**Table S2 Primer sequences used for qRT-PCR.**

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| --- | --- | --- | --- |
| Accession number | Forward primer (5’-3’) | Reverse primer (5’-3’) | qRT-PCR efficiency (%) |
| SbCys10 | CTCTACGAGGCCAAGGTCTG | CAGCATCTTTGACCACAGGA | 94.5 |
| SbCys11 | AACATCGAATCGTCCCACTC | GCCTTCTTGTTGTGCTCCTC | 92.7 |
| SbCys15 | GAGCACAACAGCAAGACCAA | CACACCTTCTCCCACACCTT | 99.3 |
| SbCys14 | GGGTGCTGCTTTGGAGTTTA | TCCTCTTGTGCAATGTCTGC | 95.1 |
| SbCys17 | AAGCTCGTGTGCTTCACAAA | GGCCTTGTACAGCTTCTTGG | 94.7 |
| SbCys12 | TACAACCGCCAGCTTAGAGG | CGACGCGGAGGTAGTACTTG | 92.4 |
| SbCys4 | ACCCCATGGCTACTACGAGA | TGGAGGACGAGCCTGTAGTT | 91.7 |
| SbCys3 | AGGCCAAGTACCAGAAGCTG | TAGATGACGGCGACGTAGG | 98.8 |
| SbCys1 | CCGGTTGAACTTGACCTTGT | TGAGAACCAGCTTCCTGCTC | 93.6 |
| SbCys7 | TGAACGAGCTGGTGATCAAG | CCACACCAGGCAGTCGTA | 91.7 |
| SbCys6 | GTCCAACAGCGAACGACAC | TTTCTTGAACTCCAGCAGCA | 96.5 |
| SbCys5 | GAACCAGCCACCTTGTTGTT | TGAATTTTAGCCCGTCGTTC | 93.4 |
| SbCys2\_2 | TGGAAGCCGATTAATGTGAA | CCGTGTCTTGTCGACCTTCT | 97.2 |
| SbCys2\_1 | GACCCCGAAATCCAAGAGAT | CCGCGTATTGTCAACCTTCT | 91.6 |
| SbCys8 | AGCTCGACGGAACTCAACAT | AAGGACAAGAGCTTCCACGA | 93.3 |
| SbCys9 | ACCATGGCAGAGACAACTCC | GCGTCGATGATGAGGTGATA | 94.1 |
| SbCys16 | GAGCACAACAGCAAGACGAA | GCATGGAGCCCCTTAAAGTT | 96.8 |
| SbCys13 | GGCTCAAGTTCGACAAGGTC | CCGGCTTAAATGACAGGAGA | 95.3 |
| AB181991\* | GTGTCGCACCAGAGGATCAT | CGCTGGCATACAAGGACAGA | 95.7 |

AB181991\*: Accession number of reference gene β-actin