**Table S1. Sequences of primers and probes used for reverse transcription and TaqMan qPCR.**

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| **Universal Reverse Primer** | 5’-GTGCAGGGTCCGAGGT-3’ |
| **hsa-miR-16-5p****(miR-16)** | RT: 5’-GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACCGCCAA-3’Forward: 5’-GCCCGTAGCAGCACGTAAATAT-3’Probe: 5’-(FAM)-GCACTGGATACGACCGCCAA-(FQ)-3’ |
| **hsa-miR-21-5p****(miR-21)** | RT: 5’-GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACTCAACA-3’Forward: 5’-GCCCGCTAGCTTATCAGACTGAT-3’Probe: 5’-(FAM)-GCACTGGATACGACTCAACA-(FQ)-3’ |
| **hsa-miR-19b-3p****(miR-19b)** | RT 5’-GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACTCAGTT-3’Forward: 5’-CGCTGTGCAAATCCATGCAA-3’Probe: 5’-(FAM)-GCACTGGATACGACTCAGTT-(FQ)-3’ |
| **hsa-miR-205-5p****(miR-205)** | RT: 5’-GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACCAGACT-3’Forward: CCTCCTTCATTCCACCGGA Probe: 5’-(FAM)-GCACTGGATACGACCAGACT-(FQ)-3’ |
| **hsa-miR-126-3p****(miR-126)** | RT: 5’-GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACGCATTA-3’Forward: 5’-CCGCTCGTACCGTGAGTAAT-3’Probe: 5’-(FAM)-GCACTGGATACGACGCATTA-(FQ)-3’ |
| **hsa-miR-25-3p****(miR-25)** | RT: 5’-GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACTCAGAC-3’Forward: 5’-CCGCCATTGCACTTGTCTCG-3’Probe: 5’-(FAM)-GCACTGGATACGACTCAGAC-(FQ)-3’ |
| **hsa-miR-125b-5p****(miR-125b)** | RT: 5’-GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACTCACAA-3’Forward: 5’-CGTCCCTGAGACCCTAACTT-3’Probe: 5’-(FAM)-GCACTGGATACGACTCACAA-(FQ)-3’ |
| **hsa-miR-183-5p****(miR-183)** | RT: 5’- GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACAGTGAA -3’Forward: 5’- CGCTATGGCACTGGTAGAA -3’Probe: 5’-(FAM)- GCACTGGATACGACAGTGAA -(FQ)-3’ |