

Table S1. Information on SSR markers used in the present study with their GenBank accession numbers, linkage groups, primer sequences and bibliographic references¹.

Locus	GenBank accession number	Linkage group	Primer sequence	
			Forward (5'-3')	Reverse (5'-3')
NSX175	scaffold_64	1	GAGCTTACTAATAGGATACGCTG	ATTGTAAGGCGAATTGGATGG
TSGR264	scaffold_2	2	GGATTTAGAGAAAGTAGAGCAACCA	TTTGAGAACCCAGCACTATAACC
NSX02	scaffold_3	3	ACATCACATCGCCAATACCC	ACTCGTGGGACTCATGACTC
NSX165	scaffold_57	3	GTTGTTCTTGTATGAAGGGGTACAG	TTTGTTTGTCTGTTGGCTTGTC
GSR3138	BUZY63372.g1	5	GCCTTTGCTAACTGGAAAGTC	CATAAGCTTGCGTGTTGTGT
GSR6101	BUZY128935.g1	7	AAAAACAGAGCAGAAGGAAC	TGCAATCCCATAATAGCAAAG

¹ Shimizu et al. [21]