

FOR THE RECORD

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Allele Frequencies for Two STR Loci D21S1436, D21S2052 in Chinese Population

POPULATION: Chinese

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Blood samples were collected from unrelated individuals of Chinese Han ethnic group in Chengdu of China. DNA was extracted using Chelex method (1). PCR amplification conditions can be accessed at <http://www.legalmed.org/dna/d21s1436.htm>. The volume of PCR reaction for each locus was 37.5 μ L. The PCR products were analyzed by horizontal non-denaturing polyacrylamide gel electrophoresis with discontinuous 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.

The complete data can be accessed at <http://www.legalmed.org/dna/d21s1436.htm>.

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

D21S1436 (N = 103)		D21S2052 (N = 102)	
Allele	Frequency	Allele	Frequency
5	0.010	17	0.010
6	0.447	18	0.029
7	0.073	19	0.093
8	0.024	20	0.132
9	0.151	21	0.260
10	0.126	22	0.324
11	0.092	23	0.098
12	0.058	24	0.034
13	0.019	25	0.020
Total	1.000	Total	1.000
HWE*	$p > 0.05$		$p > 0.05$

* Test for Hardy-Weinberg equilibrium.

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

Locus	PIC	DP	Pm	CE	H _o	H _e
D21S1436	0.72	0.911	0.089	0.574	0.786	0.736
D2S2052	0.76	0.919	0.081	0.643	0.824	0.782

* PIC (polymorphism information content), DP (power of discrimination), Pm (probability of match), H_o (observed heterozygosity), H_e (expected heterozygosity)

Reference

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