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**An anonymous human single copy genomic clone, D5S6 (M4) on chromosome 5 identifies a three allele RFLP**


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**SOURCE AND DESCRIPTION OF CLONE:** A 7.6kb Bam HI fragment cloned in the phage  $\lambda$ L47.1 isolated from a human genomic library.

**POLYMORPHISM:** Bam HI (G/GATCC) (Amersham) identifies a three allele polymorphism at 11.0kb (A1), 9.6kb (A2) or 7.6kb (A3) (Figure).

**FREQUENCY:**

	11.0kb allele (A1)	0.33
Studied 29 Caucasians;	9.6kb allele (A2)	0.52
21 female, 8 male.	7.6kb allele (A3)	0.15

**NOT POLYMORPHIC FOR:** Eco RI, Hind III, Msp I, Taq I, Pst I with a panel of 9 unrelated Caucasians.

**CHROMOSOMAL LOCALIZATION:** The probe was assigned to chromosome 5 using a panel of somatic cell hybrids (Fox & Retief, in press) and localized to 5q22-q31 by means of in-situ hybridization.

**MENDELIAN INHERITANCE:** Co-dominant segregation shown in 2 informative families (9 individuals).

**PROBE AVAILABILITY:** Available for collaborators.

**OTHER COMMENTS:** No problems on RFLP analysis under normal conditions. Low background at stringency of 0.5 x SSC.

**REFERENCE:** Fox MF and Retief AE, "Aspects of human chromosome segregation in somatic cell hybrids: establishment of a hybrid mapping panel" SAJ Science (in press).

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