

## SUPPLEMENTARY TABLE

**Supplementary Table 1. Primers for qRT-PCR analysis.**

Target gene	Forward primer	Reverse primer	RT primer
XAGE5	CTGGTTGGGCCTA TGCTTGAG	TCCGCATTCATCCCA GTCT	
COX7B2	CCAGAAATGCACT AAGCAGTCT	ACCCATGTAGCAACA CAGAAAG	
MAGEB2	GCTGCGGGTGTTT CATCCA	TGGTTAGAGGATCTT CGCTTG	
BHMT	TGCTGGAGAGATT GTGATTGGA	CTTGCTTCACTCGCA TAGAAGG	
PROM1	AGTCGGAAACTG GCAGATAGC	GGTAGTGTGTACTG GGCCAAT	
SLCO1B3	TGGAGCAACAGTA CGGTCAG	TGCTTTCGCAGATTA GAGGGAA	
RP11- 332K15.1	ACTCTCGGGATCC TGGACCT	AGTCCTCAGTGCTTG CGACA	
XX-CR54.1	TGTACTCATCAGC ACGGCATAAAGG	ATCAGAAGGCAACAC TCAGGACTG	
BC038578	AGCCAGCGAGAC CACGAACC	TAAGGTGGCGTGTCT GGAGTCTG	
LOC389023	CGTGTCTGAGATT GTGGCCTGAG	TAGGAGTTCACCGA CGTGACC	
SSTR5-AS1	GTGTTCTCCATG AAGAGCAGAGC	GCACGCAGCACACTC CTTCC	
FAM99A	GGCCTGGCTCACT CCGTGTC	TCCTCTGGCTCTCCTT CATTCTGC	
hsa_circRNA_ 103809	TCGAGACCTCTGT CAGCGAG	GCCAATACTCCCACT CGCAA	
hsa_circRNA_ 101368	AAGACTTGAGGC GAATGG	GCACAGGTGAATAGA CTTCT	
hsa_circRNA_ 102399	GTGCAGGGTCCG AGGTATT	GCCGTCGTAATACTG CCTGGT	
hsa_circRNA_ 000639	ATGCCACAGCTT TCCAA	TCTCCTTCTCCCTCT ACATT	
hsa_circRNA_ 001547	ACATACCCGTTGG CTCTC	CAGGGTCATCCACAA TCAG	
hsa_circRNA_ 400010	GTATAGGATGACT CACTGACA	GCACTCAACAATCGT TAGC	
miR-532	GCCTCCCACACCC AAGG	AGTGCAGGGTCCGAG GTATT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGC ACTGGATACGACTGCAAG
miR-130	CGCGACTCTTTCC CTGTTG	AGTGCAGGGTCCGAG GTATT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGC ACTGGATACGACGTAGTG
miR-642	GCGGTCCCTCTCC AAATGT	AGTGCAGGGTCCGAG GTATT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGC ACTGGATACGACCAAGAC
miR-329	GCGGAGGTTTTCT GGGTTTC	AGTGCAGGGTCCGAG GTATT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGC ACTGGATACGACGAAACA
miR-511	CGCGGTGTCTTTT GCTCTG	AGTGCAGGGTCCGAG GTATT	GTCGTATCCAGTGCAGGGTCCGAGGTATTCGC ACTGGATACGACTGACTG