

Information on primers in 15 nuclear simple sequence repeats(nSSR) loci

Locus	Ta(°C)	Motif	Primer Sequence (5'-3')	Allele size (bp)	Reference
QrZAG30	56	GA	TGCTCCGTCATAATCTTGCTCTGA GCAATCCTATCATGCACATGCACAT	160-208	Kampfer <i>et al.</i> 1998
PIE271	56	AG	CACACTCACCAACCCTACCC GTGCGGTTGTAGACGGAGAT	197-247	Durand <i>et al.</i> 2010
QpZAG9	56	TG	GCAATTACAGGCTAGGCTGG GTCTGGACCTAGCCCTCATG	220-274	Steinkellner <i>et al.</i> 1997
QrZAG112	56	GA	TTCTTGCTTGGTGC CGCG GTGGTCAGAG ACTCGGTAAGTATT C	98-116	Kampfer <i>et al.</i> 1998
GOT011	56	GA	CCCCACCGTCTACTCTCAA GCGTTCACCACGTCCATAAT	197-225	Durand <i>et al.</i> 2010
GOT012	56	GT	TGATGATCCC AAACCACAAA AAGGCTGCAGGACTTTCAA	204-254	Durand <i>et al.</i> 2010
GOT021	56	AT	AGAAAGTTCCAGGGAAAGCA CTTCGTCCCCAGTTGAATGT	110-150	Durand <i>et al.</i> 2010
GOT040	56	TC	AAGGCACTCGTCGCTTCTA ACCGATTGAAGCTCGAGAA	242-298	Durand <i>et al.</i> 2010
PIE163	56	TC	GAGAGGCATGTGGAACCAAG CAAGCATAGGTGGTGGAACCC	230-264	Durand <i>et al.</i> 2010
FIR026	56	GT/GA	CTTCATGCACCAATT CCTCA GGCCATGTATGTGTGCAAAA	201-215	Durand <i>et al.</i> 2010
POR025	56	AG	CACACAAACCCATATGATCTGAA TCTCTTCGATCCCTTCTGC	105-145	Durand <i>et al.</i> 2010
FIR015	56	GT	ACCCTAAAACCCCAATCACC CGGATCTCGGCTATTCTTG	112-148	Durand <i>et al.</i> 2010
QpZAG16	56	TC	CTTCACTGGCTTTCCCTCCT TGAAGCCCTTGTCAACATGC	131-189	Steinkellner <i>et al.</i> 1997
QpZAG110	56	AG	GGAGGCTTCCTCAACCTACT GATCTCTGTGTGCTGTATT T	193-267	Steinkellner <i>et al.</i> 1997
QmC02241	56	GA/TC	TCAGTGACCACACGTCACCTCTC GTTTCTGGCCATGTTTGATGG	169-211	Ueno <i>et al.</i> 2008

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