

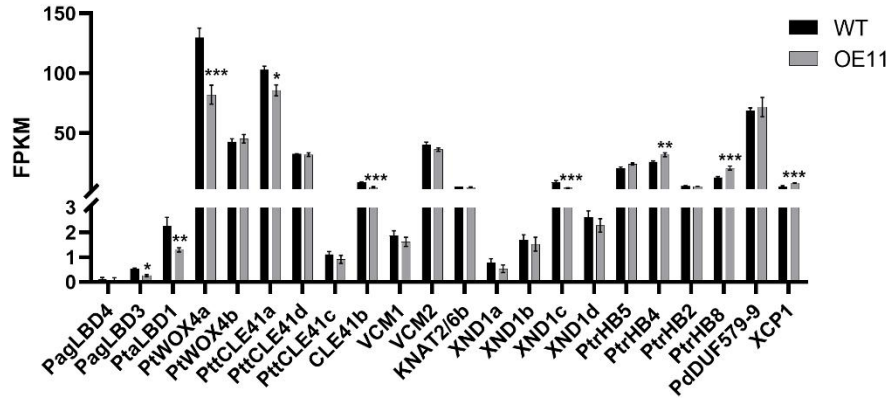


Fig. S5 Gene expression levels associated with cambium in WT and *PagAS1a* overexpression transgenic line (OE11) from RNA-seq. The average fragments per kilobase of exon per million fragments mapped (FPKM) for each gene from RNA-seq experiments are shown. Asterisks indicate significant differences between WT and *PagAS1a*-OE plants (Student's t-test): * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.