Isolation and mapping of a polymorphic DNA sequence pYNH24 on chromosome 2 (D2S44)

Y.Nakamura, S.Gillilan, P.O'Connell, M.Leppert, G.M.Lathrop, J.-M.Lalouel and R.White

The Howard Hughes Medical Institute, University of Utah Medical School, Salt Lake City, UT 84132, USA

SOURCE/DESCRIPTION: A 2.0 kb MspI fragment from cosmid YNH24 isolated by HBV-2 oligonucleotide (GGAGTTGGGGGAGGAG) (1) was subcloned into the AccI site of pUC18.

<u>POLYMORPHISM</u>: MspI identifies a >30 allele VNTR polymorphism with bands between 1.0 - 5.0 kb. TaqI, BglII, PvuII, PstI and BamHI also detect the same polymorphism.

FREQUENCY: With MspI, 97% Heterozygosity was obserbed in 108 unrelated Caucasians.

NOT POLYMORPHIC FOR: RsaI

<u>CHROMOSOMAL LOCALIZATION</u>: pYNH24 has been assigned to chromosome 2 by multipoint linkage analysis(2) with loci (APOB, D2S6) known to span this region(3,4).

MENDELIAN INHERITANCE: Co-dominant segregation of the MspI RFLP was observed in 44 three generation families.

PROBE AVAILABILITY: Contact Y.N.

<u>OTHER COMMENTS</u>: RFLPs were observed under normal hybridization and wash stringencies.

REFERENCES:

- 1. Y. Nakamura et al. Science 235:1616-1621 (1987)
- 2. G.M. Lathrop et al. Am. J. Hum. Genet. 37:482-498 (1985)
- 3. S.Povey, N.E. Morton and S.L. Sherman, Cytogenet. Cell Genet. 40:67-106 (1985)
- 4. G.M. Lathrop et al., abstract submitted to Human Gene Mapping Workshop 9. Cytogenet. Cell Genet., in press