

Table S3. Quality of microsatellite genotyping of museum samples at different loci

#	Locus	Mean QI value (M ± SE)	QI = 1		0.75 ≤ QI < 1.00		Probability (p) of correct genotyping
			Number of samples, n	Ratio of samples, %	Number of samples, n	Ratio of samples, %	
1	TGLA227	0.988 ± 0.008	22	91.7	2	8.3	***
2	BM2113	1.000 ± 0.000	24	100.0	0	0.0	***
3	TGLA53	0.961 ± 0.018	18	81.8	4	18.2	***
4	ETH10	0.992 ± 0.008	23	95.8	1	4.2	***
5	SPS115	0.967 ± 0.016	20	83.3	4	16.7	***
6	TGLA122	0.960 ± 0.019	20	83.3	4	16.7	***
7	INRA23	0.994 ± 0.006	22	95.7	1	4.3	***
8	TGLA126	0.983 ± 0.012	21	91.3	2	8.7	***
9	BM1818	0.965 ± 0.019	17	85.0	3	15.0	***
10	ETH225	0.988 ± 0.008	22	91.7	2	8.3	***
11	BM1824	1.000 ± 0.000	24	100.0	0	0.0	***

QI – quality indices (the calculation of quality indices is described in section 2.2 of the main text, \*\*\* p < 0.001)