# A POPULATION GENETIC STUDY OF THREE STRs LOCI IN A CHINESE HAN POPULATION IN CHENGDU, PR CHINA 

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We have surveyed a Chinese Han population in Chengdu, PR China with three STR loci of DHFRP ${ }_{2}$, FIBRA, and ACTBP ${ }_{2}$ located on the human chromosome $6,4 \mathrm{q} 28$ and 6 respectively by using both single PCR, PAGE and silver stain as well as triplexing out. The distribution of alleles (table), genotypes observed, H, DP, PIC, and EPP were calculated. A new allele A 9.2 of DHFRP2 with size of 177 bp was firstly found. The sequence study demonstrated that the 84th nucleotide $C$ on the basic sequence was replaced by T and A, A on both 140th, 141th positions were deleted. The number of cases studied observed genotypes, H, DP, PIC, and EPP of DHFRP 2 , FIBRA, and ACTBP 2 were as follows: 156, 19, $0.61,0.87,0.68,0.46 ; 136,44,0.90,0.96,0.86,0.86 ; 147,86,0.97,0.99,0.95$, and 0.93 . The distribution of their genotypes were in good agreement with HWE. The family study demonstrated that these three loci were in conformity with the Mendel's law. Triplexing of these three loci was achieved. The matching probability of these three loci was $7.6 \times 10^{-5}$. Tissues taken from animals of nine different kind species were used for the species specificity test. The results showed that no PCR products were found only in ACTBP2 locus. Nine species of animals tested were as follows: rabbit, pig, cow, cat, snake, chicken, rat frog, and fish. The results of 5 cases of paternity testing and personal identification of biological evidences were correct.

Table Allele distributions of DHFRP $_{2}$, FIBRA, and ACTBP 2 loci

| DHFRP | $\mathrm{DHFRP}_{2}$ | FIBRA | FIBRA | ACTBP | ACTBP |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Allele | Frequencies | Allele | Frequencies | Allele | Frequenc |
| A6 | 0.0665 | A9 | 0.0037 | A16 | 0.0034 |
| A7 | 0.4359 | A11 | 0.0184 | A17 | 0.0068 |
| A8 | 0.266 | A12 | 0.0772 | A18 | 0.0068 |
| A9 | 0.0449 | A13 | 0.0588 | A19 | 0.0102 |
| A9.2 | 0.0256 | A14 | 0.1471 | A20 | 0.0408 |
| A10 | 0.141 | A14.2 | 0.0037 | A21 | 0.0782 |
|  |  | A15 | 0.1691 | A22 | 0.0952 |
|  |  | A16 | 0.2059 | A23 | 0.0714 |
|  |  | A16.2 | 0.0184 | A24 | 0.051 |
|  |  | A17 | 0.1654 | A25 | 0.0816 |
|  |  | A17.2 | 0.011 | A26 | 0.0578 |
|  |  | A18 | 0.0846 | A27 | 0.0408 |
|  |  | A18.2 | 0.0037 | A28 | 0.051 |
|  |  | A19 | 0.011 | A29 | 0.0544 |
|  |  | A20 | 0.0147 | A30 | 0.0578 |
|  |  | A21 | 0.0074 | A31 | 0.1054 |
|  |  |  |  | A32 | 0.0578 |
|  |  |  |  | A34 | 0.0816 |
|  |  |  |  | A36 | 0.0442 |
|  |  |  |  | A38 | 0.0032 |

