



Figure S2. Statistics of Gene Ontology (GO) enrichment (left part) and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway enrichment (right part) of differentially expressed genes in trans12 vs. trans0 group (upper pictures) and trans48 vs. trans0 group (bottom pictures), respectively. p -value ≤ 0.05 and the absolute value of Log2 FC (fold-change) ≥ 1.0 are used as the threshold for significant difference expression.