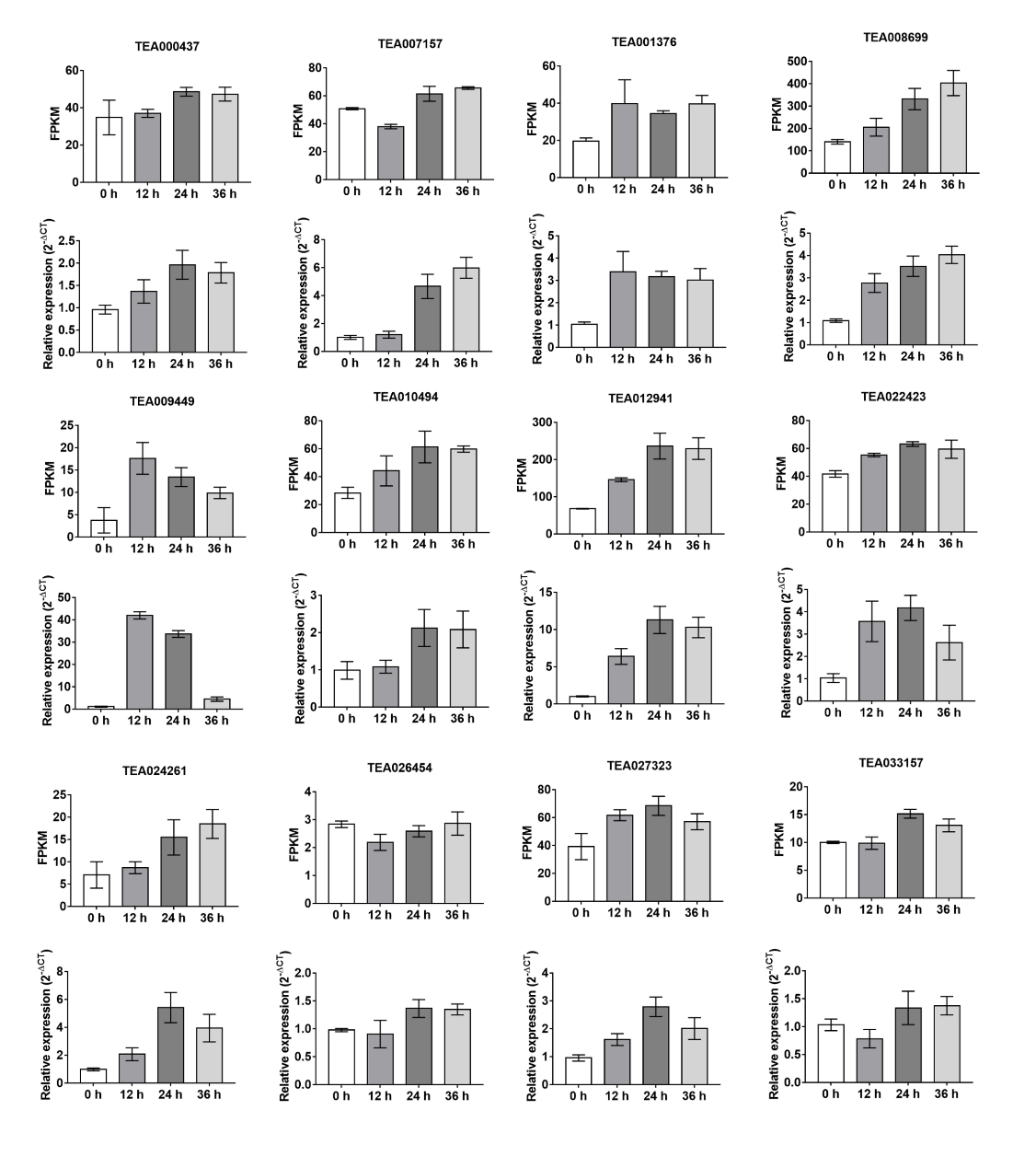
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**Fig. S4.** The expression patterns of 12 genes in α-linolenic acid metabolism and terpenoid biosynthesis pathways based on RNA-seq and RT-PCR analysis. The data represent the mean + SD (n = 3).