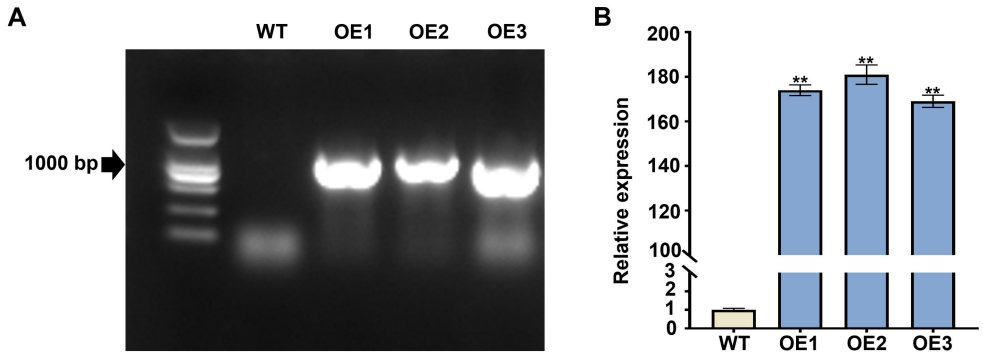


Supplementary Figure S1. Identification of *MuMYB116*-overexpressing (OE) transgenic *Arabidopsis* lines. (A) Genomic PCR validation of the MuMYB116 transgene integration in stable overexpressing *Arabidopsis* lines. (B) Relative transcript levels of *MuMYB116* in wild-type (WT) and transgenic lines (OE1, OE2, and OE3). Data are presented as mean ± SD (n = 3 biological replicates). ** $p < 0.01$, (Student's *t*-test).