A c-DNA probe for the oncogene c-MEL (pC7-1) recognises a polymorphism with NcoI

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SOURCE AND DESCRIPTION. A 2kb EcoRI fragment of the human $\underline{\text{MEL}}$ oncogene c-DNA inserted into pEMBL8.

POLYMORPHISM. NooI (C/CATGG) identifies two alleles with bands at 4.0kb and 3.8kb. Constant bands of 8kb, 4.3kb, 2.45kb, 2.1kb and 1.95kb are seen.

FREQUENCIES. Studied in 32 unrelated individuals.

4.0kb allele 73.4% 3.8kb allele 26.6%

NOT POLYMORPHIC FOR. AccI, AluI, AvaI, AvaII, BamHI, BclI, BglI, BglII, ClaI, EcoRI, HaeIII, HincII, HindIII, HinfI, HpaI, HpaII, KpnI, MspI, PstI, PvuI, PvuII, RsaI, SalI, Sau3A, SmaI, TaqI, TthIII, XbaI, XhoI.

CHROMOSOMAL LOCATION. Localized to chromosome 19 using a panel of human-hamster and human-mouse hybrids.

MENDELIAN INHERITANCE. Co-dominant segregation demonstrated in 4 families.

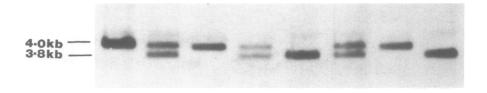
PROBE AVAILABILITY. Available for collaborators, contact D.Hughes or R.A.Padua.

OTHER COMMENTS. Wash at 65 C, 0.1 X SSC.

REFERENCES.

Padua RA, Barras N, and Currie GA (1984) Nature $\underline{311}$ 671-673. Spurr NK, Hughes D, Goodfellow PN, Brook JD and Padua RA (1986) Somat. Cell Mol. Genet. $\underline{12}$ 637-640.

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