

Genomic sequence of IGLV1S2, a human immunoglobulin variable lambda gene belonging to subgroup I

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Only three genomic sequences of human immunoglobulin variable lambda genes have been published (reviewed in ref. 1). We report the second genomic sequence of a Vλ gene belonging to subgroup I. A clone (cosmid 8.3) was isolated by screening a cosmid library made from the human tumor cell line COLO 320 HSR DNA (2) with the VλII probe pVλ2EK0.3 (3). The cosmid 8.3 clone contains a Vλ gene which was identified by cross hybridization to that probe at low stringency. This IGLV gene was shown by sequencing, to belong to subgroup I according to reference 1, and was designated as Vλ1.2 or IGLV1S2 since it represents the second genomic sequence of a VλI subgroup gene (4) ('IGLV1S' stands for 'human immunoglobulin lambda variable gene belonging to subgroup I', 'IGLV1S2' stands for Vλ1.2 following the Human Gene Mapping recommendations (HGM9)). The translation of the sequence shows a peptide leader of 19 codons

with a 109 base pair intron. The heptamer-nonamer recombination signal sequences are underlined. The IGLV1S2 gene encodes a Vλ region with a 6 amino acid CDR1 region as this is observed in the VλI LOC, MEM, NEWM and HA proteins (see ref. 1). The analysis of the deduced protein sequence of IGLV1S2 shows that the protein NEWM (5) is most probably encoded by the IGLV1S2 gene.

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GCAGCACTACGACAATCTCCAGC ATG GCC TGG TCT CCT CTC CTC ACT CTC CTC GCT CAC TGC ACA G GTGACTGGATAACAGGTCCAGGGAGGGGCCCTGG 103
M A W S P L L L T L L A H C T

AAGCCTATGGATTCTTGCTTCTCTGTTGCTCTAGAACCGAATAATGATGCTGCTCTCCACTTCCAGG 210
G S W A Q S V V T Q P

CCC TCA GTG TCT GGG GCC CCA GGG CAG AGG GTC ACC ATC TOC TGC ACT GGG AGC AGC TCC AAC ATC GGG GCA GGT TAT GAT GTA CAC TGG 300
P S V S G A P G Q R V T I S C T G S S S N I G A G Y D V H W
Kpn I
TAC CAG CAG CTT CCA GGA ACA GCC CCC AAA CTC CTC ATC TAT GGT AAC AGC AAT CGG CCC TCA GGG GTC CCT GAC CGA TTC TCT GGC TCC 390
Y Q Q L P G T A P K L L I Y G N S N R P S G V P D R F S G S

AAG TCT GCC ACC TCA GCC TCC CTG GCC ATC ACT GGG CTC CAG GCT GAG GAT GAG GCT GAT TAT TAC TGC CAG TCC TAT GAC AGC AGC CTG 480
K S G T S A S L A I T G L Q A E D E A D Y Y C Q S Y D S S L
S G

AGT GGT TCCACAGTGCTCCAGGCCGGGTGGAACTGAGACAAGAACCATCTCTGCCAGGGTGACCC 555

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