Figure S1. Venn diagram of target genes predicted in TargetScan and miRDB. Among these genes, 943 were predicted for miR-106a-5p, 438 for miR-103a-3p and 771 for miR-29a-3p in both databases.
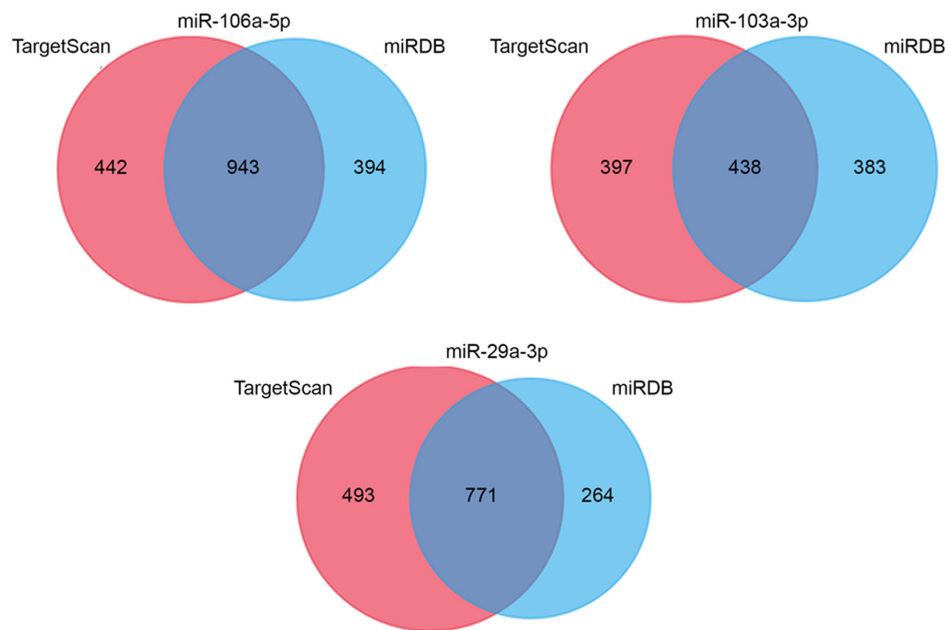
Table SI. Sequence of the forward primers for reverse transcription-quantitative PCR. The kit provides universal reverse primers.

Primer name	Sequence 5'→3'
cel-miR-238-3p	TTTGTACTCCGATGCCATTCAGA
hsa-miR-106a-5p	AAAAGTGCTTACAGTGCAGGTAG
hsa-miR-103a-3p	CAGCATTGTACAGGGCTATGA
hsa-miR-29a-3p	GTAGCACCATCTGAAATCGGTTA

miR, microRNA.