Table S3. Results of SNP typing in the mtDNA coding region using the GenoCore22 SNaPshot assay.

			I																					
		13928_R9	3594_L3'4	$10550_{\rm L}\rm K$	11467_U	4248_A	8994_W	13263_C	13368_T	11719_preHV, R0	8280de1B	4580_V	K_178	10238_N1	7028_H	$10034_{-}I$	5178_D	14766_HV	12705_R	10873_N	12612_J	2758_L2'6	10400_M	hg
rCRS		С	G	t	A	T	G	t	G	С	A	С	g	T	g	Т	g	G	g	a	a	G	g	Н
Sample	Extract #																							
aUz																								
UZOO-43	1	C	G	T	G	T	G	Т	G	T	Α	C	G	T	Α	T	G	Α	G	Α	Α	G	G	U
	2	С	G	Т	G	T	G	T	G	Т	Α	C	G	T	Α	T	G	Α	G	Α	Α	G	G	U
UZOO-46	1	C	G	T	G	T	G	T	G	T	Α	C	G	T	Α	T	G	Α	G	Α	Α	G	G	U
	2	С	G	Τ	G	T	G	Τ	G	Τ	Α	C	G	T	g/a	T	G	Α	G		Α	G	G	U
UZOO-16	1	C	G	Τ	G	T	G	Τ	G	T	Α	C	G	T	Α	T	G	Α	G	Α	Α	G	G	U
	2	C	G	Τ	G	T	G	Τ	G	Т	Α	C	G	T	g/a	T	G	Α	G	Α	Α	G	G	U
UZOO-40	1	С	G	T	G	T	G	T	G	T	Α	С	G	T	Α	T	G	Α	G	Α	Α	G	G	U
	2	С	G	T	G	T	G	T	G	T	Α	C	G	T	Α	T	G	Α	G	Α	Α	G	G	U
UZOO-70	1	С	G	Τ	G	T	G	T	G	Τ	Α	C	G	T	Α	Т	G	A	G	Α	Α	G	G	U
	2	С	G	Τ	G	T	g/a	Τ	G	Τ	Α	C	G	T	Α	Т	G	A	G	Α	Α	G	G	U
UZOO-77	1	С	G	Τ	Α	T	G	T	G	C	Α	C	G	T	G	Т	G	G	G	Α	Α	G	G	H
UZOO-7	1	С		T	Α	T	G	C	G	T	Α	С	G	T	Α	T	G			G	Α	G	Α	С
	2	С	G	T	A	T	g/a	C	G	Т	A	С	G	T	Α	T	G		A	G	A	G	Α	C
UZOO-8	1	С	G	Т	A	T	G	С	G	Т	A	С	G	T	Α	T	G	Α	A	G	A	G	A	С
	2	С	G	Т		T	g/a	С	G	Т	A		G	T	g/a	T	G			G	A	G		С
UZOO-74	1	С	G	Т	A	T	G	С	G	Т	A	С	G	Т	Α	Т	G		A	G	A	G	A	С
	2	С	G	Т	A	T	G	C	G	Т	A	С	G	T	Α	T	G		A	G	A	G	A	С

SNPs typed on the L-strand are reported in capital letters in the reference rCRS profile, whereas SNPs typed on the H-strand are reported in small letters. Missing data signifies allelic dropout or fluorescence signal below the background threshold (100 relative fluorescent units, rfu). 'g/a' indicates the presence of a mixed signal for the position interrogated. A mixed signal was repeatedly obtained at position 8994 (haplogroup W) with the detection of an additional G base. However, the rest of the profile never could support phylogenetically the presence of the G base at this particular position. For each individual, profiles were obtained from two independent extracts, except for individual BOO72-9 for which a second samples was not available and for UZOO-77, BOO57-1, BOO72-10, BOO72-4, BOO72-7, BOO72-15, and BOO72-1, for which the second individual was extracted in an independent laboratory. rCRS, revised Cambridge Reference Sequence; hg, haplogroup.

										0														
										_preHV, R0														
			4							еHУ				_				>				9		
		\\ \text{2}	L3'	_K	U_	V_	M .	C	T	_pr	lelB	>	X.	Z	⊞.	Ī	Q.	Η_	R	$\mathbf{z}_{_{\mathbf{I}}}$	\mathbf{f}^{-1}	L2'	Z _	
		3928_R9	594_L3'4	10550	11467_U	4248_	8994_W	13263	13368	11719	8280delB	4580	6371	10238	7028_H	10034	5178_D	4766_HV	12705_	10873	12612	2758 <u>_</u> L2'6	10400	
aPo			ë	=	=	4	×	Ξ	<u> </u>	1	œ	4	9	=	<u> </u>	1	N.	Ť	1	1	7	7	=	hg
Po4	1	С	G	Т	G	T	g/a		G	T	Α		G	T	A	T	G		G		Α	G		U
	2	С	G	Т	G	T	g/a	Т	G	T	Α		G	T	Α	T	G				Α	G	G	U
Po2	1	С	G	T	G	T	G	T	G	T	Α	С	G	T	Α	T	G	Α	G	Α	Α	G	G	U
	2	С	G	T	G	T	g/a	T	G	T	Α		G	T	Α	T	G		G	Α	Α	G	G	U
aBOO																								
BOO49-3	1	С	G	T	G	T	G	T	G	T	Α	С	G	T	Α	T	G	A	G	Α	Α	G	G	U
	2	C	G	T	G	T	G	T	G	T	A	C	G	T	A	T	G	A	G	Α	Α	G	G	U
BOO57-1	1	C	G	T	G	T	G	T	G	T	A	C	G	T	A	T	G	A	G	A	A	G	G	U
BOO49-1	1	C	G	T	G	T	G	T	G	T	A	C	G	T	A	T	G	A	G	A	A	G	G	U
DO072 11	2	C	G	T	G	T	G	T	G	T	A	C	G	T	A	T	G	A	G	A	A	G	G	U
BOO72-11	1 2	C	G	T	G	Т	G	T	G	Т	A	C	G	T	A	T	G	A	G	A	A	G	G G	U
BOO72-9	1	C	G	T	G	T	G	T	G	T	A	C	G	T	A	T	G	A	G	A	A	G	G	U
BOO72-9 BOO72-10	1	C	G	T	G	T	G	T	G	Т	A	C	G	T	A	T	G	A	G	A	A	G	G	U
BOO72-10 BOO72-14	1	С	G	T	G	T	G	T	G	T	A	C	G	T	A	T	G	A	G	A	A	G	G	U
BOO72-14	2	C	G	T	G	Т	G	T	G	T	A	С	G	Т	A	T	G	A	G	A	A	G	G	U
BOO72-8	1	C	G	T	G	Т	G	T	G	Т	A	C	G	T	A	T	G	A	G	A	A	G	G	U
B00720	2	C	G	T	G	T	G	T	G	T	A	C	G	T	A	T	G	A	G	A	A	G	G	U
BOO72-4	1	C	G	T	A	T	G	T	A	T	A	C	G	Т	A	T	G	A	G	A	A	G	G	T
BOO49-2	1	C	G	Т	Α	Т	G	С	G	Т	Α		G	Т	Α	Т	G			G	Α	G		С
	2	С	G	Т	Α	Т	G	С	G	T	Α	С	G	Т	Α	T	G			G		G		C
BOO49-4	1	С	G	Т	Α	T	G	C	G	T	Α	С	G	T	Α	T	G	Α	Α	G	Α	G	Α	С
	2	С	G	Т	Α	T	G	C	G	T	Α		G	T	Α	T	G			G		G	Α	C
BOO57-3	1	С	G	T	Α	T	G	C	G	T	Α	С	G	T	Α	T	G		Α	G	Α	G	A	C
	2	С	G	T	Α	T	G	C	G	T	Α	С	G	T	Α	T	G	Α	Α	G	Α	G	Α	C
BOO72-2	1	C	G	T	Α	T	G	C	G	T	Α	С	G	T	Α	T	G	Α	Α	G	Α	G	Α	C
	2	С	G	T	Α	T	G	C	G	T	Α	С	G	T	A	T	G	Α	Α	G	Α	G	Α	C
BOO72-7	1	C	G	T	A	T	G	C	G	T	A	~	G	_	A	T	G			G	A	G	A	C
BOO72-12	1	C	G	T	A	T	G	C	G	T	A	С	G	T	A	T	G		A	G	A	G	A	C
DO072.5	2	C	G	T	A	T	G	C	G	T	A		G	T	A	T	G		A		A	G	A	C
BOO72-5	1 2	C	G	T	A	T	G	C	G	T	A	C	G	T	A	T	G		A	G	A	G	A	C
BOO72-6	1	C	G	T	A	T	G	C	G	T	A	C	G	T	A	T	G	A	A	G G	A	G G	A	C
BUU/2-0	2	C	G	T	A	T	G	C	G	T	A	C	G	T	A	T	G	Α	A	G	A	G	A	C
BOO49-6	1	C	G	T	A	T	G	T	G	T	A	C	G	Т	A	T	T	A	A	G	A	G	A	D
50079-0	2	\vdash	G	T	A	T	G	T	G	T	A	C	G	T	A	T	T	A	11	G	A	G	11	D
BOO72-13	1	С	G	T	A	T	G	T	G	T	A	C	G	T	A	T	T	A	A	G	A	G	A	D
23372 13	2	C	G	T	A	Т	G	T	G	T	A	C	G	T	A	T	T	A	A	G	A	G	A	D
BOO72-15	1	C	G	T	A	T	G	T	G	T	A	C	G	T	A	T	T	A	A	G	A	G	A	D
BOO49-5	1	٦	G	T	A	T	G	T	G	T	A	C	G	T	A	T	G		A	G	A	G	A	M
11111	2	С	G	T	A	T	G	T	G	T	A	_	G	T	A	T	G	A	A	G	A	G	A	M
BOO72-3	1	C	G	T	Α	T	G	T	G	T	A	С	G	T	A	Т	G			G	A	G	Α	M
	2	С	G	Т	Α	T	G	Т	G	T	Α	С	G	T	Α	T	G	A	Α	G	Α	G	Α	M
BOO72-1	1	С	G	T	Α	T	G	T	G	T	Α		G	T	Α	T	G			G	Α	G	Α	M

SNPs typed on the L-strand are reported in capital letters in the reference rCRS profile, whereas SNPs typed on the H-strand are reported in small letters. Missing data signifies allelic dropout or fluorescence signal below the background threshold (100 relative fluorescent units, rfu). 'g/a' indicates the presence of a mixed signal for the position interrogated. A mixed signal was repeatedly obtained at position 8994 (haplogroup W) with the detection of an additional G base. However, the rest of the profile never could support phylogenetically the presence of the G base at this particular position. For each individual, profiles were obtained from two independent extracts, except for individual BOO72-9 for which a second samples was not available and for UZOO-77, BOO57-1, BOO72-10, BOO72-4, BOO72-7, BOO72-15, and BOO72-1, for which the second individual was extracted in an independent laboratory. rCRS, revised Cambridge Reference Sequence; hg, haplogroup.