

First nucleotide sequence of a human immunoglobulin variable λ gene belonging to subgroup II

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The first nucleotide sequence of a human variable lambda gene belonging to subgroup II and designated as V λ 2.1 is reported. This gene was isolated from the genomic λ 275 library (clone λ 275V λ 2). That library was made with the DNA of a member of the family 275, characterized by two exceptional Gm haplotypes (1). The peptide leader of 19 codons contains a 118 base pair (bp) intron. The heptamer-nonamer recombination signal sequences are underlined and the cysteins involved in the intrachain disulfide bond are circled. The additional cystein in the 5' region of the gene belongs to the third hypervariable region (Chuchana, P. and Lefranc, M.-P., unpublished data). The translation of the sequence shows that this gene belongs to subgroup II (2). The subclone pV λ 2EK0.3, containing a 310 bp fragment Eco 47 III-Kpn I (restriction sites overlined in the sequence below) represents a specific V λ II probe which only detects the V λ genes belonging to subgroup II in high stringency conditions.

TGCTGATTTCATGAGCCCGCCTCTCTAGCAAGGGATAGACAGGCTGGGCAGGGCCATGCTGGGTCAAAAGAGGCAAGCCTCTGGGACGTCCTCCACC	100
ATG GCC TGG GCT CTG CTG CTC ACT CTC CTC ACT CAG GAC ACA G GTGACGCCCTCCAGGGAAAGGGTCTGGGACCTCTGG	184
M A W A L L L L T L L T Q D T	15
GCTGATCTTGGTCTCTGCTCTCAGGCTCACCGGGCCAGCACTGACTCACTGGCATGTGTTCTCCCTCTTCCAGGG TCC TGG GCC CAG	278
-1 +1 G S W A Q	20
TCT GCC CTG ACT CAG CCT GCC TCC GTG TCT GGG TCT CCT GGA CAG TCG ATC ACC ATC TCC TGC ACT GGA ACC AGC	353
S A L T Q P A S V S G S P G Q S I T I S C T G T S	45
AGT GAT GTT GGG AGT TAT AAC CTT GTC TCC TGG TAC CAA CAG CAC CCA GGC AAA GCC CCC AAA CTC ATG ATT TAT	428
S D V G S Y N L V S W Y Q Q H P G K A P K L M I Y	70
GAG GGC AGT AAG CGG CCC TCA GGG GTT TCT AAT CGC TTC TCT GGC TCC AAG TCT GGC AAC ACG GGC TCC CTG ACA	503
E G S K R P S G V S N R F S G S K S G N T A S L T	95
ATC TCT GGG CTC CAG GCT GAG GAC GAG GCT GAT TAT TAC TGC TGC TCA TAT GCA GGT AGT AGC ACT TTA C CACAGT	579
I S G L Q A E D E A D Y Y C C S Y A G S S T L	118
GGTCCAAGTTCATGGGGACTGAGACCAAAACCTGCCAGGGCTTCAGACTTCCCTTGCTCTGAAGATGCTTCTCACCOGGTGCAAGGGCTTGCTG	679
CAGCGAGGCCCTTGAGAATTG	699

References: (1) Lefranc, G. et al (1982) J. Immunogenet. 9, 1-9.

(2) Kabat, E.A. et al. (1987) ed. Nat. Inst. Heath, Bethesda, pp 63-77.