Table S2. Information of primer sequence

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| Gene | GenBank No. | Sequences (5’-3’) | Length/bp |
| β-actin | XM\_024980020.1 | F: CTAGGCGGACTGTTACTGAGC  R: ATGTTTGCTCCAACCAACTGC | 239 |
| CDC45 | NM\_001161623.2 | F: TATACGCTGGTTCCGGTTTC  R: TCTTCCTGTTTCGCTCCACT | 235 |
| RPL9 | NM\_011292.2 | F: GACGTTCTCTCTTTGCCCCA  R: GTGATTGAAGTCCCTCCGCA | 152 |
| SGK1 | NM\_001161845.2 | F: GATGGGCCTGAACGATTTTA  R: CCCTTTCCGATCACTTTCAA | 225 |
| ID3 | NM\_008321.2 | F: GCCCGAGAGAAGGACTGAAC  R: CGACACCCCATTCTCGGAAA | 129 |
| FOS | NM\_010234.3 | F: TACTACCATTCCCCAGCCGA  R: GCTGTCACCGTGGGGATAAA | 113 |
| SCD2 | NM\_009128.2 | F: CGCCTTCCCCTACGACTACT  R: GGAACAGGAACTGCAAGACC | 201 |
| JUN | NM\_010591.2 | F: GCACATCACCACTACACCGA  R: GGGAAGCGTGTTCTGGCTAT | 127 |
| IER3 | NM\_133662.2 | F: CGGACTATGCGCTGGATCTT  R: GAAAGTCTCTGTCCGTCCCC | 222 |
| HMOX1 | NM\_010442.2 | F: GAACCCAGTCTATGCCCCAC  R: GGCGTGCAAGGGATGATTTC | 117 |
| BIRC5 | NM\_001012273.1 | F: TGAGCAAATTCTGCCCCCAG  R: CCATGTCCCCAAGCCTCTAC | 171 |
| SKA2 | M\_025377.3 | F: AACTGCAGAGATTGGGCGAG  R: ATGCATCACGATCCTCTCGG | 85 |
| DNAJB9 | NM\_013760.4 | F: TCTCGGATGCCAATAGTCGG  R: CTAGAACCATCCTGGCGTGT | 217 |