

**Supplementary Table S2.** Table information of six pairs of EST-SSR primers and polymorphsim results

Accession	SSR sequence	Sequence of primer(5'-3')	Observed number of alleles ( <i>Na</i> )	Effective number of alleles ( <i>Ne</i> )	Nei's gene diversity ( <i>H</i> )	Shannon's Information index ( <i>I</i> )	polymorphic Loci
ZB23	(CCAC) <sub>5</sub> (CA) <sub>8</sub>	F:GGAGCTTCAGACTTGGTGCA R:ACCAGGATCAGGTTCCACGA	2	1.1559	0.1086	0.1866	13
ZB26	(AGG) <sub>5</sub>	F:ATTCACATCCACTGCGAGCA R:GCCTTGCTGCTCTCCTTACA	2	1.3371	0.2164	0.3443	7
ZB28	(CTT) <sub>8</sub>	F:TGATAGCCACAGCCACAGTG R:ACAGTGTGTGCCATCCTCAG	2	1.1969	0.1498	0.2612	5
ZB53	(GCAACA) <sub>7</sub>	F:GCACGGCAACAGTTCGTATC R:CCTGCCCCATTGAATTCCT	2	1.5337	0.3205	0.4884	6
ZA26	(GGA) <sub>5</sub> (GGT) <sub>5</sub>	F:GCAGAACGATGAAGGGTGGA R:TGAACCACAAACCACCACCA	2	1.2738	0.1608	0.2543	13
ZA51	(TCCAAG) <sub>5</sub>	F:GGTGCCATGTCACAGGATCA R:GTTTGTGACTGCTGCAAGGG	2	1.4961	0.2855	0.4287	7