

Supplementary File S1.

No.	Name	Sequence	No.	Name	Sequence
RAPD primers					
1	A02	TGCCGAGCTG	11	B07	GGTGACGCAG
2	A03	AGTCAGCCAC	12	B09	TGGGGGACTC
3	A08	GTGACGTAGG	13	B10	CTGCTGGGAC
4	A09	GGGTAACGCC	14	C02	GTGAGGCGTC
5	A15	TTCCGAACCC	15	C06	GAACGGACTC
6	A16	AGCCAGCGAA	16	C07	GTCCCGACGA
7	A17	GACCGCTTGT	17	C08	TGGACCGGTG
8	A18	AGGTGACCGT	18	C10	TGTCTGGGTG
9	B03	CATCCCCCTG	19	C14	TGCGTGCTTG
10	B04	GGACTGGAGT	20	C16	CACACTCCAG
ISSR primers					
1	814	(CT) ₈ TG	8	HB8	(GA) ₆ GG
2	844A	(CT) ₈ AC	9	HB9	(GT) ₆ GG
3	844B	(CT) ₈ GC	10	HB10	(GA) ₆ CC
4	17898A	(CA) ₆ AC	11	HB11	(GT) ₆ CC
5	17898B	(CA) ₆ GT	12	HB12	(CAC) ₃ GC
6	17899A	(CA) ₆ AG	13	HB13	(GAG) ₃ GC
7	17899B	(CA) ₆ GG	14	HB14	(CTC) ₃ GC
NUCmer primers					
Rx1	F10337514	TGAGAAGGGTTTAGTTTGCAC CTGTGAATAGGCAGAAAGGTC	Rx18	M12975375	TGATGGACTGGACTGTGAC CCATTTCTCTGATGCCTGC
Rx2	F11806232	TGCTTGTAAAGGGAGCCACG ACCCACCCAAAAACAGAGAAG	Rx19	M13446011	GCCACAAGAACAAGATATTGC GAGGAACCCATTGATAAAGC
Rx3	F12098636	TGAACTCAAATATCTGGAGCTG CCAAGTAACCAATGGAGATAC	Rx20	M15402427	ATGTAAATCGGGTAGTGAATCG CAACATAACAACCACGAAGTTC
Rx4	F12109005	GCCTCGAAATGACTGCAAG CATGCAGCACTGAATCTCG	Rx21	M1704617	GAAAAAAGCTCTTCTTTCCGC AGGCATAAAGTGCCACAAC
Rx5	F20713033	TGAAATTGGTGCTCTTCCAAC CCTCGTCAAGATTTGTTTCCG	Rx22	M17432556	AGAACAGGCAAGAATGTCTATC CTTCTCGAACAGCAGTAGC
Rx6	F2310454	CGAGGAAATCAAATGGAGCATC CCATAGACTGCGAAAACCC	Rx23	M19428070	GGAGTTTAACTACGGTAGGG ACCGTGCATACTCAAAG
Rx7	F25581559	GGTTACAGATGTGCAGGATG ACAAGATGCGTGAGAGCAG	Rx24	M21206660	CAAACCAAAGGATGAACTCAC TGATTAATGTTTCTTTGGCCC
Rx8	F26050110	GCTTAAGATTTGATGCTCTTGC GAATCTTTCCCTTGCCCTG	Rx25	M22110263	AGTGGGTTTGGGGGATTTG TTCAGTACAACCTACCAAGCAC
Rx9	F36548753	CGAGACAAGCTAAGTCTGAAC GACAAGATTCAACCCTAGCTTC	Rx26	M22146179	AGCTGCAAACAACATGGAC AGAGGCGAAACTTTTCTTAAC
Rx10	F3663458	TTGAAACAAGCTGGATATAGCC CAAAAATACGACTTTCCATGCC	Rx27	M24152935	AGCAATCCTACGTGACAGC TCTAGATTGAAAGAGGGGTG
Rx11	F37007195	TGATTTCTCATCAAGCAACAC TTAAGTTCTTTGGGTCCGAC	Rx28	M25938614	TTGAGGCATACCTCTCACC TGTGGGATGATTTGAAGACAC
Rx12	F44091788	TCTACTCCAATTGCAACGATG GGGCTTGTGTTGACTTTG	Rx29	M2696007	TCAAGAGTGCTTGGCACAG CTTGAATTTACGCAGCATCAG
Rx13	F5567539	CGCATTTTATGGTGCTGC GCTCCATCCAATCTTCCAGG	Rx30	M27531625	AGGGAGAAGTGAATGGTGGC ATCTACGAAACCACCTGTAACC
Rx14	F5883398	TGCAGTTTCCAAAGTCGTTAG CAATCCTGCTCATTAAGCTCG	Rx31	M31773413	GCTCCTATGTTCTGGCACAC CCGTTATGATAGATCGGATTGC
Rx15	F5936656	GTCACAATGGAGCAATTTTACC ACAGCTTGATTCAATCTGGTC	Rx32	M32128495	AGATGGCCTAGTGATATACGC GTTTAGTACGACTACCTTCCAC
Rx16	F6118470	ACATCTGGGAATCAACTTCAC CCCAAACGAGAAAAGAAGCTG	Rx33	M35343149	TGTGACCTAAGAGTGTCTC TGACATGCTCACATGAAATAGC
Rx17	F6386942	CATCTTTTGTCTTTGGGATTCG CCATGAAACAGGTCCATGAG	Rx34	M44553680	TGTTTCTAAAGCCTAGTTGCC TTTGCTCTCGGTTGACAC
			Rx35	M6969018	ACTTGCCAGACTCTAAGAAAAG TCCACTATTCCATAAACATGC