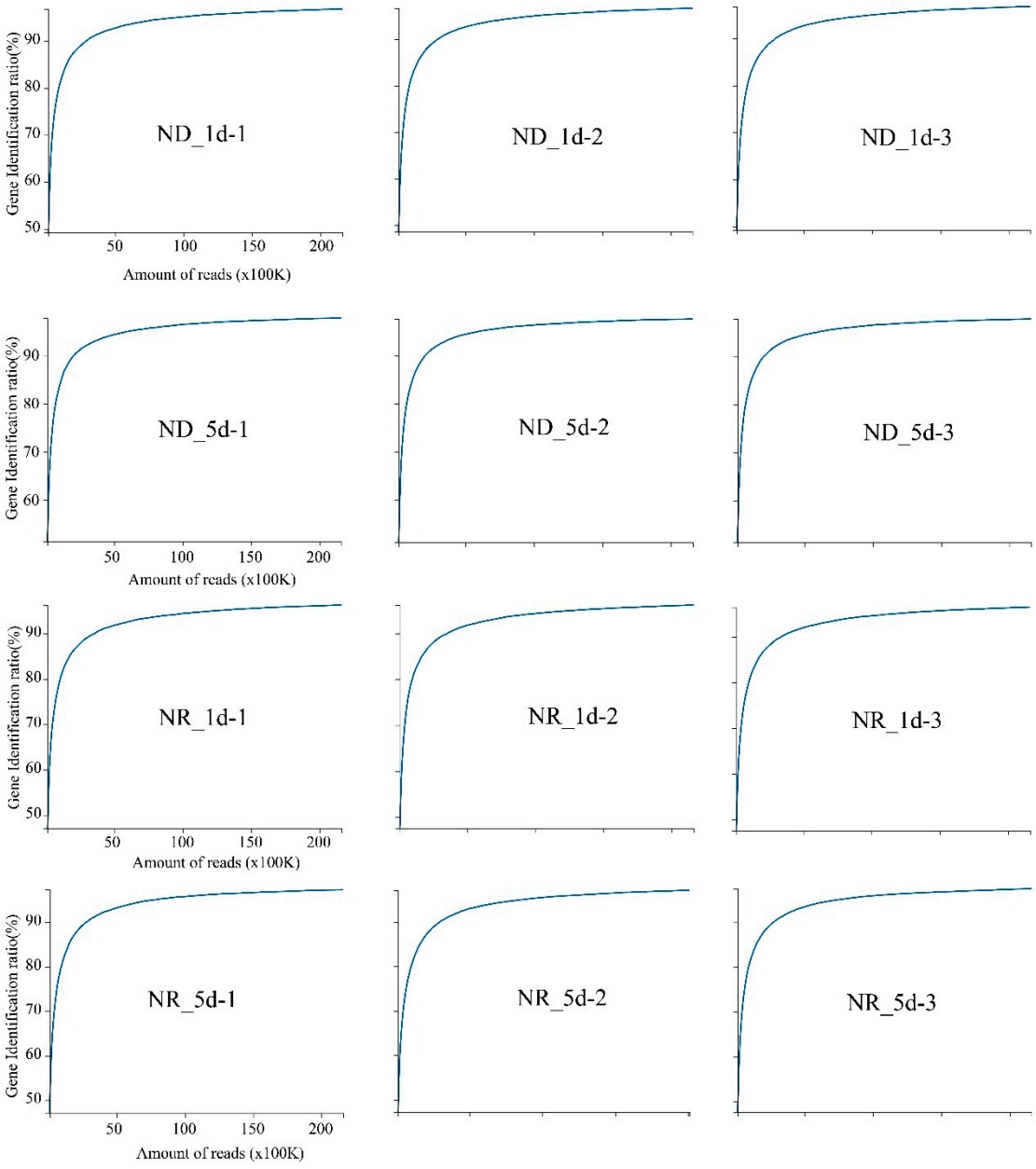


Figure S1. Sequencing saturation curves of different samples.

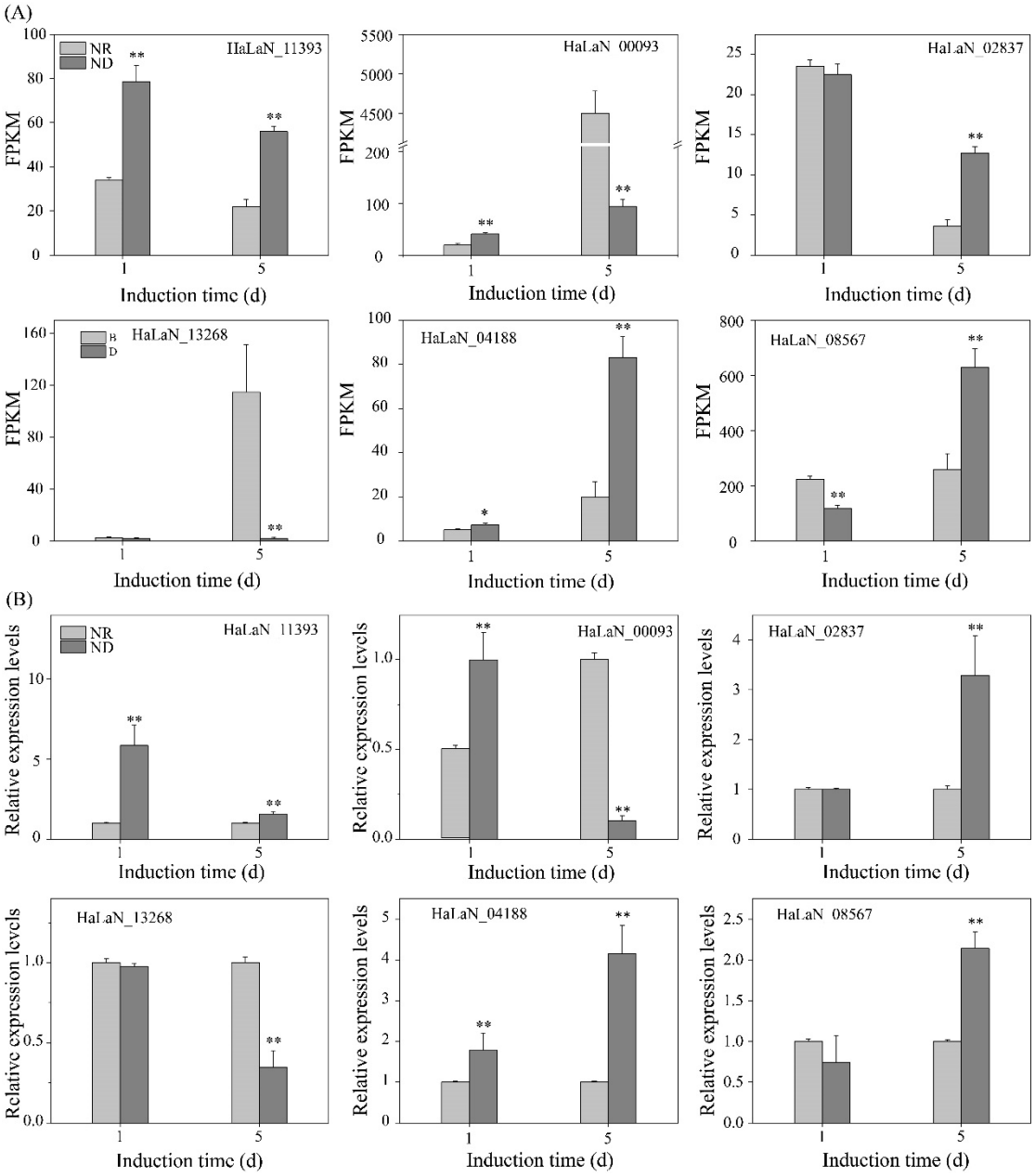


Figure S2. The gene expression of six genes with randomly selected from different metabolic pathways by RNA-seq (A) and qRT-PCR (B).