

Multiplex Fluorescent PCR Analysis in Japanese Population

Masaki Hashiyada, Masayuki Nata, Jun Kanetake, Noboru Adachi, Guijin ji and Masato Funayama
Department of Forensic Medicine, Tohoku University School of Medicine, Sendai, Japan



Introduction

Short tandem repeat (STR) analysis is a useful tool in forensic sciences to get information of individual identification. Recently, several STR loci can be amplified in one tube using multiplex PCR STR kit which are commercially available.

Material and Methods

We investigated 10 STR loci in Japanese population living in Sendai by two multiplex PCR kits, *GenePrint*[®] PowerPlex[™] Fluorescent STR System (Promega, USA) and AmpF/STR[™] Profiler[™] (Perkin-Elmer, USA). Genomic DNA was extracted from EDTA whole blood and oral swab using SDS-Proteinase K or Chelex treatment followed by the phenol/chloroform extraction. PCR was performed in accordance with manufacture's protocols. Electrophoresis was carried out on an ABI 377 sequencer and the alleles were determined by GeneScan[®] 2.0.2 software (Perkin-Elmer, USA).

Results and Discussion

The statistical data of analyzed 10 STRs are shown in Table 1. In all loci, statistical parameters indicated relatively high rate, and no significant deviation from Hardy-Weinberg Equilibrium was detected. We apply this STR system for the paternity test and forensic casework, e. g., personal identification in rape cases. This system is very effective in the paternity test. However, in some rape cases, PCR products failed to amplify several loci in high molecular range because of DNA degradation.

Locus	Number of Individuals	Alleles Observed	Het.	PD	PIC
<i>D7S820</i>	321	8	0.771	0.914	0.738
<i>D13S317</i>	322	9	0.818	0.941	0.792
<i>D5S818</i>	322	9	0.786	0.922	0.754
<i>CSF1PO</i>	246	7	0.734	0.888	0.693
<i>TPOX</i>	248	7	0.691	0.851	0.637
<i>TH01</i>	249	8	0.725	0.875	0.676
vWA	249	8	0.800	0.930	0.770
<i>D16S539</i>	221	7	0.777	0.916	0.742
<i>D3S1358</i>	107	7	0.705	0.860	0.651
<i>FGA</i>	107	10	0.831	0.930	0.770

Het.: heterozygosity,
PD: power of discrimination,
PIC: polymorphic information content