



**Supplementary Fig. 4.** Gene ontology of the biological process of JM1 cells knocked down (A) antimir-1307-5p and (B) antimir-193-5p. Gene ontology of top 20 biological process in miRNAs knocked down. FDR was computed by Benjamini-Hochberg method. Fold enrichment is the measure of enrichment; higher values indicate higher enrichment.