

Table S1

PCR primers.

STR	Primer forward seq 5'-3'	Primer reverse seq 5'-3'	Concentration (nM)	Multiplex
STR 1	ATTCCCTGGCTGTGAATG	GCACATGGGAGAAACAAACA	15	2
STR 2	ACTGGTCACCAAGTCCATCC	GCTTGGCAACATCCCTGTAT	5	2
STR 3	GCTATAAACCTGCCCCTGCT	CAAGCAGCTGAGATTACTGGTG	305	3
STR A	AACGATAAGATGGCCCTAAGC	CCCTGTCAATCAAGAGCACA	75	1
STR B	TTTCTTGGCCAGTTGCATATC	CCAGCTTGGGTGAGAAAGAG	200	1
STR 4	AGATATCTGTGCGGCCATGT	TTAGCATTCAAGTTCGTCCA	305	2
STR 5	GGGAATCACTTGAACCCACA	TCCCTGAGACTTCCACTTTGA	10	2
STR 6	CTCCAGCCTAGGTGGCATAG	GCATCTTGGTTTGGCAATTC	5	3
STR 8	AGGGTTGTGTGAGGGATTTCT	TTGGCTATTCGATGCCACTT	3.5	3
STR 9	GGGCAACAGAGCAAGATGTAG	TGGAGATAATCCAAATCAGAAGAA	15	3
STR 10	CTGAGAGGCTGAAGCAGGAA	CCTTGAACTCCAGGGCTCTA	3.5	3