

TableS1.Primer sequence for qRT-PCR analysis

Gene name	accession number	Forward primer(5'-3')	Reverse primer(5'-3')
<i>BnPIP1;1</i>	XM_009118558.3	5'-GCTAAGAACAGACCAAAAAGTG-3'	5'-GGCATTCGGTGGTATGATA-3'
<i>BnPIP1;2</i>	XM_013892758.2	5'-GAATGGTCCAACCCAAAAG-3'	5'-TGTTCCCATTCTTGCACCT-3'
<i>BnPIP1;3</i>	XM_009120575.3	5'-CTGGAAGCCTTTGACAACG-3'	5'-CATAAACCAGCGGTGACA-3'
<i>BnPIP2;2</i>	XM_013836821.2	5'-GGAGTCTCGGGCATTCTT-3'	5'-CGATGGATACAGCACAGGG-3'
<i>BnPIP2;5</i>	XM_009105687.3	5'-GTGTAGTAAGAAGACTGGAAGGA-3'	5'-GAAAGTGACGTTGGTGAGG-3'
<i>BnPIP2;7</i>	XM_013815142.2	5'-TGGGATGGTAGCCAAATGC-3'	5'-TTCTCCGCGACTGACCCT-3'
<i>BnTIP1;1</i>	XM_009143362.3	5'-TTTTTGTCTTTTGCCTTTGTG-3'	5'-GGAAATCAACAGAAAAGGATTCA-3'
<i>BnNIP5;1</i>	XM_013880108.2	5'-ACGTCCCAGATCCATTTTCTT-3'	5'-CCAGGACTAGTTTCACCAAACAG-3'
<i>BnNCED</i>	XM_013732216.1	5'-GTCGCGTCAACCTCCAAGCT-3'	5'-TCTGTTTCTCCCCGGAGAGG-3'
<i>BnNAC55</i>	XM_009124259.3	5'-CACTCGCACCGCCTCAAAAC	5'-CCCATGGACGCAGAAAAGAATAAG-3'
<i>BnERF2</i>	XM_033280981.1	5'-GGGTTTGGTTAGGGACGTTTG	5'-GGAACCACGCATCCTAAAAGC-3'
<i>BnRbohF</i>	XM_013825233.2	5'-AGAACGTTGAAGGGTGGAAC 3'	5'-AGCTCCAATGCAAACCTCCTT-3'
<i>BnRbohD</i>	XM_013788801.2	5'-TGTAGAGCCGTCTCTCCCTT-3'	5'-ACGGTCCTGAGCTTACGAGT-3'
<i>BnMYB108</i>	NM_001315555.1	5'-GCCTCCGCCGCATCATTA-3'	5'-GTTTCGGTTTTGGAGCGTTGTAGTA-3'
<i>BnWRKY40</i>	XM_013732543.1	5'-CATTCACTCGTTTAATTCGTCCAC-3'	5'-CGCGCATATACGTACATACTCACAA-3'
<i>BnADH</i>	XM_013832947.2	5'-CTAAGTTCATCATGTCTACGACCGG-3'	5'-TCGGTGTGACAGAGAGAAGTGAAG-3'
<i>BnLDH</i>	XM_013886552.2	5'-ATTGTGTAATGGCGAACAACGG-3'	5'-ATGACGGAGACCTTGGTGCA-3'
<i>BnPDC</i>	XM_013852099.2	5'-TGGAGGGTACACCATTGAAGTGG-3'	5'-CGCAGTCCAGCATTTACCTTCTC-3'
<i>BnGAPDH</i>	XM_013801897.2	5'-TTAGACTCGAGAAAGCTGCGACC-3'	5'-CTGTTGTACCAACGAAGTCAGTTG-3'
<i>BnGLB</i>	XM_009124578.3	5'-ATTGGTGAGGACACTGAAAGAA-3'	5'-TCTCGGCCTTAATGGCTAAAG-3'
<i>BnCPK12</i>	XM_013869200.2	5'-AGTGGCTGCATTCTTTCTTTGAC-3'	5'-ATCCCTCCACTCCCATTTCCTT-3'
<i>BnCPK11</i>	XM_009146872.3	5'-TGGTGGCAGCATTTTCGTATTTT-3'	5'-TTCTCCCAACTCCATCTCCTTTCTT-3'
<i>BnPPR</i>	XM_013831783	5'-TGGTGTGCGATAAGTGTGAGA-3'	5'-GGTGTCCATCTGTTCTTCTTGG-3'
<i>Bnactin7</i>	XM_013858992.2	5'-CCCTGGAATTGCTGACCGTA-3'	5'-TGGAAAGTGCTGAGGGATGC-3'