

Supplemental Figure 5

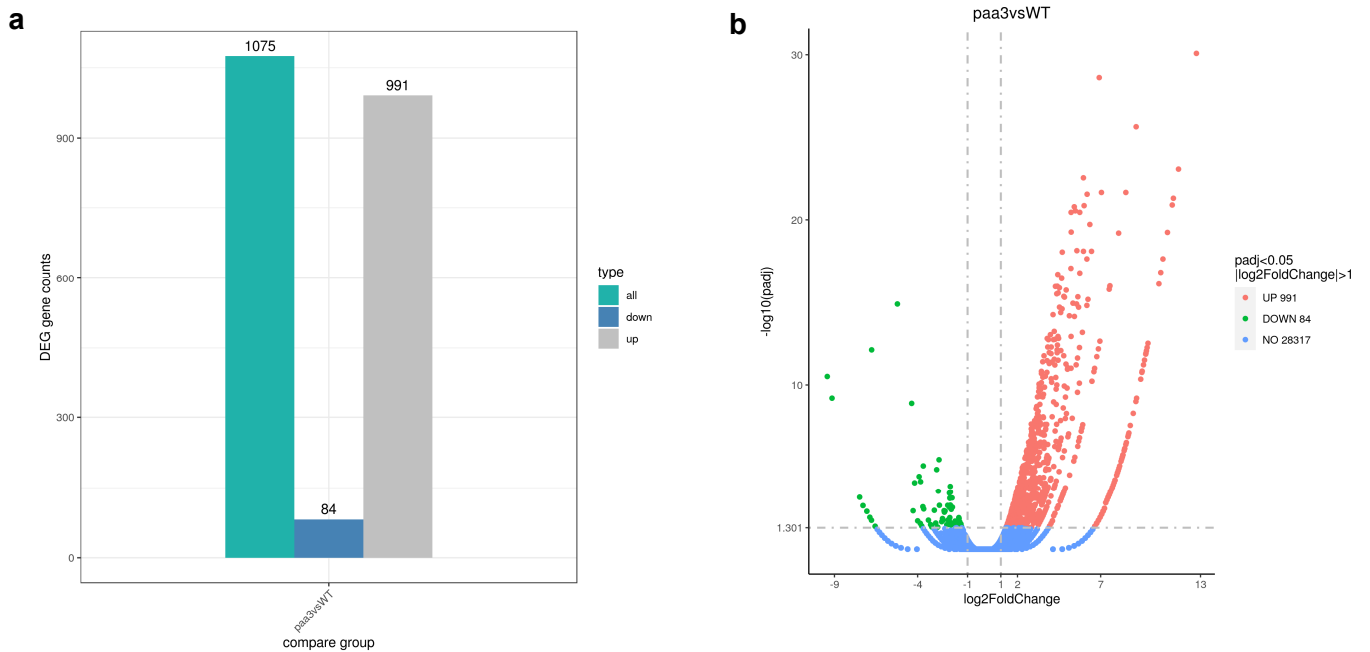


Fig. S5, transcriptome analysis of *paa3*. **a**, Showed that there were 1075 differential genes in WT and *paa3* samples, among which 991 genes were up-regulated and 84 genes were down-regulated. **b** The volcano figure showed an overall overview, including 28,317 genes with no change in expression, 991 up-regulated genes and 84 down-regulated genes.