**Supplementary Table S2.** Primers

| | Primer Sequence 5'→3' |
|---|---|
| U6-Forward | CTCGCTTCGGCAGCACA |
| U6-Reverse | AACGCTTCACGAATTTGCGT |
| miRNA consensus primer | GTGCAGGGTCCGAGGT |
| hsa-miR-100-RT | GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACcacaag |
| hsa-miR-100-AS | GCAACCCGTAGATCCGAACTT |
| hsa-miR-145-RT | GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACaaggga |
| hsa-miR-145-AS | GTCCAGTTTTCCCAGGAATCC |
| hsa-miR-182-RT | GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACtgtgag |
| hsa-miR-182-AS | AGCTTTGGCAATGGTAGAACTC |
| hsa-miR-210-RT | GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACtcagcc |
| hsa-miR-210-AS | TGTGCGTGTGACAGCGGC |
| FRAP1-R | TCCGGCTGCTGTAGCTTATTA |
| FRAP1-L | CGAGCATATGCCAAAGCACT |
| GAPDH-L | CGACCACTTTGTCAAGCTCA |
| GAPDH-R | AGGGGTCTACATGGCAACTG |
| hsa-mir-100-XhoI-F | CTCGAGAAAACAATTGAATGGTAGACTCT |
| hsa-mir-100-MluI-R | ACGCGT ACTCTTAAAGGAAACTAAGGGG |