**Supplemental table 2** Primer sequences for 3’-RACE analysis

|  |  |  |
| --- | --- | --- |
| *Gene* | *GSP2 (5′ to 3′)* | *NGSP2 (5′ to 3′)* |
| *EjMYB3* | AGCCGTTTCAGTCCAAACCAACTCG | CAGGTTGAATTTCAGAGCAGCGAAGC |
| *EjMYB4* | CCAACCAATTGGACTCGACCCAGAG | ATCATGCAGCAGCAGCCCAATGTAG |
| *EjMYB5* | GGACCGGAAAAAGCTGCAGATTGAG | CTCCGCATCTCCTTCTCTGTCACCA |
| *EjMYB6* | GAGAAGGCCTCTGTGGGATCCAATG | AGAGTTCCTGCAACCGTCCATGTCA |
| *EjMYB7* | GCATTCTCCCATCCCAGCTGCTAAA | CACCAACACTTCGGAGCCCATTTCT |
| *EjMYB8* | AGCAGGGAAGAAGAAGACGCCATCA | CATATGTCACAACGAGCGACGACGA |
| *EjMYB9* | GGGAAACCCCTGATCGTTGGTGTAA | CTCCAGCGGTAGACACAGCACCAAT |
| *EjMYB10* | AAGGGGTGCCATGGACAGAAGAAGA | GAAGTTGGCCAGCATGAGCATTGAC |
| *EjMYB11* | GAACTTGCAAGGGCCGAACTCACTC |  |
| *EjMYB12* | CATTCGTTTGAAGAGGCCGAGGATG | AGGATGCTCACCCTTCAGCCTTCAA |
| *EjMYB13* | CCACACTCTCCAACACCAGCTCCAC | TCGGATAGCAGGAGCTGGAAGGGTA |
| *EjMYB14* | GCAGTGGAAAGTTCAGGAGAGAAGCA | CTGCAGTTCACGGAAGGGGTTTGTG |