

SNP marker	Chr	Forward Primer (5'-3') Reverse Primer (5'-3')	Enzyme	PCR product size	Site	Confirmed fragment sizes	Acc
2828298	I	TCCATTGATTCTCTGGTAAC <u>CCG</u> GCTTTGAGCCACACTAAGCTC	<i>Sma I</i>	300	20,80	100,200 200,80,20	<i>ebi-1</i> Ws-2
2833639	I	TGAAACCCATTTGTCACGTC TTTTGAAGCTTCTAAAACATCTTGTC <u>AG</u>	<i>Alu I</i>	363	335	363 335, 28	<i>ebi-1</i> Ws-2
3976656	I	TCTTGATCGGAAAGGAATGG TTCTCCTTCTCAAAC <u>CGAGCT</u>	<i>Sac I</i>	357	355	357 355, 22	<i>ebi-1</i> Ws-2
4211791	I	TACAAGACGGTTTCCCGCACGTGCCT <u>I</u> GCAGTTGATTGCGCTACTGA	<i>Pst I</i>	300	27	300 283, 27	<i>ebi-1</i> Ws-2
6411	V	CTGGCCCAACGTTCTTTGCAAACCTCCGGTGA <u>ATT</u> GACCCAAGACTGGAGAACCA	<i>EcoR I</i>	266	32	266 234,32	<i>ebi-1</i> Ws-2
169909*	V	CTTCAATTCCTTCAGGACAG TATGCGCTATGAAATTTGTG	<i>Hinf I</i>	173	153	153, 20 173	<i>ebi-1</i> Ws-2
639679 ^a	V	ATCTTTTGGCTCCAGGGCCAGATCACAGT <u>G</u> TCTA CCACCTGCTGAGAAGGAGAC	<i>Acc I</i>	215	35	215 35,180	<i>ebi-1</i> Ws-2
1694415 ^b	V	GGACACCTATGGTAGCCACAACCAAGAT <u>I</u> ATGCTTAATTGCGGGAAACA	<i>Mbo I</i>	185	28	185 28, 157	<i>ebi-1</i> Ws-2
1694415 ^{b*}	V	CCACAACCAAGAACTTTAT ACATGTATGCGAGAGAGGAT	<i>Apo I</i>	159	14	14, 145 159	<i>ebi-1</i> Ws-2
1938340	V	AATTCTTCCAATGGTTAAGAGTC <u>G</u> A CGGCAACAGATTTTAATTTT	<i>Sal I</i>	240	23	240 23, 217	<i>ebi-1</i> Ws-2
2363266	V	ACGGGCTAGTTTCTCCTGCT TATGGTTTTAGCAATTTCTGTGTTGCAAAC <u>CCGG</u>	<i>Sma I</i>	360	325	360 325, 35	<i>ebi-1</i> Ws-2
2388648	V	GCGTCAATCCTGGAATTTTGGTAGGTCC <u>CGA</u> TGCATCTACATGGGAAC TTATTT	<i>Mbo I</i>	273	32	32, 241 273	<i>ebi-1</i> Ws-2
2586630 [^]	V	ATAAACTTTATCTTTGGATTCTTCTTTGTATA <u>A</u> AGGATTTGCGCATGCCTAT	<i>Acc I</i>	354	32	354 322,32	<i>ebi-1</i> Ws-2
2895501	V	TACTCTTTAGTGAACCTCAGACCTCAC <u>GCG</u> TTTGAGTTATGAATTATTTGTAGCGTA	<i>Mlu I</i>	189	30	189 159, 30	<i>ebi-1</i> Ws-2
3263815 [^]	V	GCAATGGCTGTCCAAAATCT ATGGAAGAAAAATACCACAATAACAAAATTAC <u>CC</u>	<i>Sma I</i>	304	272	304,272 272	<i>ebi-1</i> Ws-2

4174053	V	AGGAATGATCTTGATCCGGAAAGACGGCGTTC CCGGTTACGAAGCAGAGTGT	<i>XbaI</i>	224	33	33, 191 224	<i>ebi-1</i> Ws-2
4318679	V	AGCGACACCGTTTTGGTAAG CTAAAATCACTCACTGCCTCTTCG	<i>Taq I</i>	198	65,167	65,103,30 65,133	<i>ebi-1</i> Ws-2
4507737	V	AATCACTCGGCTTCTGCAAT GTTTTGGATCTCTTGCATCATCTTCTACGTCTGCA	<i>PstI</i>	210	177	210 33, 177	<i>ebi-1</i> Ws-2
4615934	V	ATAGCTATGGCCCTAATCATAAAGCT GCCAATACAATGACATCACA	<i>HindIII</i>	246	23	23, 223 246	<i>ebi-1</i> Ws-2
5883034^	V	AGGACACGGTCGAATATGTTA GCTATACGAATTTGCAAATTTTATACATTTTC	<i>EcoR II</i>	368	219	368 368	<i>ebi-1</i> Ws-2
6337369	V	AACCGTCGAGTGTGATCG CTCGGAGCTTCAAAAATTGG	<i>Nru I</i>	200	20, 60	120,60,20 60,140	<i>ebi-1</i> Ws-2
6383466	V	AAACGATAGTTTACGTGTATAGAATT CCTGCGAAATTATATGGAAA	<i>EcoRI</i>	206	23	206 23, 183	<i>ebi-1</i> Ws-2
6855641	V	AGGCGTAAGAACTGCACTTGAAGTCGA ATTTTCAGCTCAAGCACCTG	<i>SalI</i>	200	25	200 25, 175	<i>ebi-1</i> Ws-2
7283071	V	AGCTGCCATTGTCTGTATCT AATTGTTTCCTGCTACCTCTGACA	<i>SpeI</i>	246	220	246 26, 220	<i>ebi-1</i> Ws-2
9193463^	V	CGAAATATAGCATTCTGAATTTTGTTCAGC CGAAATATAGCATTCTGAATTTTGTTCAGC	<i>Alu I</i>	349	30	349 349	<i>ebi-1</i> Ws-2