Flowering retardation by high temperature in chrysanthemums: involvement of *FLOWERING LOCUS T-like 3* gene repression. *Yoshihiro Nakano, Yohei Higuchi, Katsuhiko Sumitomo, Tamotsu Hisamatsu* 

## SUPPLEMENTARY DATA

**Figure S1.** Representative *C. seticuspe* exposed to 30 °C for 1 week at different growth stages. Plants were grown at 20 °C with a 10-h photoperiod and transferred to 30 °C at the first, second, third, fourth or fifth week. The plants were photographed after 50 SDs.





**Fig. S2.** Expression analysis of *CmFTL3* in the leaves of *C. morifolium* 'Mona Lisa' (A) and 'Kurarisu' (B) by using qRT-PCR in grafting experiment. Grafted plants were grown at 20 °C or 27 °C with a 10-h photoperiod. The leaves were harvested from the stock after 3 weeks. Open and closed horizontal bars represent light and dark periods, respectively. *CmACT* was used as an internal standard. Values are means  $\pm$  SE (n = 3).



**Fig. S3.** Expression analysis of *CsTFL1* in the shoot tip of *C. seticuspe* by using qRT-PCR. Shoot tips (3 mm) were harvested every 1 week from plants grown at 20 °C or 30 °C with a 10-h photoperiod. *CsEF1a* was used as an internal standard. Values are means  $\pm$  SE (n = 3).



**Fig. S4.** Expression analysis of *CsFTL1* in the leaves of *C. seticuspe* by using qRT-PCR. Plants were grown at 20 °C or 30 °C with a 10 h-photoperiod. The leaves were harvested after 2 weeks. (A) Comparison with NB condition at light on. (B) Diurnal expression changes at 20 °C and 30 °C. Open and closed horizontal bars represent light and dark periods, respectively. *CsACT* was used as an internal standard. Values are means  $\pm$  SE (n = 3).

Target	Accession no.	Forward primer	Reverse primer
CsFTL3 CmFTL3	AB679272 AB545937	GAAGGCGTAACAAATAAAAATAGC	CAAGTCTTACAATTTGGTACTGTCG
CsACT CmACT	xxxxxxxx xxxxxxxx	CAGGATGAGCAAGGAAATCACC	AGGTGCTGAGTGATGCAAGGAT
CsM8	XXXXXXXX	TATCATGCCCCCATGGATGG	CCATTTCACATCCATGGTCG
CsM19	XXXXXXXX	CATGCAATTGCTACTGGAGC	ATGAACAAGGATAACGAGCC
CsM37	XXXXXXXX	GGTCGAAACTATCTTCAGTCG	CCAAGAAACCTTAAGTCCTTC
CsM41	XXXXXXXX	CTCGAGCTCATCATGTCTCC	GGTCTTACGTGTATATGATGC
CsM44	XXXXXXXX	CCATCGGATTCATTAACGGC	TTACTGCAACTTGGCCCTGC
CsM86	XXXXXXXX	GATTACCAAGCTCATGAACC	ATTACAAACCTTGGGCAACG
CsTFL1	XXXXXXXX	GCAGCCAGAAGACGTTAACC	GAAGATCCTCACAATTAGGCC
CsFTL1	AB679270	CATATTGAAAGCACGCACACAT	ACACGGGTTGGCTGTAGCATT

 Table S1.
 Primer sequences used in this study