



Supplementary Fig. 4. Gene Ontology (GO) enrichment analysis of downregulated differentially expressed genes (DEGs) in So-cheong2 (SC2) at each time point. The 30 most significantly overrepresented GO terms of each DEG set are presented. The color and size of each dot indicate the $-\log_{10} P\text{-value}$ and enrichment score, respectively. DP, Daepung.