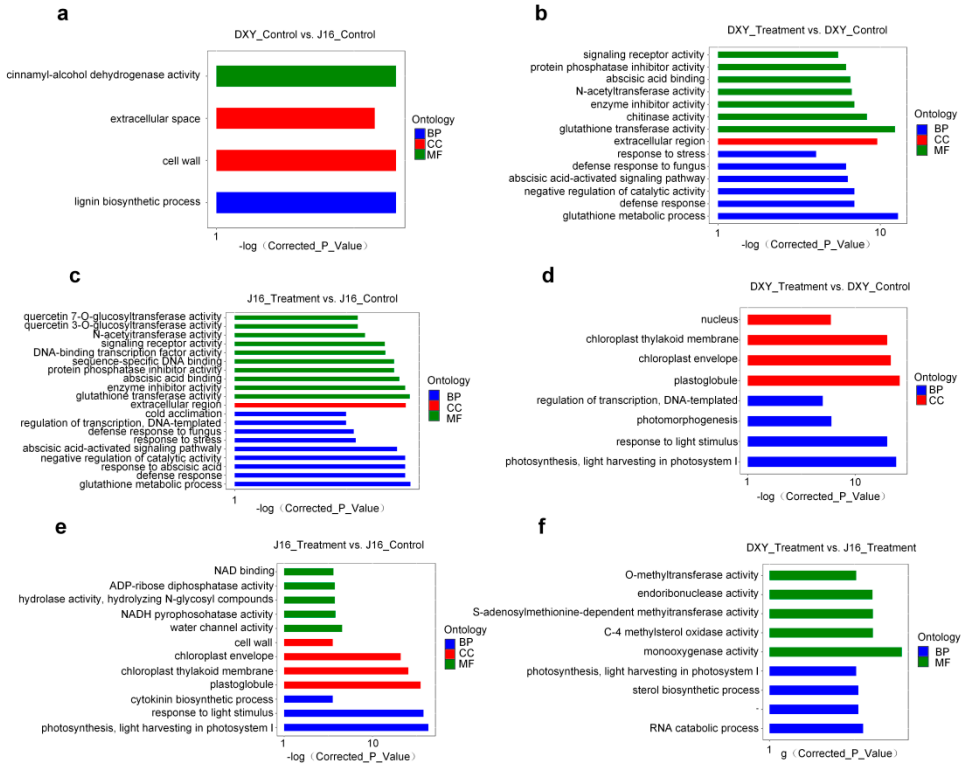


Figure S2. Gene Ontology (GO) enrichment analysis of differentially expressed genes (DEGs) across different comparison groups

(a-c) Enriched GO terms for upregulated DEGs in the following comparison groups, respectively: DXY_C vs J16_C (a), DXY_T vs DXY_C (b), J16_T vs J16_C (c). (d-f) Enriched GO terms for downregulated DEGs in the following comparison groups, respectively: DXY_T vs DXY_C (d), J16_T vs J16_C (e), DXY_T vs J16_T (f).