



Supplementary Figure S4: qRT-PCR verification of three selected DEGs and Pearson relationship with FPKM and qRT-PCR. Comparison of RNA-seq data (green line) with qRT-PCR data (black bar). The normalized expression level (FPKM) of RNA-seq is indicated on the right y-axis. The relative qRT-PCR expression level is shown on the left y-axis. eEF-1a gene was used as the internal control.