

Table S1. Primer sequences.

	Primer name	Primer sequence (5' - 3')	Positions of the targeted fragment	Fragment size (base pair)	Reference		
Hypervariable Region I							
	L15996	CTCCACCATTAGCACCCAAAGC	15977 - 16161	185	[1]		
	H16142	ATGTACTACAGGTGGTCAAG			[2]		
	L16117	TACATTACTGCCAGCCACCAT	16097 - 16237	141	[3]		
	H16218	TGTGTGATAGTTGAGGGTTG			[4]		
	L16209	CCCCATGCTTACAAGCAAGT	16190 - 16322	133	[4]		
	H16303	TGGCTTTATGTACTATGTAC			[4]		
	L16287	CACTAGGATACCAACAAACC	16268 - 16429	162	[4]		
	H16410	GCGGGATATTGATTCACGG			[4]		
Hypervariable Region II							
	L00205	ACAGGCGAACATACTACTAAAGTG	00181 - 00261	105	This study		
	H00261	GTTATGATGTCTGTGTGGAAAGTGG			This study		
	L00270	TGAATGTCTGCACAGCCACTTTCCA	00246 - 00350	105	This study		
	H00327	TTGGCAGAGATGTGTTTAAGTGCT			[5]		
Coding Region							
	Haplogroup	Targeted SN	Primer name	Primer sequence (5' - 3')	Positions of the targeted fragment	Fragment size (base pair)	Reference
	C1f	8577	L8562	TTCGCTTCATTCATTGCCCCC	08542 - 08624	110	This study
			H8624	AGTCGGTTGTTGATGAGATATTTGGAGG			This study
	C	9545	L9524	CTGAGCCTTTTACCACTCCAGCCTA	09500 - 09559	83	This study
			H9559	GATTTAGCGGGGTGATGCCTGTTG			This study
	C1f	11605	L11595	ACAAGCTCCATCTGCCTACGACA	11573 - 11631	80	This study
			H11631	ACTACGAGGGCTATGTGGCTGA			This study
	C1f	12217	L12197	TGACAACAGAGGCTTACGACCCC	12175 - 12197	68	This study
			H12223	TGGGGGCATGAGTTAGCAGT			This study

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