

Table S2 Primer sequences and product sizes for RT-qPCR data. ^a

| Locus #/ transcript # | Forward Primer | Reverse Primer | Product Size |
|----------------------------------|--------------------------|---------------------------|-------------------------|
| 58798/1 | AAAGATCTGCTGGCTGACCCTGA | AACACCAACCAATTGCCTCACCC | 134 |
| 102514/1 | AAGTGACCCGTGCGTTCTCAA | TGCGTTTGGGTTGGGAATGTGT | 148 |
| 95010/1 | TTTGACATGCTGCGGAACTGCT | AATGGCCACGACGTGTTGAAGG | 225 |
| 125065/1 | TGTCAGTGGCGTTGCACAGTCTT | ACATTGCCAATTCTTGCGCGGT | 159 |
| 77179/1 | GAAATGGCGGAAAAAGCATA | GGTGAAATTGTGTCCATC | 225 |
| 95925/1 | CAAAGCCTGGACATCGACGCAA | CAATGACACAGGCCCGCAGAAA | 194 |
| 86800/1 | AGCTGGAGGGAAGGCACCAATAA | TGGGAGCTGTCAATCAACTGGGA | 110 |
| 65589/1 | TTTGCCGGGAACACGTGCATT | TGAGCGCCGAGTGATGTAGGA | 177 |
| 70728/1 | ACCAACGGATTCCCATTCTCGTCA | TTTGCGGGCAGCAGTGTGT | 110 |
| 95114/1 | TGTCGCGCTGTTCCTTTGTT | TGGCCAAAGCAAGGCGTTTGTGA | 187 |
| 101012/1 | GGTCAGCACGCATGAAAGCATTGT | AAGCAATCCAGATGGCAAAGGCAG | 171 |
| 66644/1 | TCCAAGACCAAGTGTGGTGGACT | TGATCCAAGTCAGGGACAGGCAA | 110 |
| 101000/1 | TCTGTCTGGACACTGCTGTTGA | ATCCAACCGAATTCTCCGTGGT | 189 |
| 105631/1 | ACCGTGAACACTTCTTGAGAGCCA | GCCTCGGTTGAATGCTTTGTTCTGT | 210 |
| 125822/1 | ACCTCGCGCCTTGCTTATCAA | AATGGGACTGTTAAGCGGTTCTGT | 225 |
| 27493/1 | GGTTTGCTGCATCTCACAGGTCA | AGAAACAGCTGGCGACTAAGCTCT | 133 |
| 12296/1 | AGCCAAGGTCTTGAGCAGCTTA | TTGGGCCTTGACAGTACAGTGAACA | 125 |
| 119098/1 | ACTGCAGTCCACGATGCTATCCTT | GTCTGTTGTGCTTTGTCGAGATGC | 125 |
| 12335/1 | TGAAACCTCCTTTCAGCCTCCCA | TCACTTCACTCATCTCGGCAGCA | 172 |
| 84201/1 | AGCAGTTGGTAAGTCTGCACAA | GTAACCATGGTAGCAGCATGAA | 105 |
| 58671/1 | AACAGCTTTGGCAGCACCTGTAGA | TGCTTTCACAGCAACCCAGAAGAC | 114 |
| 77428/1 | AAGGCAAGCGGTAACGAGGTTT | TGCTTTCCTTCTGTCAGCCAGT | 177 |
| 21845/2 | TCATGGCAAGGACGACGAGTGAA | TCACCCATGGCAGTAAAGAGCGA | 156 |
| 59465/1 | TCGGCAGGATTGTGTCCAAGTGA | AAACGAGCGACACAACGTCAGCA | 197 |
| 13527/1 | AGACACCCAAGTTCCTTCCCA | ACACGCCGTAAGTAAACGCCAA | 212 |
| 12461/1 | AGCAAAGGGCACGAACAACCAAC | TTGACTCGTATGGCCGCTAACA | 125 |
| 431/2 | TGGCCTTCAACAAACCTTACGCT | ACGTTTGTAGTCCCAGCCAGTCA | 238 |
| 1568/1 | AAGTTCGTTGGAGGGTACTGCGA | CCACCAAAGACTTCACACAGCCA | 110 |
| 20440/1 | AATGGCGGAGTTTGTCAAGACGG | TGCCGATGCATTTGCTGAGTT | 118 |

^a Transcripts are listed in the same order as in Table S1.