



Supplementary Fig. 3. Gene Ontology (GO) enrichment analysis of differentially expressed genes (DEGs) downregulated at 6 and 12 h after inoculation (HAI) compared with 0 HAI in Socheong2 (SC2) and Daepung (DP) cultivars. The 30 most significantly over-represented GO terms from each DEG set are shown. The color and size of each dot represent the $-\log_{10} P$ -value and enrichment score, respectively.