

Figure S1 The 18 most significantly enriched KEGG pathways based on the coregulated DEGs in pericarp and seeds. The dot color and size indicate the P-value and gene number, respectively, as shown on the right.

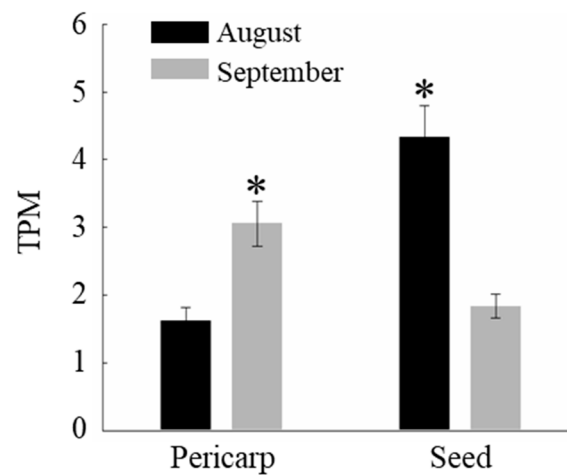


Figure S2 Transcriptional level of gene *TEA024158*.