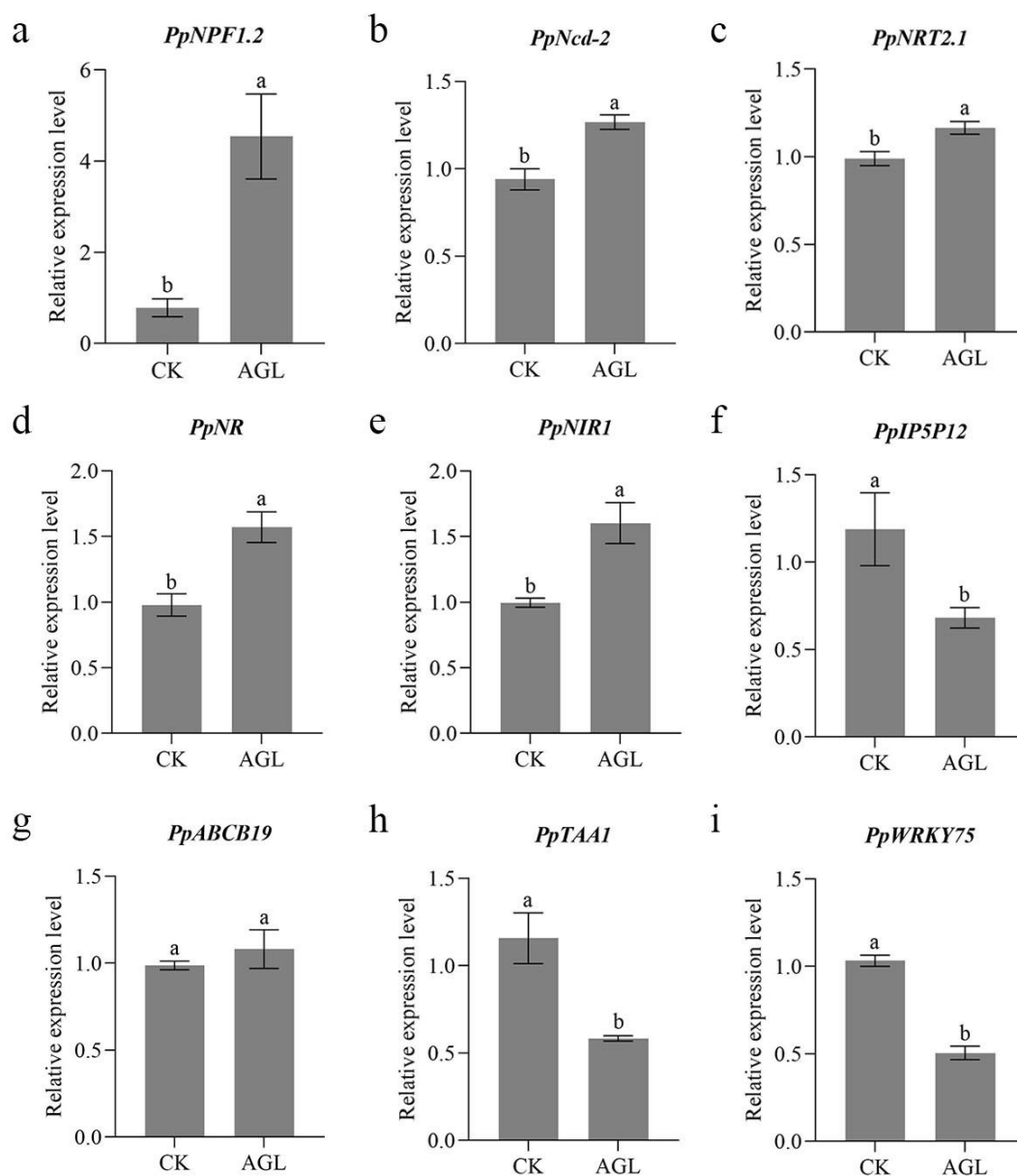




Supplementary figure.S3 RT-qPCR validation of relative expression levels of related genes within peach roots after homologous overexpression of *PpAGL24*. (a-i) *PpNPF1.2*, *PpNcd-2*, *PpNRT2.1*, *PpNR*, *PpNIR1*, *PpIP5P12*, *PpABCB19*, *PpTAA1* 和 *PpWRKY75*.