





Additional file 15. Validation of small RNAs and microarray data by qRT-PCR. All the values are calculated by fold change (*pap2*/WT, OE7/WT or OE21/WT). Columns in white indicate small RNAs reads or microarray data, black columns indicate real-time RT-PCR data. Asterisks indicate significant difference compared with WT. * P < 0.05, ** P < 0.01.