

Table 8. Primer sequences used in this study

Locus	5' flanking primers	3' flanking primers
INT01	5'-CTTATGTAACGTTTTTGGTCATGTAARGT-3'	5'-TTATGATGTGGTTATTAATGATTATGAAA-3' 5'-TATTAATGATTATGAAAARGCCAAGTCRCT-3'
INT0657	5'-AGAGCTCCYACAGACACCCTGATG-3'	5'-GAAAGTAATGGCTGATGTGGTTCCAGKTCC-3'
INT0990	5'-ATCTACTGCCAAAACCTGTGCCTKTTG-3'	5'-RAGCATCTTGCCATAGCCCTGTCTCAT-3'
INT1026	5'-GATRCTGTACACATTATCTCGACCCA-3'	5'-TGCAWCAGACCATCCTGTGCCATCAG-3'
INT1067	5'-GTTACCTCYCTGAAAAGTATATAATTGGTTGCTAA-3'	5'-GACTCTGYACATCTATCCCATGACTCTC-3'
INT1075	5'-TTCCTTTGATCGAACCTGTGACCACAT-3'	5'-TGCTGGGCAGGTACTGRTAGCTG-3'
INT1095	5'-AAGCSATTTAACTGGWGGTTTCACAAGAA-3'	5'-CAAKKGACTTTGAAAAMAGCATGAAGAGATATTTG-3'
INT1111	5'-AGGAGGCCAGGCATGCTGTACAAAA-3'	5'-ATAYTTAACACAGCCAATTTACAAGGGAGGAAG-3'
INT1124	5'-TGCCATRACCTCAGACATGATCTCTCCA-3'	5'-TTACGARAAAAATGCTTAAGCTGCGMCGVGAT-3'
INT1167	5'-ARGGTCTTCCGTGTCWGSCTGGTACAG-3'	5'-GMWGACCATCTGATCCAGGGTTTCCC-3'
INT1189	5'-TTCACTTCCGAGAGAACKGATCGCT-3'	5'-GAACARTTRAAAGAGATYCCATACACTAAARTAGAGT-3'
INT1068	5'-TGCTRAAGAAGCCAAACTATAAGAARCTAGAACT-3'	5'-TTCCATCTTCTAGDACTTAAATTTGTTTTCCT-3'
INT1098	5'-ATGGHTGTGAAAATGGYCGCTGTGTGAG-3'	5'-ACTTGGCRTTCTTGACAGAGACATCCG-3'
INT1122	5'-AGAAGGAGTTGGAACCAAGGTGATGGAC-3'	5'-CTCCATATACGGCCACTGATCGAAGATTAT-3'
INT1151	5'-CASKTCTTCAGATTGCGTTTGGTAGTT-3'	5'-AAATGAACAAAAGAGAATGAAGRTRTGMAAATCA-3'
INT1233	5'-AAGTGATGGGCCCTCAGAGACCTAGAT-3'	5'-YAAAAGGTATTTCTTCTCCAGTATCCAGGTTCTTA-3'
INT0904(_Af)	5'-CTTTTCTCTCTTGATGTGCAAYGGTTCAT-3'	5'-TACTGGAGAATTTGGCTGARGAYAACCTT-3'
INT0957(_Af)	5'-GTGGCAATTTGAGAAATCYAGACAA-3'	5'-GTGGAGCCAAYACACCTTATGARAAAAG-3'
INT0979	5'-GTTTAYTCTGTGCWGGCTACTGTGTGG-3'	5'-GCACTCTCTCTTATTRATGCCAGGAAAC-3'
INT1060	5'-CCATKATGANTGCTGGGTTCACTCCA-3'	5'-GCCCCAATTTACTATGCAAATAGCGACTT-3'
INT048	5'-GTGTTTCACCAATAGARTCTCCACCTCG-3'	5'-CAGGCTCAAAAATATGGGATCCTACACC-3'
INT0964	5'-TGTCTCGTCATAAGCATCTAGCCAAGGA-3'	5'-TSACTAACTCTACCACTGCATTCCCACCATA-3'
INT276	5'-GCYGTGTTAGGTTTGTAGAAGATTGGATTGTC-3'	5'-CTATHATTGAGGCCATGCCAGTCCCTTGA-3'
INT0894	5'-GTGBTCATAAGCCACAAATTCATTGAAGAA-3'	5'-CAAGAAGAYCCCAAYTCTATWATTGCAGGCAT-3'
INT251	5'-ATAGTAGAGAACACCCAGYTCATTGAGGG-3'	5'-ATGTTYGACTACCCACAGCTGAGTTTGA-3'
INT285	5'-GAAATAGAACTATTAARCGGCARCAGAA-3'	5'-TTTTTAAGATATGAACAAGCTGAACATTTTAGGAA-3'
INT315	5'-AGTAAACTTGGGTATAACTGTTGTGTTGTT-3'	5'-TGAGATTKCCTTMMACCTTCACAACTC-3'
INT713	5'-TGTCAGTYCCTTCATGRTAATGTTCAA-3'	5'-ATCTTATTTAGTCCACCATGGGTACTGT-3'
INT810	5'-ACTYTCCATCCCTTTGCTGGCATA-3'	5'-AGCACCACAGTTATCCAAAATGTAATA-3'
INT875	5'-CACTRAAFAARATGATGGAGGTGGATGT-3'	5'-ACCAACATTGCTGAGACATCCATAACCA-3'
INT942	5'-TTAGCAACCAAAGCAMGAGAGGGTAAACT-3'	5'-GAYAGGATCTATBACACCATGAGGRAACT-3'
	5'-TACTCCTGCAGGACTCATCACACCTATTGT-3'	5'-CCTCTGAAGCACCAATTGCCAAAATAC-3'
INT980	5'-ATGGAGAAGAGCTTGTAAAAATTGAAAAG-3'	5'-TTAGTTTAGATTCTCGATAAGGAACATGAG-3'
INT999	5'-AGAAGGGCTTTTTAGGAARTCAGGATCTGTT-3'	5'-GAAAAGTGCTTCATGCAAATCAGCTGG-3'
INT138	5'-GTGGCGAGTAAGAAAAGAAGCYATGATGGG-3'	5'-AAGCATTTCATTCTCTCTCTGTTTCCAGGT-3'
INT226	5'-GCAGTTCGWCCYACTCGATGGATRTAATC-3'	5'-AGCAARCGCCTAGGATCCCTTAATAAGTT-3'
INT237	5'-GGTTAGATACATCGGCATGATCCTTYCTGGT-3'	5'-ATATCACTACCCCATCCCTGACYTGACT-3'
INT165	5'-ATCTGATGTCYAGTGACTGAATTTCTGAA-3'	5'-TTGCAGATAAAATGGTACTGTGACAGAAAATA-3'
INT265	5'-AGGCTCTCCMTGRCCACAGCAATCA-3'	5'-CTGTCTTTTACATCCATGTTGGGACCA-3'
INT382	5'-CATAAAATGTGTATCTATCCATAAAGRTGG-3'	5'-TAGCTAGTAGCCAAGTGGGAGAAGATTC-3'
INT391	5'-AATGYTTCTGCTTYCTCAAATGTATGGAG-3'	5'-ACCTGTACAATCCTYTCAAACATATGA-3'
	5'-ACCCYAAAGGAGCCATGCTRACCCAT-3'	
INT189	5'-TGGCTATCATCATARCCAGTTGTAAGGTC-3'	5'-AGAATTTGTCTTTGTCTCAGCARCTAAA-3'
INT283	5'-GAAYGTGGTTGAAAACTGAATGCCAATGT-3'	5'-CTTGGCCCTTGACCATTACTCTGGCTC-3'