecular Sciences, 2021, 22, 9544.	4.1	22
8	Optimization of Ultrasound Assisted Extraction (UAE) of Kinsenoside Compound from Anoectochilus roxburghii (Wall.) Lindl by Response Surface Methodology (RSM). Molecules, 2020, 25, 193.	3.8	19
9	Genome-Wide Identification and Expression Analysis of Auxin Response Factor (ARF) Gene Family in Longan (Dimocarpus longan L.). Plants, 2020, 9, 221.	3.5	18
10	Virus-induced gene silencing (VIGS) in Chinese narcissus and its use in functional analysis of NtMYB3. Horticultural Plant Journal, 2021, 7, 565-572.	5.0	16
11	Identification of Genes Involved in Flavonoid Biosynthesis of Chinese Narcissus (Narcissus tazetta L.) Tj ETQq1 1 0.784314 rgBT /Over	1.8	13
12	Discovery of genes involved in anthocyanin biosynthesis from the rind and pith of three sugarcane varieties using integrated metabolic profiling and RNA-seq analysis. BMC Plant Biology, 2021, 21, 214.	3.6	13
13	Genome-Wide Identification and Expression Analysis of Sugar Transporter (ST) Gene Family in Longan (Dimocarpus longan L.). Plants, 2020, 9, 342.	3.5	11
14	NtbHLH1, a JAF13-like bHLH, interacts with NtMYB6 to enhance proanthocyanidin accumulation in Chinese Narcissus. BMC Plant Biology, 2021, 21, 275.	3.6	9
15	Isolation and characterization of GI and FKF1 homologous genes in the subtropical fruit tree Dimocarpus longan. Molecular Breeding, 2017, 37, 1.	2.1	8
16	High efficiency in vitro plant regeneration from epicotyl explants of Ponkan Mandarin (Citrus) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 142	2.1	3
17	Complete chloroplast genome sequence of Syzygium samarangense (Myrtaceae) and phylogenetic analysis. Mitochondrial DNA Part B: Resources, 2022, 7, 977-979.	0.4	3
18	Integrative mRNA and Long Noncoding RNA Analysis Reveals the Regulatory Network of Floral Bud Induction in Longan (Dimocarpus longan Lour.). Frontiers in Plant Science, 0, 13, .	3.6	3