

FOR THE RECORD

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Autosomal STR Frequencies in Afghanistan Population

POPULATION: Afghanian ($n = 130$)

KEYWORDS: forensic science, DNA typing, population genetics, STR frequencies, Afghanistan

Buccal cells were collected by buccal brushes (Sterile Omni Swab, Whatman International Ltd.) from healthy, random Afghanian individuals. DNA was extracted by Chelex method (1). DNA samples were quantified using AmpF ℓ STR[®] Quantifiler (Applied Biosystems, Foster City, CA) by Real time 7000 (Applied Biosystems, Foster City, CA), one nanogram of the samples obtained from the previous phase was amplified by GeneAmp[®] PCR System 9600 (Applied Biosystems, Foster City, CA). Simultaneous amplifications of 16 STRs loci (multiplexed PCR) were performed by using AmpF ℓ STR[®] Identifiler[™] (Applied Biosystems, Foster City, CA) according to the user manual recommendations (2).

The sixteen loci analyzed in this study are D3S1358, vWA, FGA, D8S1179, D21S11, D18S51, D5S818, D13S317, D7S820, TH01, TPOX, CSF1PO, D19S433, D2S1338, D16S539 and the gender determination marker Amelogenin.

STRs typing was performed on 16-capillary ABI Prism[®] 3100 Genetic Analyzer (Applied Biosystems, Foster City, CA) capillary electrophoresis system according to the user manual and data were analyzed by GeneMapper v. 3.2 Software (Applied Biosystems, Foster City, CA).

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STRs frequencies were calculated using the statistical software Poppene v. 1.31 (3). Each locus was tested for Hardy-Weimberg equilibrium by the Chi-square test (χ^2).

The complete data are available to any interested researcher upon request.

References

1. Walsh PS, Metzger DA, Higuchi R. Chelex 100 as a medium for simple extraction of DNA for PCR-based typing from forensic material. *Biotechniques* 1991 Apr;10(4):506-13. [\[PubMed\]](#)
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TABLE 1—*STR allele frequencies in Afghanistan population (n = 130).*

Allele	D8S1179	D21S11	D7S820	CSF1PO	D3S1358	THO1	D13S317	D16S539
6	0.2692
7	0.0077	...	0.1769	0.0077	...
8	0.0308	...	0.1538	0.1462	0.1615	0.0615
9	0.0846	0.0308	...	0.1692	0.0846	0.1846
9.3	0.2308
10	0.0769	...	0.2231	0.2385	...	0.0077	0.1077	0.0923
11	0.0692	...	0.2846	0.2462	0.2385	0.3077
12	0.1308	...	0.2000	0.4308	0.3000	0.2231
12.2
13	0.2615	...	0.0385	0.0462	0.0846	0.1154
13.2
14	0.1923	...	0.0077	...	0.0769	...	0.0154	0.0154
14.2
15	0.1538	...	0.0077	...	0.3077
15.2
16	0.0846	0.2769
16.2
17	0.2308
18	0.1077
19
20
20.2
21
21.2
22
22.2
23
23.2
24
24.2
25
26
27	...	0.0231
28	...	0.1385
28.2
29	...	0.2231
29.2
30	...	0.2154
30.2	...	0.0231
31	...	0.0385
31.2	...	0.0923
32	...	0.0077
32.2	...	0.1308
33
33.2	...	0.1077
Ho	0.800	0.846	0.7846	0.6615	0.6462	0.8154	0.7385	0.7385
He	0.841	0.851	0.8030	0.6992	0.7637	0.7990	0.8070	0.8018
P	0.649	0.571	0.998	0.294	0.161	0.416	0.985	0.253

Ho: Observed Heterozygosity; He: Expected Heterozygosity; P: Probability value of Chi-square test for Hardy-Weimberg equilibrium.

TABLE 1—Continued.

Allele	D2S1338	D19S433	vWA	TPOX	D18S51	D5S818	FGA
6
7	0.0154	...
8	0.5308
9	0.0615	...	0.0846	...
9.3
10	0.1077	...	0.0615	...
11	...	0.0077	...	0.2154	0.0156	0.2385	...
12	...	0.0615	...	0.0846	0.0781	0.4769	...
12.2
13	...	0.2769	0.1562	0.1231	...
13.2	...	0.0231
14	...	0.2538	0.0923	...	0.2812
14.2	...	0.0846
15	...	0.1000	0.0692	...	0.1250
15.2	...	0.1077
16	0.0154	0.0615	0.2846	...	0.1250
16.2	...	0.0231
17	0.0923	...	0.3000	...	0.0781
18	0.1692	...	0.1615	...	0.0547	...	0.0231
19	0.1923	...	0.0692	...	0.0469	...	0.0769
20	0.0846	...	0.0154	...	0.0234	...	0.1000
20.2	0.0154
21	0.0231	...	0.0077	0.1154
21.2
22	0.0462	0.0156	...	0.0923
22.2
23	0.1154	0.2385
23.2
24	0.1000	0.2000
24.2
25	0.1615	0.1077
26	0.0308
27
28
28.2
29
29.2
30
30.2
31
31.2
32
32.2
33
33.2
Ho	0.8923	0.8000	0.8615	0.5846	0.7812	0.5385	0.7538
He	0.8731	0.8278	0.7906	0.6544	0.8535	0.6947	0.8587
P	0.907	0.415	0.977	0.269	0.348	0.170	0.164

Ho: Observed Heterozygosity; He: Expected Heterozygosity; P: Probability value of Chi-square test for Hardy-Weimberg equilibrium.