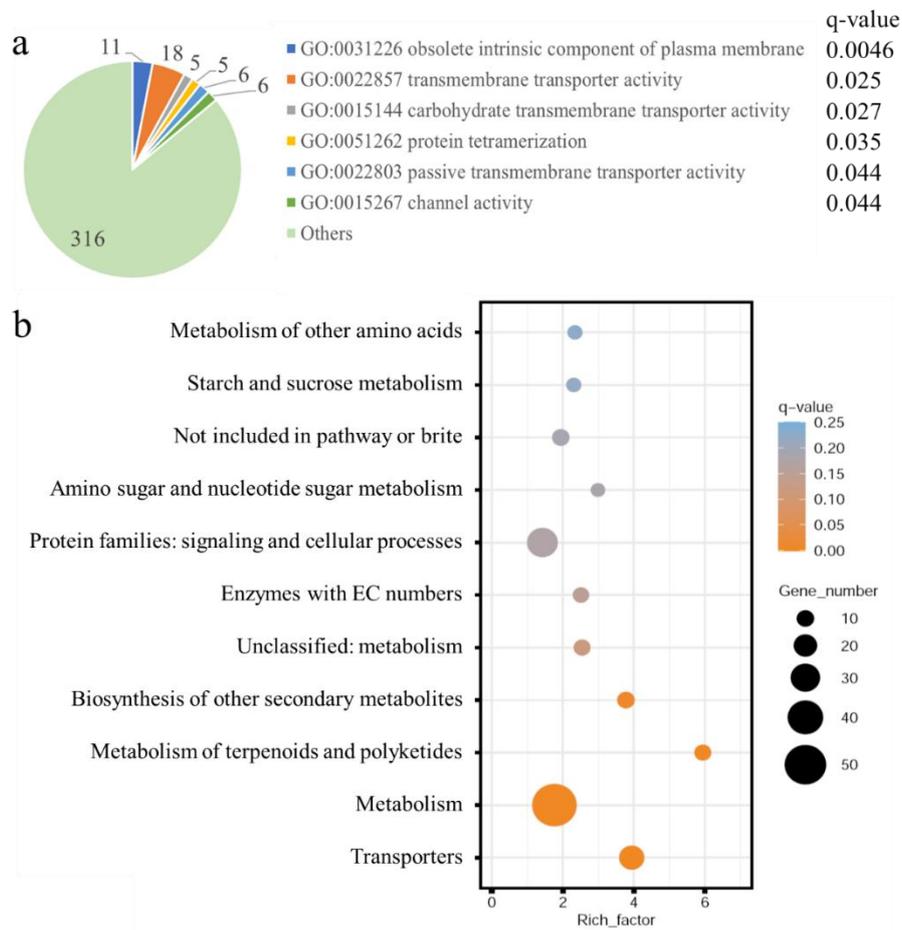


The size of each bubble indicates the number of genes associated with the pathway, while the color gradient represents the significance level (q value).



Supplementary Fig. S4 Functional enrichment analysis of hybrid-specific differentially expressed genes (DEGs).

(a) The most significantly enriched Gene Ontology (GO) terms for hybrid-specific DEGs. For each GO term, the numbers of associated genes is displayed within the pie chart, and the corresponding q-value is listed to the right of the term. (b) KEGG pathway enrichment of the hybrid-specific DEGs. **The bubble size and color indicates the gene count and significance level, respectively.**