

Table 2. Autosomal STR profiles and Amelogenin gene typings of the 10 individuals studied

Sample	PCRs, no.	Extractions, no.	Bone /Tooth samples, no.	<i>AMEL</i>	D3S1358	D8S1179	D5S818	vWA	D21S11	D13S317	FGA	D7S820	D18S51
SZA23 1	6	2	2	X/Y	14	13/14	11	14/16	26/29.2	11/13	22/23	7/8	-
SZA23 2	8	2	2	X	15/17	12	11	19	-	-	-	-	-
SZA23 3	6	2	2	X/Y	15/17	13/14	12/13	14/17	28/29	-	-	-	-
ELT 2	2	2	2	X/Y	15/16	12/17	12	14/(17)	30/31	8/9	24	-	-
DR2	8	3	2	X/Y	15/17	13/15	11	18/19	30/31	-	-	-	-
DEB 1	6	2	1	X	-	-	-	14	-	-	-	-	-
DEB 3	7	3	2	X	18	11/13	12	-	-	-	-	-	-
DEB 4	6	3	2	X/Y	18/19	13/14	11/13	16/17	-	9	-	-	-
KRE 1	9	4	2	X	16/17	11/15	12	16	28/29	-	-	-	-
KRE 2	7	3	2	X/Y	15/17	9/14	11/12	16	28	-	-	-	-

Consensus alleles were established if genotypes were reproduced in at least three amplifications. *AMEL*, amelogenin gene. Dashes indicate cases with fewer than three successful reproducible typings.