



Figure S5 Transcriptome sequencing analysis of *PagbZIP60*. (A) Differentially expressed genes (DEGs) with an adjusted $|\log_2(\text{FoldChange})| \geq 1$ and a $\text{padj} < 0.05$ were identified between OE and WT. (B-C) KEGG and GO enrichment analysis of DEGs. OE: *PagbZIP60* overexpression lines, WT: non-transgenic poplar.