

Supplementary Table S3 Polymorphism levels of the 8 SSRs over the entire sample set

Locus	N	Na	Ne	He
MA4	427	19	3.692	0.729
MA7	426	5	1.180	0.152
MA14	423	13	3.834	0.739
MA12	426	6	2.295	0.564
MA19	426	5	1.696	0.410
MA51	418	11	6.327	0.842
MA21	422	15	4.530	0.779
MA13	429	3	1.141	0.124
Mean	424.63 (1.22)	9.63 (2.03)	3.09 (0.65)	0.54 (0.10)

N = sample size; Na = No. of Different Alleles; Ne = No. of Effective Alleles;
He = Expected heterozygosity. The standard error is indicated in parentheses.