Table S2, ABCC5 3’UTR mRNA presents a perfect match with the seed sequence of mir-128 by using TargetScan

	predicted consequential pairing of target region (top) and miRNA (bottom)	seed match	site-type contri-bution	3' pairing contri-bution	local AU contri-bution	position contri-bution	context score	context score percentile
Position 373-379 of ABCC5 3' UTR hsa-miR-128	5' ...UUAUAUUAAAAUAAGCACUGUGC... 3' UUUCUCUGGCCAAGUGACACU	7mer-m8	-0.161	0.067	-0.082	-0.003	-0.17	70