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| **miRNA** | **Stem-loop primers for reverse transcription** | **Forward primers for**  **RT-qPCR** | **Universal reverse primer for**  **RT-qPCR** | **Universal Probe Library probe #21** | **Mature miRNA sequences** |
| hsa-miR-26b-5p | 5' - GTTGGCTCTGGTGCAGGGTCCGAGGTATTCGCACCAGAGCCAAC ACCTAT - 3' | 5' - GTTTGGGTTCAAGTAATTCAGG - 3' | 5' - GTGCAGGGTCCGAGGT - 3' | 5' - TGGCTCTG - 3' | 5' - UUCAAGUAAUUCAGGAUAGGU - 3' |
| hsa-miR-451 (5’) | 5' -GTTGGCTCTGGTGCAGGGTCCGAGGTATTCGCACCAGAGCCAAC AACTCA - 3' | 5' - GTTTGGAAACCGTTACCATTAC - 3' | 5' - GTGCAGGGTCCGAGGT - 3' | 5' - TGGCTCTG - 3' | 5' - AAACCGUUACCAUUACUGAGUU - 3' |
| hsa-RNU-43 | 5' - GTTGGCTCTGGTGCAGGGTCCGAGGTATTCGCACCAGAGCCAAC AATCAG - 3' | 5' - GTGAACTTATTGACGGGCG - 3' | 5' - GTGCAGGGTCCGAGGT - 3' | 5' - TGGCTCTG - 3' | 5' - GAACUUAUUGACGGGCGGACAGAAACUGUGUGCUGAUUGUCACGUUCUGAUU - 3' |
| hsa-miR-223-3p | 5' - GTTGGCTCTGGTGCAGGGTCCGAGGTATTCGCACCAGAGCCAAC TGGGGT - 3' | 5' - GTTGGGTGTCAGTTTGTCAAAT - 3' | 5' - GTGCAGGGTCCGAGGT - 3' | 5' - TGGCTCTG - 3' | 5' - UGUCAGUUUGUCAAAUACCCCA - 3' |

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| **mRNA** | **Forward primers for RT-qPCR** | **Reverse primers for RT-qPCR** | **Amplicon lenght** |
| SELP | 5' - CCATTGTCTAGAGGGCCAGT - 3' | 5' - GGGCTTCCTGGATAGTCAATG - 3' | 119 nt |
| IL1B | 5' - AGCCAGGACAGTCAGCTCTC - 3' | 5' - AGAGGCCTGGCTCAACAA - 3' | 65 nt |
| DICER | 5' - TGTTCCAGGAAGACCAGGTT - 3' | 5' - ACTATCCCTCAAACACTCTGGAA - 3' | 76 nt |
| ITGA2B | 5' - AGCCTCAATGTGTCCCTACC - 3' | 5' - AGTCCAGGACGATTCGTGTC - 3' | 103 nt |
| RPLP0 (36B4) | 5' - ATGCAGCAGATCCGCATGT- 3' | 5' - TCATGGTGTTCTTGCCCATCA - 3' | 64 nt |

**Suppl. Table 3.**