

Genome-wide identification and characterization of<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.	1.1	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.	1.6	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.	1.7	31
5	Evolving Tale of TCPs: New Paradigms and Old Lacunae. <i>Frontiers in Plant Science</i> , 2017, 8, 479.	1.7	38
6	Transcriptome “ Scale characterization of salt responsive bean TCP transcription factors. <i>Gene</i> , 2018, 642, 64-73.	1.0	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 Abscisic Acid Treatments. <i>Journal of Plant Growth Regulation</i> , 2018, 37, 1286-1299.	2.8	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.	1.7	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.	1.5	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.	1.8	41
11	Genome-wide identification and transcript analysis of TCP transcription factors in grapevine. <i>BMC Genomics</i> , 2019, 20, 786.	1.2	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.	0.8	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.	1.3	13
14	Genome-Wide Identification and Characterization of the TCP Gene Family in Cucumber ( <i>Cucumis</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 16	1.0	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	1.7	2
16	Evolution and development of three highly specialized floral structures of bee-pollinated <i>Phalaenopsis</i> species. <i>EvoDevo</i> , 2020, 11, 16.	1.3	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.	1.0	20
18	Overexpression of TCP transcription factor OsPCF7 improves agronomic trait in rice. <i>Molecular Breeding</i> , 2020, 40, 1.	1.0	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.	3.5	10
20	Orchid Bsister gene PeMADS28 displays conserved function in ovule integument development. <i>Scientific Reports</i> , 2021, 11, 1205.	1.6	8
21	Phalaenopsis Genome and Transcriptome Exploitation and Its Application for Breeding. <i>Compendium of Plant Genomes</i> , 2021, , 49-65.	0.3	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.	2.4	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.	1.8	9
24	OrchidBase 4.0: a database for orchid genomics and molecular biology. <i>BMC Plant Biology</i> , 2021, 21, 371.	1.6	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.	1.1	5
28	Comprehensive In Silico Characterization and Expression Profiling of TCP Gene Family in Rapeseed. <i>Frontiers in Genetics</i> , 2021, 12, 794297.	1.1	13
29	Organ-Specific Gene Expression Reveals the Role of the Cymbidium <i>ensifolium</i> -miR396/Growth-Regulating Factors Module in Flower Development of the Orchid Plant <i>Cymbidium ensifolium</i> . <i>Frontiers in Plant Science</i> , 2021, 12, 799778.	1.7	9
30	Molecular genetic insights into orchid reproductive development. <i>Journal of Experimental Botany</i> , 2022, 73, 1841-1852.	2.4	10
31	Transcriptome atlas of <i>Phalaenopsis equestris</i> . <i>PeerJ</i> , 2021, 9, e12600.	0.9	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.	1.6	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.	1.2	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, .	1.7	1
40	Evolutionary analyses and expression patterns of TCP genes in Ranunculales. <i>Frontiers in Plant Science</i> , 0, 13, .	1.7	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, .	1.7	6
42	Genome-Wide Identification and Characterization of TCP Gene Family Members in <i>Melastoma candidum</i> . <i>Molecules</i> , 2022, 27, 9036.	1.7	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.	2.8	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.	1.0	3
45	The function of BoTCP25 in the regulation of leaf development of Chinese kale. Frontiers in Plant Science, 0, 14, .	1.7	0