

FOR THE RECORD

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Allele Frequencies for Fifteen STR Loci in Guinea-Bissau Population

POPULATION: Guinea-Bissau population, ethnical groups: Balanta, Beafada, Cassanga, Fula, Mancanha, Mandinga, Manjaco, Nalú, Papel, $n = 92$

KEYWORDS: forensic science, DNA typing, population genetics, Occidental Africa, Guinea-Bissau population, D8S1179, D21S11, D7S820, CSF1P0, D3S1358, TH01, D13S317, D16S539, D2S1338, D19S433, vWA, TPOX, D18S51, D5S818, FGA

The ethnical variability is probably the most important characteristic of Guinea-Bissau population, in which 99% of the individuals constitute African ethnical groups, and European and mulatto are less than 1% (1).

Bloodstains were obtained from 92 unrelated and healthy known African individuals from Guinea-Bissau. The DNA was extracted using the Chelex® method (2). PCR amplification of D8S1179, D21S11, D7S820, CSF1P0, D3S1358, TH01, D13S317, D16S539, D2S1338, D19S433, vWA, TPOX, D18S51, D5S818 and FGA loci was performed using the AmpFℓSTR® Identifiler™ PCR Amplification Kit (Applied Biosystems) following the manufacturer's protocol (3). The amplified fragments were separated and detected using the ABI Prism® 310 Genetic Analyzer (PE Applied Biosystems) (4). Typing was made with GeneScan® Analysis Software (PE Applied Biosystems) and by comparison with allelic ladders (5). The data were analyzed using the program of statistical analysis Genepop (6). With this program it was confirmed the existence of Hardy-Weinberg equilibrium. Statistical parameters showed high discriminating power of these fifteen loci.

The complete data are available to any interested person upon request by contacting the corresponding author at joanapvp@clix.pt.

References

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TABLE 1—Allele frequencies for fifteen STR loci of Guinea-Bissau population ($n = 92$).

Allele	D8S1179	D21S11	D7S820	CSF1PO	D3S1358	TH01	D13S317	D16S539	D2S1338	D19S433	vWA	TPOX	D18S51	D5S818	FGA
6					0.1087						0.1304				
7				0.0761		0.3098					0.0272		0.0109		
8			0.2228	0.0598		0.3315	0.0217	0.0380			0.2011		0.0924		
9			0.1413	0.0326	0.0054	0.1522	0.0217	0.2283			0.2174	0.0054	0.0380		
9.3					0.0761										
10	0.0054		0.2989	0.2065		0.0217	0.0163	0.1087		0.0054		0.0870		0.0217	
10.2													0.0054		
11	0.0435		0.2174	0.1794	0.0054		0.3152	0.2554		0.0380		0.3152	0.0217	0.2011	
11.2											0.0054				
12	0.0815		0.1033	0.3044			0.4130	0.2283		0.1359	0.0054	0.0163	0.0652	0.3967	
12.2										0.0380					
13	0.2554		0.0163	0.1250	0.0380		0.1848	0.1250		0.2663	0.0217	0.0054	0.0707	0.2283	
13.2										0.0815			0.0054		
14	0.3315			0.0163	0.1250		0.0272	0.0109		0.2065	0.0870		0.0489	0.0109	
14.2										0.0652			0.0054		
15	0.2337				0.3043			0.0054	0.0054	0.0326	0.1630				
15.2										0.0870					
16	0.0435				0.3044				0.0598	0.0109	0.2663		0.1467		
16.2										0.0217					
17	0.0054				0.1630				0.0815		0.2391		0.1630		
17.2										0.0054					
18					0.0544				0.0543		0.125		0.1467		0.0054
18.2														0.0109	
19									0.1087		0.0598		0.0815		0.0870
20									0.0978		0.0272		0.0435		0.0380
21									0.2065		0.0054		0.0109		0.0598
22									0.1033				0.0054		0.1576
23									0.0652				0.0109		0.2228
24									0.1033						0.1902
25									0.0870						0.0978
26	0.0054								0.0217						0.0489
27	0.0163														0.0544
28	0.1576														0.0054
29	0.1685														0.0109
30	0.2446														0.0054
30.2	0.0380														
31	0.1141														
31.2	0.0272														0.0054
32	0.0109														
32.2	0.1359														
33	0.0054														
33.2	0.0489														
34.2	0.0054														
35	0.0163														
36	0.0054														
<i>p</i>	0.9062	0.0620	0.6079	0.6644	0.5677	0.4926	0.4839	0.1226	0.5965	0.4584	0.2978	0.6441	0.3793	0.3936	0.2931
H	0.7597	0.8501	0.7828	0.8062	0.7681	0.7529	0.6939	0.8015	0.8891	0.8449	0.8173	0.7873	0.8827	0.7394	0.8615
PD	0.9039	0.9606	0.9189	0.9366	0.9105	0.8999	0.8528	0.9313	0.9778	0.9592	0.9426	0.9236	0.9749	0.8933	0.9662
PE	0.5400	0.7028	0.5731	0.6230	0.5554	0.5312	0.4422	0.6071	0.7773	0.6976	0.6412	0.5867	0.7639	0.5174	0.7196

NOTE: *p*: Hardy-Weinberg equilibrium exact test; H: heterozygosity.