

FOR THE RECORD

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Genetic Polymorphism of Two Novel STR Loci AY639920 and AY639923 in a Chinese Population

POPULATION: Chinese.

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Blood samples were collected from unrelated individuals of the Chinese Han ethnic group in Chengdu of China. DNA was extracted using the Chelex method (1). PCR amplification conditions can be accessed at <http://www.legalmed.org/dna/AY639920.htm> and <http://www.fayi.cn/dna/AY639920.htm>. The volume of PCR reaction for each locus was 37.5 μ L. The PCR products were analyzed by vertical nondenaturing polyacrylamide gel electrophoresis with a continuous buffer system and visualized by silver staining (2,3). Data were analyzed using POWERSTATS program (4). The genotype distribution was analyzed for Hardy–Weinberg equilibrium according to Hou’s method (5) and no deviation from Hardy–Weinberg equilibrium was observed (Tables 1 and 2).

The complete data can be accessed at <http://www.legalmed.org/dna/AY639920.htm> and <http://www.fayi.cn/dna/AY639920.htm>.

TABLE 1—Allele frequencies of two STR loci in Chinese population.

AY639920 (N = 105)		AY639923 (N = 101)	
Allele	Frequency	Allele	Frequency
11	27.6%	14	1.5%
14	1.4%	15	7.4%
15	5.7%	16	30.2%
16	2.9%	17	39.6%
17	48.6%	18	11.9%
18	8.1%	19	8.9%
19	5.2%	20	0.5%
20	0.5%		
Total	1.000	Total	1.000
HWE*	$p > 0.05$		$p > 0.05$

*Test for Hardy–Weinberg equilibrium.

TABLE 2—Population genetics and forensic data of two STR loci.

Locus	PIC	DP	P_m	CE	H_o	H_e
AY639920	0.630	0.845	0.155	0.291	0.600	0.677
AY639923	0.680	0.883	0.117	0.497	0.743	0.731

PIC, polymorphism information content; DP, power of discrimination; P_m , probability of match; H_o , observed heterozygosity; H_e , expected heterozygosity.

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