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Rat inv (193-256) V L T K C Q E E V S H I P D V H P G A F R P K C D E N G N
Mou inv (192-255) V L T K C Q E E V S H I P A V Y P G A F R P K C D E N G N
Hum inv (193-256) V L T K C Q E E V S H I P A V H P G S F R P K C D E N G N
Hum Tg (97-160) Q L Q K Q Q I L L S G Y I N S T D T S Y L P Q C Q D S G D

Y M P L Q C H G S T G Y C W C V F P N G T E V P H T K S R G R H N C S
Y L P L Q C H G S T G Y C W C V F P N G T E V P H T K S R G R H N C S
Y L P L Q C Y G S I G Y C W C V F P N G T E V P N T R S R G H H N C S
Y A P V Q C D V Q H V Q C W C V D A E G M E V Y G T R Q L G R P K R C
    
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Fig. 3: Alignment of the amino acid sequence encoded by the additional segment in rat invariant chain cDNA with the sequence encoded by exon 6b of mouse and human invariant chain genes and the sequence preceding and including the second type I repetitive unit of human thyroglobulin. Amino acids identical in 2 or more sequences are boxed. The amino acid homology between rat and human invariant chain proteins in this region is 88% compared with 75% for the rest of the protein.

presence of eight extra nucleotides at the extreme 5' end of our clone. Also, we have noted four differences between the protein sequences at positions 16, 17, 136 and 147 where Henkes *et al.* have incorrectly translated the nucleotide sequence.

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