

Apa I polymorphism of a human immunoglobulin VH3 subclass locusMichael A. Walter*, Peter S. Linsley⁺ and Diane W. Cox**Research Institute, Hospital for Sick Children and Departments of Paediatrics and Medical Genetics, University of Toronto, Toronto, Ontario M5G 1X8, Canada and ⁺Oncogene, Seattle, WA, USA**SOURCE/DESCRIPTION:**

a 2.2 kb Eco RI fragment derived from λ VH26 (1), flanking a human immunoglobulin VH3 subclass gene

POLYMORPHISM:

a polymorphic Apa I (GGCC/C) fragment of 11.8 kb
(a 9.4 kb fragment is constant)

FREQUENCY: studied in DNA from 45 random Canadian caucasians:

frequency of 11.8 kb fragment: 0.54 ± 0.07
frequency of the absence of the fragment: 0.46 ± 0.07

NOT POLYMORPHIC FOR:

digests of DNA from 10 unrelated individuals with Xba I or Bam HI,
did not reveal polymorphisms

CHROMOSOMAL LOCALISATION:

14q32.3 to 14qter (2)

MENDELIAN INHERITANCE:

co-dominant segregation shown in 4 two generation families and
4 three generation families

PROBE AVAILABILITY:

write to D. W. Cox

OTHER COMMENTS:

wash stringency of 0.1% SDS, 0.1 x SSC; at 65°C

REFERENCE:

- 1) Matthyssen G., Rabbitts T.H. (1980) P.N.A.S. USA. 77, 6561-6565.
- 2) Cox D.W., Markovic V.D., Teshima I.E. (1982) Nature. 297, 428-430.

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