

Table 1 Polymorphism information of SSR primers amplification

Primer	Observed number of alleles (N_a)	Effect number of alleles (N_e)	Observed heterozygosities (H_o)	Expected heterozygosities (H_e)	Polymorphism information content (PIC)	Shannon's information index (I)
SSR02	10	7.562 5	0.727 3	0.909 1	0.853 8	2.145 8
SSR03	11	6.722 2	0.818 2	0.891 8	0.836 2	2.137 5
SSR05	8	5.500 0	0.636 4	0.857 1	0.793 5	1.843 9
SSR08	9	4.938 8	0.545 5	0.835 5	0.775 6	1.870 0
SSR09	10	6.540 5	0.727 3	0.887 4	0.830 9	2.074 5
SSR12	10	4.745 1	1.000 0	0.826 8	0.773 1	1.940 1
SSR13	9	5.902 4	0.545 5	0.870 1	0.810 6	1.960 8
SSR15	9	5.902 4	0.727 3	0.870 1	0.810 6	1.960 8
SSR16	10	7.117 6	0.727 3	0.900 4	0.844 3	2.110 6
SSR18	8	5.902 4	0.272 7	0.870 1	0.809 8	1.909 2
SSR21	10	6.540 5	0.363 6	0.887 4	0.830 9	2.074 5
SSR23	8	4.653 8	0.727 3	0.822 5	0.758 7	1.767 8
SSR24	8	5.761 9	0.272 7	0.865 8	0.805 1	1.897 7
SSR26	10	6.722 2	0.727 3	0.891 8	0.835 8	2.089 9
SSR30	8	5.260 9	0.363 6	0.848 5	0.787 0	1.854 1
Mean	9.2	5.984 9	0.612 1	0.869 0	0.810 4	1.975 8
Total	138	89.773 2	-	-	-	-