

## SUPPLEMENTARY FIGURE

	Predicted consequential pairing of target region (top) and miRNA (bottom)	Site type	Context++ score	Context++ score percentile	Weighted context++ score	Conserved branch length	PCT	Predicted relative $K_D$
Position 285-291 of ANGPTL4 3' UTR	5' ... AGUUGGGGACUCAGAGGGACCAC ...	7mer-m8	-0.39	95	-0.39	4.416	0.85	-4.676
hsa-miR-133a-3p.1	3' GUCGACCAACUUCUCCUUGUUU							

Supplementary Figure 1. Predicted consequential pairing of target region of ANGPTL4 and miR-133a-3p.