

Genome-wide identification and characterization of<i>Phalaenopsis equestris</i>
development of<i>Phalaenopsis equestris</i>

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Transcriptome-Wide Identification and Expression Analysis of DIVARICATA- and RADIALIS-Like Genes of the Mediterranean Orchid <i>Orchis italica</i> . <i>Genome Biology and Evolution</i> , 2017, 9, 1418-1431.	2.5	22
2	Phylogeny and expression pattern analysis of TCP transcription factors in cassava seedlings exposed to cold and/or drought stress. <i>Scientific Reports</i> , 2017, 7, 10016.	3.3	33
3	Post genomics era for orchid research. , 2017, 58, 61.		29
4	Evolution and Expression Patterns of TCP Genes in Asparagales. <i>Frontiers in Plant Science</i> , 2017, 8, 9.	3.6	31
5	Evolving Tale of TCPs: New Paradigms and Old Lacunae. <i>Frontiers in Plant Science</i> , 2017, 8, 479.	3.6	38
6	Transcriptome “ Scale characterization of salt responsive bean TCP transcription factors. <i>Gene</i> , 2018, 642, 64-73.	2.2	24
7	Genome-Wide Identification of the TCP Transcription Factor Family in Chickpea (<i>Cicer arietinum</i> L.) and Their Transcriptional Responses to Dehydration and Exogenous Absciscic Acid Treatments. <i>Journal of Plant Growth Regulation</i> , 2018, 37, 1286-1299.	5.1	5
8	Genome-Wide Identification of TCP Family Transcription Factors in <i>Medicago truncatula</i> Reveals Significant Roles of miR319-Targeted TCPs in Nodule Development. <i>Frontiers in Plant Science</i> , 2018, 9, 774.	3.6	29
9	Transcriptome-Wide Analysis Reveals the Origin of Peloria in Chinese <i>Cymbidium</i> (<i>Cymbidium sinense</i>). <i>Plant and Cell Physiology</i> , 2018, 59, 2064-2074.	3.1	14
10	Evolutionary and Comparative Expression Analyses of TCP Transcription Factor Gene Family in Land Plants. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3591.	4.1	41
11	Genome-wide identification and transcript analysis of TCP transcription factors in grapevine. <i>BMC Genomics</i> , 2019, 20, 786.	2.8	47
12	Evolution of RADIALIS and DIVARICATA gene lineages in flowering plants with an expanded sampling in non-core eudicots. <i>American Journal of Botany</i> , 2019, 106, 334-351.	1.7	24
13	The crystal structure of the TCP domain of PCF6 in <i>Oryza sativa</i> L. reveals an RHH-like fold. <i>FEBS Letters</i> , 2020, 594, 1296-1306.	2.8	13
14	Genome-Wide Identification and Characterization of the TCP Gene Family in Cucumber (<i>Cucumis</i>) Tj ETQq1 1 0.784314 rgBT/Overlock 2.4 16	2.4	16
15	Expression and Function Studies of CYC/TB1-Like Genes in the Asymmetric Flower <i>Canna</i> (Cannaceae,) Tj ETQq0 0 0 rgBT/Overlock 10 T	3.6	2
16	Evolution and development of three highly specialized floral structures of bee-pollinated <i>Phalaenopsis</i> species. <i>EvoDevo</i> , 2020, 11, 16.	3.2	9
17	Genome-Wide Identification of YABBY Genes in Orchidaceae and Their Expression Patterns in <i>Phalaenopsis</i> Orchid. <i>Genes</i> , 2020, 11, 955.	2.4	20
18	Overexpression of TCP transcription factor OsPCF7 improves agronomic trait in rice. <i>Molecular Breeding</i> , 2020, 40, 1.	2.1	8

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19	Evolution of Class II <i>TCP</i> genes in perianth bearing Piperales and their contribution to the bilateral calyx in <i>Aristolochia</i> . <i>New Phytologist</i> , 2020, 228, 752-769.	7.3	10
20	Orchid Bstsr gene PeMADS28 displays conserved function in ovule integument development. <i>Scientific Reports</i> , 2021, 11, 1205.	3.3	8
21	Phalaenopsis Genome and Transcriptome Exploitation and Its Application for Breeding. <i>Compendium of Plant Genomes</i> , 2021, , 49-65.	0.5	0
22	The ancestral duplicated <i>DL/CRC</i> orthologs, <i>PeDL1</i> and <i>PeDL2</i> , function in orchid reproductive organ innovation. <i>Journal of Experimental Botany</i> , 2021, 72, 5442-5461.	4.8	18
23	Extending the Toolkit for Beauty: Differential Co-Expression of DROOPING LEAF-Like and Class B MADS-Box Genes during Phalaenopsis Flower Development. <i>International Journal of Molecular Sciences</i> , 2021, 22, 7025.	4.1	9
24	OrchidBase 4.0: a database for orchid genomics and molecular biology. <i>BMC Plant Biology</i> , 2021, 21, 371.	3.6	10
25	Proteomic profiling uncovered the cytosolic superoxide dismutase BsSOD1 associated with plant defence in the herbal orchid <i>Bletilla striata</i> . <i>Functional Plant Biology</i> , 2020, 47, 937.	2.1	5
28	Comprehensive In Silico Characterization and Expression Profiling of TCP Gene Family in Rapeseed. <i>Frontiers in Genetics</i> , 2021, 12, 794297.	2.3	13
29	Organ-Specific Gene Expression Reveals the Role of the Cymbidium ensifolium-miR396/Growth-Regulating Factors Module in Flower Development of the Orchid Plant Cymbidium ensifolium. <i>Frontiers in Plant Science</i> , 2021, 12, 799778.	3.6	9
30	Molecular genetic insights into orchid reproductive development. <i>Journal of Experimental Botany</i> , 2022, 73, 1841-1852.	4.8	10
31	Transcriptome atlas of <i>Phalaenopsis equestris</i> . <i>PeerJ</i> , 2021, 9, e12600.	2.0	1
32	Genetic insights into the regulatory pathways for continuous flowering in a unique orchid <i>Arundina graminifolia</i> . <i>BMC Plant Biology</i> , 2021, 21, 587.	3.6	11
38	Genome-Wide Identification and Expression Pattern Analysis of the TCP Gene Family in Radish (<i>Raphanus sativus</i> L.). <i>Horticulturae</i> , 2022, 8, 656.	2.8	1
39	Transcriptome mining of hormonal and floral integrators in the leafless flowers of three cymbidium orchids. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	1
40	Evolutionary analyses and expression patterns of TCP genes in Ranunculales. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	3
41	Genome-wide analysis of the TCP gene family and their expression pattern in <i>Cymbidium goeringii</i> . <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	6
42	Genome-Wide Identification and Characterization of TCP Gene Family Members in <i>Melastoma candidum</i> . <i>Molecules</i> , 2022, 27, 9036.	3.8	8
43	PeGRF6-PeGIF1 complex regulates cell proliferation in the leaf of <i>Phalaenopsis equestris</i> . <i>Plant Physiology and Biochemistry</i> , 2023, 196, 683-694.	5.8	2

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44	Advances in Research on the Regulation of Floral Development by CYC-like Genes. Current Issues in Molecular Biology, 2023, 45, 2035-2059.	2.4	3
45	The function of BoTCP25 in the regulation of leaf development of Chinese kale. Frontiers in Plant Science, 0, 14, .	3.6	0
46	A novel MORN-motif type gene GmMRF2 controls flowering time and plant height of soybean. International Journal of Biological Macromolecules, 2023, 245, 125464.	7.5	0
47	Genome-Wide Identification of TCP Gene Family in Dendrobium and Their Expression Patterns in Dendrobium chrysotoxum. International Journal of Molecular Sciences, 2023, 24, 14320.	4.1	1
48	Genome-wide analysis reveals the TCP-miR159-miR319 module is crucial for rice (Oryza sativa L.) growth and response to drought and salinity.. Plant Stress, 2023, 10, 100215.	5.5	0
49	Evolution and development of fruits of Erycina pusilla and other orchid species. PLoS ONE, 2023, 18, e0286846.	2.5	1
50	PeCIN8 expression correlates with flower size and resistance to yellow leaf disease in Phalaenopsis orchids. BMC Plant Biology, 2023, 23, .	3.6	0
53	Sweet cherry TCP gene family analysis reveals potential functions of PavTCP1, PavTCP2 and PavTCP3 in fruit light responses. BMC Genomics, 2024, 25, .	2.8	0
54	Genome-wide identification and characterization of TCP gene family in Dendrobium nobile and their role in perianth development. Frontiers in Plant Science, 0, 15, .	3.6	0
55	Genomic survey, bioinformatics analysis, and expression profiles of TCP genes in Liriodendron chinense and functional characterization of LcTCP4. Trees - Structure and Function, 2024, 38, 287-302.	1.9	0