Table A2: Characteristics of microsatellite loci applied in the study. All were run in the same panel; i.e. run in the same capillary.

Locus information			Amplification details			Genetic diversity		
Reference[a]	*Locus*[b]	*Label*[c]	*MP*[d]	*Primer(µM)*[e]	*PCR*[f]	*Na*[g]	*Allele-range*[h]	*Uhe*[i]
Genebank: AF151370	BFRO13	FAM	MP1	0.09	58	5	235-251	0.66
(Junge *et al*., 2009)	213	FAM	single	0.38	60	11	283-327	0.78
(Junge *et al*., 2009)	414	FAM	MP2	*0.20*	60	7	389-413	0.71
(Junge *et al*., 2009)	309	FAM	MP4	0.57	59	2	447-451	0.47
(Sušnik *et al*., 2000)	BFRO10	VIC	MP1	0.08	58	5	96-128	0.33
(Sušnik *et al*., 1999b)	BFRO15	VIC	MP1	0.04	58	2	144-154	0.50
(Sušnik *et al*., 1999b)	BFRO18	VIC	MP1	0.04	58	5	181-195	0.59
(Junge *et al*., 2009)	207	VIC	MP1	0.10	58	2	216-224	0.47
(Sušnik *et al*., 1999a)	BFRO9	VIC	MP1	0.05	58	2	243-247	0.16
(Junge *et al*., 2009)	438	VIC	single	0.34	60	10	261-297	0.79
(Sušnik *et al*., 2000)	BFRO11	NED	MP3	0.30	59	2	86-102	0.43
(Junge *et al*., 2009)	313	NED	MP2	0.18	60	6	180-200	0.75
(Olsen *et al*., 1998)	Ogo2	NED	MP1	0.07	58	4	233-245	0.63
(Junge *et al*., 2009)	433b	NED	MP3	0.18	59	8	291-319	0.69
(Junge *et al*., 2009)	445	NED	MP4	0.13	59	13	374-422	0.85
(Junge *et al*., 2009)	419a	PET	single	0.24	58	2	106-110	0.29
(Junge *et al*., 2009)	415	PET	MP3	0.33	59	10	189-225	0.76
(Junge *et al*., 2009)	211	PET	MP2	0.25	60	5	232-240	0.08
(Junge *et al*., 2009)	214	PET	MP1	0.14	58	3	292-313	0.53

[a]Publication of the locus or Genebank accession number if unpublished.
[b]Common name of the locus
[c]Label, Fluorescent dye used
[d]MP, Multiplex group or single locus PCR amplification
[e]Primer concentration in the PCR-mix
[f]Annealing temperature profile of the PCR program used

[g]Na, Number of alleles found in the locus
[h]*A*-range, Range of the length (bp) of alleles
[i]H_E, Unbiased expected heterozygosity in the locus

Marker information and methodological details are described in detail in Junge *et al*. (2009) and Koskinen and Primmer (2001).

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