

Primer Name	Sequence (5'-3')	Length (bp)
Mt244-23-F3	GGACAGTTTTCTGCAGTCAT	20
Mt244-23-B3	GCCTATTGTGTCTTATCAATGT	23
Mt244-23-FIP	ACAAACAGATCAAGTCCCCTCAAACAGAACAAG TGAAAATATTGC	46
Mt244-23-BIP	ATTCCAGGAATACAAATGAGAGGAGCACTGTGCA GATTGGAAAT	44
Mt244-1-F3	GCATTATCATAAAATGAAAGGAGA	25
Mt244-1-B3	TTTGGTCCAAAGGCTGTG	18
Mt244-1-FIP	ACTGCAGAAAACGTCCTTAGAAAATTTACGAA GGCTTCAAC	44
Mt244-1-BIP	ATTACCTTICATTGAGAGGGACTTGCTGGAATTGCA GGCACTG	42
Mt244-1-FLOOP	ATTGTCGTAATAAGCTGT	19
Mt244-1-BLOOP1	ATCTGTTGTAAATAGACGTCG	21
Mt244-1-BLOOP2	TTGTAATAGACGTCG	15
Mt244-12-F3	GGAAAGGAGATTTTCACGAA	20
Mt244-12-B3	TTTGGTCCAAAGGCTGTG	18
Mt244-12-FIP	ACTGCAGAAAACGTCCTTAGAAAGGCTTCAACT TCCAAAATCAC	46
Mt244-12-BIP	ATTACCTTICATTGAGAGGGACTTGCTGGAATTGCA GGCACTG	42
Mt244-43-F3	GAACAAGTGAAAATATTGCTCC	22
Mt244-43-B3	GTIGATGCCTATTGTGTCTT	21
Mt244-43-FIP	TCCTGGAATTGCAGGCACTGATATTACCTTATTG AGAGGGA	42
Mt244-43-BIP	CAAATGAGAGGAGGCACAGCCTCTTTAACACTG TGCAGAT	42
Mt244-43-FLOOP	CCGACGTCTATTACAAACAGATCAA	25
Mt244-43-FLOOP2	CCGACGTCTATTACAA	16
Mt244-43-BLOOP	GGTTTCAGTATCATGGAGGT	20
NBS5-LRR-8-F3	GAACGAGAGCGAGACGGA	18
NBS5-LRR-8-B3	CCATCTCGACGCACACCT	18
NBS5-LRR-8-FIP	TCGTCGAGCAGGACAACGAACCTGAGGGAGCGATC CACCTT	40
NBS5-LRR-8-BIP	AGTTCCAGTTGGCGGACGTGAGGATCAGCTTACA CCCGT	39
NBS5-LRR-8-FLOOP	CTCCTCAGGGCCCTGCG	17
NBS5-LRR-8-BLOOP1	AATCCCCGCGCCAAGCA	17
NBS5-LRR-8-BLOOP2	CCGCGCCAAGCA	12
NBS5-LRR-6-F3	GATCATCGCCAATCGGCT	18
NBS5-LRR-6-B3	CCATCTCGACGCACACCT	18
NBS5-LRR-6-FIP	TCTTCTCCTCAGGGCCCTGGAACGAGAGCGAGA CGGA	38
NBS5-LRR-6-BIP	AGTTCCAGTTGGCGGACGTGAGGATCAGCTTACA CCCGT	39
NBS5-LRR-6-FLOOP	AGAAAGGTGGATCGTCCCT	20
RPS2-1-F3	GGTTGTTGCTCCAAGGGC	18
RPS2-1-B3	CAACCTCTCGGACAATGA	20
RPS2-1-FIP	ACAGCCACATCGTCTCATCTTCAGGAGCAGCAA GAGATCG	41
RPS2-1-BIP	AGATTGCCTCCTGTCTCTCCCATAGAGTCGGCATC TCCGC	40
RPS2-1-FLOOP	TCTGCTCTGATGATGCTTAT	20
RPS2-1-FLOOP2	TCTGCTCTGATGAT	14
RPS2-1-BLOOP	AAGAGGGACCGACAACGTGG	20
RPS2-5-F3	TGACGACGATGTGGCTGTA	19
RPS2-5-B3	CGCCTACTGCATCATCTGC	19
RPS2-5-FIP	TAGAGTCGGCATCTCCGCCGAGATTGCCTCCTGTC TCTCC	40
RPS2-5-BIP	CCTGGTCTGCCAGGGAGTTCGTCTCAGGCACA CCTC	38
Mt151-131-F3	CACAGAGGAGGAGGTCTT	18
Mt151-131-B3-AI	TGATACAATAACCTCTGTATGT	23
Mt151-131-B3-T	CAGATCAGGATGAATACTCTGTATG	25
Mt151-131-FIP	CGAGGAGGGTCTCCACCATTAGGATGTATGGTTT CTTGAAC	42
Mt151-131-BIP	GCTGGGCTGGACTATGTCTCAACTACAGGAAAAA AGTGCAAAC	43
Total		1,365