

Fig. S1 Quantitative RT-PCR analysis of *PbELF4a* and *PbELF4b* expression in 10-day-old seedlings from control plants and transgenic lines. *AtACT* served as the internal control for gene expression analysis. Data are presented as means  $\pm$  SD (n = 3). The sampling times were ZT1, ZT8, and ZT15.

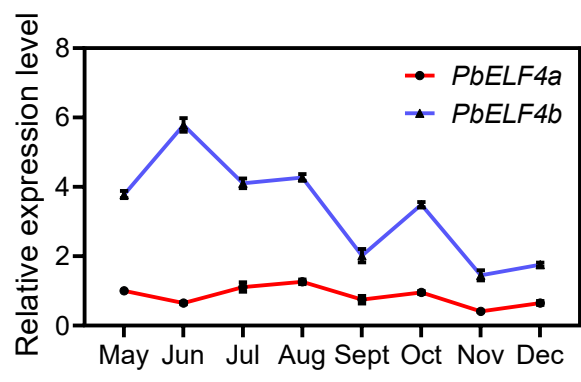


Fig. S2 Dynamic expression patterns of *PbELF4a* and *PbELF4b* during pear leaves development. The expression of *PbELF4a* in sample May was set to "1." *PbUBQ* was used as an internal control. Data are presented as means  $\pm$  SD (n = 3).