



**Figure S2.** Pathway analysis of the target genes of differentially expressed miRNAs of *X. sorbifolium*. The abscissa is the enrichment factor (Number of target genes of differential miRNAs enriched in this pathway/ Number of background genes in the pathway), the ordinate is the corresponding path description, the bubble size is the number of differential genes, and the bubble color is the enrichment significant P value. (**A**): Down-regulation in Fa vs. Fb; (**B**): Up-regulation in Fa vs. Fb; (**C**): Down-regulation in Ma vs. Mb; (**D**): Up-regulation in Ma vs. Mb; (**E**): Up-regulation in Ma vs. Fa; (**F**): Down-regulation in Mb vs. Fb; (**G**): Up-regulation in Mb vs. Fb.