### SUPPLEMENTARY TABLE

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

| Target gene | Forward primer | Reverse primer | RT primer |
|-------------|----------------|----------------|-----------|
| XAGE5       | CTGGTTGGGCCT    | TCCGCATTCC      | GTCT      |
|             | TGCTTGAG        |                |           |
| COX7B2      | CCAGAAATGCAGC   | ACCCAAGTGAGA    | CAGAAAAG  |
|             | AAGCAGTCTC      |                |           |
| MAGEB2      | GCTGCGGAGAGTT   | TGGTTAGAGGAGCTT | CGCTTG    |
|             | CATCCA          |                |           |
| BHMT        | TGCTGAGGAGATT   | TCTGCTCTACGAGC | TAGAAGG   |
|             | GTGATTGGA       |                |           |
| PROM1       | AGTCGGAAACTG    | GGTAGTGTTG      | GGCCAAT   |
|             | GCAGATAGC       | GTCTG          |           |
| SLCO1B3     | TGGAGAACAGATA   | TGGTTTCG        | GAGGGAA   |
|             | CGGTAGC         |                |           |
| RP11-332K15.1 | ACTCTCGGGATCC  | AGTCTCTCAGTG    |            |
|             | TGGACCT         | AGTCTCTCAGTG    |            |
| XX-CR54.1  | TGTACTCTACAGC   | ATCAGAAGGCCAACC | TCAGGACTG |
|             | ACGGCATAAGG     |                |           |
| BC038578    | AGCCAGCGAGAC    | TAAAGTTGGG      | GGAATGC   |
|             | CACGAACC        | CTGAGGCTAGG     |            |
| LOC389023   | CGTGTCTGAGATT   | TAGGAGTTCCA      |            |
|             | GTGGCTGAGC      | CGTGGACC        |            |
| SSTR5-AS1   | GTGCCCTCAGTG    | GCACGCAGCAGCACT | CTCC      |
|             | AAGAGCAGAGC     |                |           |
| FAM99A      | GGCTGGCTCAG     | TCTCTGGTG       |            |
|             | CCGTGC          | CATCTGTG        |           |
| hsa_circRNA_103809 | TCGAGACCTCTGT | GCCAATACTCCACT |            |
|             | CAGCGAG         | GTCTTCTTCTCCTT |            |
| hsa_circRNA_101368 | GAATGG         | CTTCTTCTCCTCCTT |            |
|             | AAGACTTGAGGC    | ACACAGTGAATAGA |            |
| hsa_circRNA_102399 | GTGCAGGGGTCCG | GCGCTGTAATACGT |            |
|             | AGGTATT         | CCTGCT         |            |
| hsa_circRNA_000639 | ATGCCACAGCCTT | TCTCTTCTCCTCCT | ACATT     |
|             | TCCAA           |                |            |
| hsa_circRNA_001547 | ACATACCGTGTTTG | CAGGCTCATT CCACA |            |
|             | CTCTC          | TCAG           |            |
| hsa_circRNA_400010 | GTATAAGGATGACT | GCACCTCAACAAATCGT | CAGC |
|             | CACTGACA        |                |            |
| miR-532     | GCGTTCACACACC   | AGTGCAGGAGGAGGC | GTCGTATCCAGTGCAGGCGAGGTTATTCGAG | ACTGGATACGACTGCAAG |
|             | AAGG           | GTATT          |            |
| miR-130     | CGCGACTCTTTTCC | AGTGCAGGAGGAGGCG | GTCGTATCCAGTGCAGGCGAGGTTATTCGAG | ACTGGATACGACTGCAAG |
|             | CTGTTCG        | GTATT          |            |
| miR-642     | GGGGGTCCCCCTCC | AGTGCAGGGGAGGCG | GTCGTATCCAGTGCAGGCGAGGTTATTCGAG | ACTGGATACGACTGCAAG |
|             | AAAATT         | GTATT          |            |
| miR-329     | GCGGAGCTTTTCTCT | AGTGCAGGGGAGGCG | GTCGTATCCAGTGCAGGCGAGGTTATTCGAG | ACTGGATACGACTGCAAG |
|             | GGGTTTC        | GTATT          |            |
| miR-511     | CCGGTTGTCTTTTCT | AGTGCAGGGGAGGCG | GTCGTATCCAGTGCAGGCGAGGTTATTCGAG | ACTGGATACGACTGCAAG |
|             | GCTCTG        | GTATT          |            |