An anonymous X-chromosomal clone identifying a frequent RFLP at Xp21-22 (HGM8 provisional no. DXS207)

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SOURCE AND DESCRIPTION OF CLONE: pPA4B, a 3.5 kb HindIII fragment subcloned in pAT153 from a human genomic Charon 21A library enriched for X-chromosomal sequences (Kunkel et al. Nucl. Acids Res. <u>10</u> 1557 (1982).

POLYMORPHISM: Xba1 (T/CTAGA) identifies a simple two-allele polymorphism with either one band at 12.0 kb (A1) or two bands at 9.5 kb and 2.5 kb (A2).

FREQUENCY:Studied in 54 European Caucasians, 31 females, 23 males: 12.0 kb allele (A1) 0.68 9.5 kb allele (A2) 0.32

NOT POLYMORPHIC FOR: BstEII, EcoRI, EcoRV, HindIII, MspI, TaqI, using a panel of 5 unrelated Caucasians (9 X-chromosomes).

CHROMOSOMAL LOCALISATION: Between Xp21 and Xp22 using a panel of translocationhybrids (generous gift from Dr K.H. Grzeschick, Münster, FRG, and Dr H.-H.Ropers, Nijmegen, Holland).

MENDELIAN INHERITANCE: Codominant X-linked segregation shown in 5 informative families, 47 individuals.

PROBE AVAILABILITY: Available for collaboration.

OTHER COMMENTS: The 2.5 kb band (not shown in figure) is very faint, probably because only a short sequence is homologous to the clone.

REFERENCE: Ahrens et al.(1985) Cytogenet, Cell Genet, 40 567.

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