新Y-STR DYS605在山西汉族人群中的多态性分布 Study of Polymorphism at New Y-STR DYS605 in a Chinese Han Population of Shanxi

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摘要

为了应用更多的Y染色体特异性STR基因座以用于法医学和人类遗传学研究, 本文用PCR结合PAGE技术检测128例山西 汉族无关男性DYS605等位基因分布状况。结果显示:山西地区汉族男性DYS605基因座观察到22,21,20,19,18 共5个等位基因,基因频率分别为0.0156; 0.1797; 0.4531; 0.2891; 0.0625。等位基因20和19之间的电泳距离在 ▶ <u>本刊中 包含"法医学"的</u> 非变性胶上非常接近,要有足够的电泳距离才能区分。测序表明该基因座包括3个串联重复区,其中一个为可变重 复区。20例女性DNA未发现扩增产物。Abstract: We study the polymorphism at DYS605 ,a new tetranucleotide Y-STR locus, in a Chinese Han population of Shanxi to meet the need of more genetic markers in forensic practice and genetic analysis. DNA were extracted from 128 unrelated male venous blood, and amplified using GDB primers. PCR products were detected using non-denaturing polyacrylamide gel electrophoresis and silver staining. Five alleles named 22, 21, 20, 19, 18 were observed with frequency of 0.0156; 0.1797; 0.4531; 0.2891; 0.0625. PCR products were not found in female DNA. Using a long enough gel for a long electrophoresis time is strongly encouraged because the rung between allele 20 and allele19 is smaller than expected. Allele sequences show that the repetitive units of DYS605 were composed with the variant units and non-variant units.

法医学 Y染色体短串联重复序列 遗传多态性 山西 汉族人 DYS605 Key words forensic science Y-STR Shanxi Chinese Han population DYS605

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Abstract

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