S3 Fig. Enrichment analysis of day 6 and 12 compared to NP *DN*, *UP* and *OTHER* subsets. GSEA analysis provided statistical analysis of gene expression biases in the *DN* (**A, D**), *UP* (**B, E**) and *OTHER* (**C, F**) subsets constructed as described in **S2 Fig.** and Material and Methods section. A, E: *DN* subset shows significant enrichment for down-regulated genes (q-value<10⁻⁷). B, E: *UP* subset shows significant enrichment for up-regulated genes (q-value<10⁻⁷). C, F: *OTHER* subset shows no enrichment (q-value>0.05).