| ID          | Sequence(5’-3’)                  |
|-------------|----------------------------------|
| GAPDH F     | ACCACAGTCCATGCCATCAC             |
| GAPDH R     | TCCACCACCCCTGTTGCTGTA            |
| hsa_circ_104566 F | GTCCATTGCTTTACCTGTGC       |
| hsa_circ_104566 R | CTGTCCAGCTGATAAAGGCG     |
| PSD3 F      | GTGCCCTGATGTACCATACCA            |
| PSD3 R      | GCCAATTCGGACGATGATTCTA           |
| miR-338-3p F | GCAGTCCAGCATCAGTGA              |
| miR-338-3p R | GTCCAGTTTTTTTTCAACA             |
| FOXP1 F     | TGACAAACAACCAGCTCTTCA           |
| FOXP1 R     | TGAGGGCTCAGCCTTTGTT             |
| U6 F        | CTCGCTTCGGCAGCAGACATA           |
| U6 R        | AACGATTACGAATTTGCGT             |