## FOR THE RECORD

*Weibo Liang*,<sup>1</sup> *Ph.D.; Meili Lv*,<sup>1</sup> *Ph.D.; Yongchun Xu*,<sup>1</sup> *M.D.; Miao Liao*,<sup>1</sup>; *Bin Zhou*,<sup>1</sup> *M.D.; Yi Jia*,<sup>1</sup> *Ph.D.; and Zhang Lin*,<sup>1,2</sup> *Ph.D.* 

## Allele Frequency Distribution of STR Loci D5S814 in Four Populations\*

**POPULATIONS:** 100 unrelated Chinese volunteer donors, 100 unrelated Thai volunteer donors, 100 unrelated Mongol volunteer donors, 100 unrelated Germany volunteer donors.

KEYWORDS: forensic science, D5S814, short tandem repeat, Chinese, Thai, Mongol, Germany

 TABLE 1—Allele frequency distributions of D5S814 in four populations.

Populations Chinese Menggu Thai Germany Allele (n = 100)(n = 100)(n = 100)(n = 100)23 24 25 0.250 . . . 0.005 0.020 0.155 0.160 0.160 0.185 26 0.350 0.290 0.240 0.280 27 28 0.285 0.395 0.385 0.330 0.175 0.145 0.155 0.170 29 0.025 0.040 0.010 0.015 DP 0.882 0.858 0.900 0.893 Het 0.780 0.730 0.660 0.730 PE 0.562 0.476 0.369 0.476 0.690 0.700 0.710 PIC 0.660 HWE test\* 0.995 0.954 0.953 0.982

\* Probability values.

DP: power of discrimination.

Het: heterozygosity.

PE: power of exclusion.

PIC: polymorphism information content.

Blood Specimens were obtained from 100 unrelated Chinese volunteer donors, 100 unrelated Thai volunteer donors, 100 unrelated Mongol volunteer donors, 100 unrelated Germany volunteer donors respectively. DNAs were extracted from blood specimens using Chelex- $100^{(1)}$ . Genotyping were carried out by PCR in a PE9600 cycler. The components of a 20 µL reaction mixture were as follows:

Genotypes	Populations			
	Chinese $(n = 100)$	$\begin{array}{c} \text{Menggu} \\ (n = 100) \end{array}$	Thai $(n = 100)$	Germany $(n = 100)$
24–25				1
25-25	3	2	4	3
23-26			2	
24-26				1
25-26	13	14	6	8
26-26	10	5	8	12
23-27			2	
24-27	1			2
25-27	10	12	10	13
26-27	19	24	16	17
27-27	7	16	18	9
23-28			1	
25-28	3	1	7	8
26-28	15	10	6	6
27-28	12	10	8	14
28-28	2	4	4	3
25-29		1		1
26-29	3		2	
27-29	1	1	5	2
28-29	1		1	

TABLE 2—Genotype distributions of D5S814 in four populations.

template DNA 20 ng, primer 0.2  $\mu$ mol/L each, dNTPs 200  $\mu$ mol/L each, KCl 50  $\mu$ mol/L, Tris-HCl(pH 8.3)10 mmol/L, MgCl<sub>2</sub> 1.5 mmol/L, Taq polymerase 1U. Primer sequences: D5S814: 5'-tg-gacttttccagcacagat-3',5'-ctctacaaagaagttaaatcgagc-3'. PCR conditions: start at 94°C for 4 min, followed by 36 cycles consist of 35 s at 94°C, 40 s at 58°C, 50 s at 72°C followed by a 10 min extention at 72°C. The amplified products were electrophoresed in 6% polyacrylamide gel by using 100 bp ladder and allelic markers as size markers, followed by silver staining. The amplified products were analyzed by The Promega Software, POWERSTATS. The

100

100

100

Total

100

<sup>&</sup>lt;sup>1</sup> College of Forensic Medicine, Sichuan University, Chengdu 610041, P. R. China.

 $<sup>^2\,{\</sup>rm Key}$  Lab of Biotherapy of Human Diseases, Ministry of Education, P. R. China.

<sup>\*</sup> The research was supported by grants from the Chinese Natural Sciences Foundation (No. 30171033) and State Ministry of Education (No. 01143) as well Sichuan Youth Foundation (2001–2).

## 2 JOURNAL OF FORENSIC SCIENCES

complete dataset is available to any interested researcher upon request from the corresponding author.

## Reference

1. Singer-Sam J, Tanguary RL, Riggs AD. Use of Chelex to improve the PCR signal from a small number of cells. Amplification 1989:3;11.

Additional information and reprint requests: Professor Zhang Lin, Ph.D. College of forensic medicine Sichuan University Chengdu, 610041, Sichuan, P. R. China Tel: 86-28-85460532 Fax: 86-28-85405541 E-mail: kjc@scu.edu.cn