**S1 Table**. PCR primer sequences used in the current study.

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| Primer set | Primer sequence |
| Cloning primer | (BTB1-F) 5’-CAGAATACGCGAGTCAATCTACC-3’(BTB1-R) 5’-GGTGACCACAAACCCTCAA-3’(BTB2-F) 5’-TACCCTCGGTTCTTCTCAGT-3’(BTB2-R) 5’-AGCTGAGAGTTTCAACGTACC-3’ |
| 5’-RACE primer | (5’RACE outer) 5’-GCTGATGGCGATGAATGAACACTG-3’(5’RACE inner) 5’-CGCGGATCCGAACACTGCGTTTGCTGGCTTTGATG-3’(BTB1-5’outer) 5’-TCCTGAACCACATTCACCTCAC-3’(BTB1-5’inner) 5’-TCGGCGTAGCGAATATCTTTCA-3’(BTB2-5’outer) 5’-CGATTTCTGGCCTGATTCGTTTG-3’(BTB2-5’inner) 5’-ACAGTGTCGGTAGTTGGTCTTG-3’ |
| 3’-RACE primer | 3’RACE outer) 5’-GCGAGCACAGAATTAATACGACT-3’(3’RACEinner) 5’-CGCGGATCCGAATTAATACGACTCACTATAGG-3’(BTB1-3’outer) 5’-GGTCTGATCTCGGGTGTTTCAA-3’(BTB1-3’inner) 5’-CGGTGTGGAAACCTCTTTCACTA-3’(BTB2-3’outer) 5’-CATGTCGCAACAGAACAGGAAG-3’(BTB2-3’inner) 5’-GGAAAGCGGCGTCGAAGATAA-3’ |
| qRT-PCR | (actin-Frt) 5’-ACATCAAGGAGAAGCTCTGC -3’(actin-Rrt) 5’-CCTCGGGACAACGGAAAC-3’(GAPDH-Frt) 5’-CCGTTCATCGATGTCACCTA-3’(GAPDH-Rrt) 5’-GATGTTGGTGGGGTCAGACT -3’ (BTB1-Frt) 5’- ATCATCACGCTGGAAGAAGG-3’(BTB1-Rrt) 5’- GAGGGAAGAGCGAAGAGATTG-3’(BTB2-Frt) 5’-AAGATCAAAGGTCTCGCAGAA-3’(BTB2-Rrt) 5’-GACGACGAATGGTGTCCATTAT-3’ |
| dsRNA synthesis\* | (BTB1-dsF) 5’-TAATACGACTCACTATAGGGAGGTACAAGGGTGAGGTGA ATGT-3’(BTB1-dsR) 5’-TAATACGACTCACTATAGGGAGGAGTTCAGATCCGTCGA GTTATT-3’(BTB2-dsF) 5’-TAATACGACTCACTATAGGGAGCAAACGAATCAGGCCAGA AAT-3’(BTB2-dsR) 5’-TAATACGACTCACTATAGGGAGTTCGAACTCGGCTTCTTCT C-3’ |

\*T7 polymerase promoter sequence (**TAATACGACTCACTATAGGGAG**) was added to 5’ end of the oligonucleotide sequence to facilitate subsequent dsRNA transcription as described in manufacturer’s protocol (MEGAScript, Ambion).