

FOR THE RECORD

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Population Data of Two STR Loci in Chinese Population

POPULATION: Chinese

KEYWORDS: forensic science, Han in Sichuan, China, DNA typing, short tandem repeats, polymerase chain reaction, D8S1128, D7S3058

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/D8S1128.htm>. The volume of PCR reaction for each locus was 20 μ L. The PCR products were analyzed by horizontal non-denaturing polyacrylamide gel electrophoresis with discontinuous buffer system and visualized by silver staining (2). Data of population genetics and forensic science were analyzed using POWERSTATS program (3). The genotype distribution was analyzed for Hardy-Weinberg equilibrium

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

| Allele | Frequency | |
|--------|-------------------|-------------------|
| | D8S1128 (N = 100) | D7S3058 (N = 114) |
| 7 | 0.391 | |
| 8 | 0.260 | 0.225 |
| 9 | 0.083 | 0.189 |
| 10 | 0.109 | 0.081 |
| 11 | 0.078 | 0.095 |
| 12 | 0.057 | 0.113 |
| 13 | 0.016 | 0.126 |
| 14 | 0.005 | 0.099 |
| 15 | | 0.059 |
| 16 | | 0.014 |
| Total | 1.000 | 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 | PD | Pm | PE | H _o | H _e |
|---------|------|-------|-------|-------|----------------|----------------|
| D8S1128 | 0.72 | 0.882 | 0.118 | 0.683 | 0.848 | 0.851 |
| D7S3058 | 0.84 | 0.956 | 0.044 | 0.797 | 0.901 | 0.963 |

PIC: polymorphism information content; PD: power of discrimination; Pm: probability of match; PE: power of exclusion; H_o: observed heterozygosity; H_e: expected heterozygosity.

according to Hou's method (4). No deviation from Hardy-Weinberg equilibrium was observed. The complete data can be accessed at <http://www.legalmed.org/dna/D8S1128.htm>.

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