



**Figure S4.** Significantly enriched GO terms for the target genes of peach miRNAs in DPA50 vs. DPA20 (A) and DPA75 vs. DPA50 (B) libraries; enriched factors were calculated according to the numbers of differentially expressed genes which were mapped to GO term divided by the numbers of all genes in that GO term; the top 20 enriched GO terms are listed.

DPA, days post anthesis