

Table S6 Summary of TCP proteins in ornamental plants

Family	Species	Number of TCP genes	References
Ginkgoaceae	<i>Ginkgo biloba</i>	13	Yu et al., 2022
Hypoxidaceae	<i>Hypoxis decumbens</i>	15	Madrigal et al., 2017
Orchidaceae	<i>Cattleya trianae</i>	18	Madrigal et al., 2017
	<i>Dendrobium chrysotoxum</i>	14	Li et al., 2023; Huang et al., 2023
	<i>Dendrobium officinale</i>	24	Li et al., 2023
	<i>Dendrobium nobile</i>	23	Li et al., 2023; Wei et al., 2024
	<i>Erycina pusilla</i>	15	Tang et al., 2024
	<i>Phalaenopsis equestris</i>	23	Lin et al., 2016
	<i>Dactylis glomerata</i>	22	Wang et al., 2023
Poaceae	<i>Panicum virgatum</i>	41	Zheng et al., 2019
	<i>Phyllostachys edulis</i>	16	Liu et al., 2018
	<i>Prunus mume</i>	19	Zhou et al., 2016
Rosaceae	<i>Prunus mira</i>	19	Zhang et al., 2024
	<i>Prunus amygdalus</i>	19	Zhang et al., 2024
	<i>Prunus fruticosa</i>	19	Zhang et al., 2024
	<i>Rosa chinensis</i>	16;18	Hou et al., 2022; Cheng et al., 2023
	<i>Melastoma candidum</i>	35	Li et al., 2022
Amaranthaceae	<i>Amaranthus hypochondriacus</i>	16	Wei et al., 2023
Convolvulaceae	<i>Ipomoea nil</i>	28	Wei et al., 2023
Solanaceae	<i>Petunia axillaris</i>	66;28	Zhang et al., 2020; Wei et al., 2023
Apocynaceae	<i>Catharanthus roseus</i>	15	Hao et al., 2023
Lentibulariaceae	<i>Utricularia gibba</i>	44	Wei et al., 2023
Oleaceae	<i>Osmanthus fragrans</i>	39	Zheng et al., 2023
Asteraceae	<i>Chrysanthemum lavandulifolium</i>	39	Wu et al., 2023a
	<i>Helianthus annuus</i>	34	Wu et al., 2023b
	<i>Taraxacum officinale</i>	33	Wei et al., 2023
	<i>Cynara cardunculus</i>	33	Wei et al., 2023
	<i>Carthamus tinctorius</i>	33	Wei et al., 2023
Phrymaceae	<i>Mimulus guttatus</i>	26	Wei et al., 2023
Caprifoliaceae	<i>Lonicera japonica</i>	20	Wei et al., 2023

References

- Wei X, Judith R, Lidija B, Tao Z, Henri van de G, Carla O, Marco B, Julie F de C, Ingrid M. van der M, Koen J. F. V, et al. 2023. Phylogenomic analysis provides insights into MADS-box and TCP gene diversification and floral development of the Asteraceae, supported by de novo genome and transcriptome sequences from dandelion (*Taraxacum officinale*). *Front. Plant Sci* **14**:1198909.
- Li Y, Li L, Yang J, Niu Z, Liu W, Lin Y, Xue Q, Ding X. 2023. Genome-Wide identification and analysis of TCP gene family among three *Dendrobium* species. *Plants* **12**:3201.
- Wang C, Feng G, Xu X, Huang L, Nie G, Li D, Zhang X. 2023. Genome-Wide identification, characterization, and expression of TCP genes family in Orchardgrass. *Genes* **14**:925.
- Huang Y, Zhao XW, Zheng QY, He X, Zhang M.-M., Ke SJ, Li YY, Zhang CL, Ahmad S, Lan S, et al. 2023. Genome-Wide Identification of TCP Gene Family in *Dendrobium* and Their Expression

Patterns in *Dendrobium chrysotoxum*. *Int. J. Mol. Sci.* **24**:14320.

Yu L, Chen QW, Zheng JR, Xu F, Ye JB, Zhang WW, Liao YL, Yang XY. 2022. Genome-wide identification and expression pattern analysis of the TCP transcription factor family in *Ginkgo biloba*. *PLANT SIGNALING & BEHAVIOR* **17**:e1994248.

Zheng AQ, Sun FL, Cheng TT, Wang YF, Xie KL, Zhang C, Xi YJ. 2019. Genome-wide identification of members of the TCP gene family in switchgrass (*Panicum virgatum* L.) and analysis of their expression. *Gene* **702**:89-98.

Zhang Q, Qian C, Li LL, Li W, Li YH, Zhao H. 2024. Genome-Wide Identification and Characterization of TCP Genes in Eight *Prunus* Species and Their Expression Patterns Under Cold Stress in *P. tenella* var. *tenella*. *Genes* **15**:1443.

Liu HL, Wu M, Li F, Gao YM, Chen F, Xiang Y. 2018. TCP Transcription Factors in Moso Bamboo (*Phyllostachys edulis*): Genome-Wide Identification and Expression Analysis. *Frontiers in plant science* **9**:1263.