

Nucleotide sequence of a rabbit genomic DNA encoding mature endothelin-3

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Endothelin, a potent vasoconstrictor peptide, was isolated and its cDNAs from human and porcine libraries were cloned (1, 2). Analysis of human genomic DNA suggested the existence of an ET-family (ET-1, ET-2, ET-3) (3). Using ET-3 specific antibody, we have recently revealed the ET-3 expression in the brain, intestine and placenta of rat (4). We have also isolated a human cDNA clone for ET-3 (5). To examine the species specificity of ET-3 genes, we have examined a rabbit ET-3 gene and present here its partial DNA sequence.

The rabbit genomic library (EMBL-3) was constructed from liver chromosomal DNA and screened (6) with a 45-nucleotide synthetic DNA probe corresponding to mature ET-1. The predicted amino acid sequence of rabbit ET-3 (boxed in the figure) is identical to human and rat sequences (3,7), and the prepro-ET-3 region is highly conserved, suggesting its important role in animals. Compared with human ET-3 gene (unpublished data) or its cDNA (5,9), this sequence includes the entire exon

containing mature region and its 3'-exon (probably exon 1). The predicted splice sites and junction consensus sequences in the rabbit gene are indicated by arrows and underlining, respectively. The first methionine at position 317 is thought to be the translation initiation site. The length of the 3'-portion of rabbit prepro-ET-3 is the same as the human type except for the 24bp repeat sequence which exists in both the human genome and cDNA but not in the rabbit genome.

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TGTGTTGGCGCGACCCCGGGCGCCCTGGTCAAAGGCCCGCGCAGCTCCAGCCCCCTCCGGGGGGCGGGAGGCAGGGGG	90
GTGGTGGAGGCCAGAAAAGCCCGAGCCCACAGCCGGGAGTCCTCTGGCGGGGATGGCGACGGCGCGCTGAAAGTTGGTACCCGGCGCAC	180
CCAACTGCGCGCTGCAGCCCGAGGACCGAGCGAGCCAGGGAGCCGGCGGGCTCGAACCCCCACGGCGAGCCCGCGGGCGCTGTACCTGG	270
CCACCCAGCGGGGACCTGCGCCCGGGTGCTCCCGCGCTGATCCGGGTCATGGAGCCGGCTGTGGATCCTTTGGGGCTCACAGTG	360
M E P G L W I L L G L T V	13
ACCGCCGCCCG <u>AGGTAA</u> GGCCCGGGCGCGCGCTGTCGGCGCGAGCGCACACAAAAGGACCTGGCGGGGAGGTGGCGCGTCGC	450
T A A A	17
GGGGAGGGCCCGCACCCCTGGAGGGCGCTGGCGGGGCGACAGCTCAGCGCAGGGCGCTGCAGCTGTGGCTATGGGGCTG	540
GTCAGCTACTGGTCAGTGTCTCGAAGGCTCTGCAGACTGCAGAACTGCTAGCCAAGTTTCAGTGGCCGAGCAGAGTGCCTGCAA	630
GTTTCAGGGAGTTAGATGGCTTGAGGCTCTGCAGGGAGTGGCTGGAGGAAGCTGCAAGCGCTTCACACGGCATGCACGGGCTTGG	720
AACATTTCTGAAACTGTGTTAGGGCTGTGCTGGCTGGAGGCTTGGAGGCTTGGAGGATGTGAGACAATTGGAGATAACTTGC	810
AGGAGTTGCTGCCCGCCCGAACCCAGGTGGCTGCCGGGTGGCGCTGAGGGCTCTGCTCTGGCTCAGGAG <u>CCCTGGTCTTGG</u>	900
<u>CTCCCTGAGGATTCTGCGCTTGGCCAGCCCGACTGGGGGCTGGCAGGACCAGCGTGCCTGGGGGGGGGGGGGGGGGGGGGGGGGG</u>	990
G F V P C P Q T G G A G R T S V P R A P R V A G S E G	44
GAAGTGTGAAAGACTCTGGCCAGCCCTAGAAGGCAGACTGTGGCCCCAACGGCAGGGCAAGGGCCAGGCCCTGGAAAGCCCTGGCGGGGG	1080
D C E D S V A S P R R Q T V A P T A G K G P S P G S P G R G	74
CAGGGGGGGAGGGGAGCCGGGGCACCGCCGTGTGGCAGCTGCACCTGCTACCTACA <u>AAAGACAAAGAGTGGCTACTACTGCCCC</u>	1170
Q A A E G D P G H R R R C T C F T Y K D K E C V Y Y C H	104
CTGGACATCTGGATCAACACTCCGGAGTGAGTCAGGCCAGCCCCCTGCTCACTGTCCCCGTCCAGGGGGCTGATGCCACC	1260
L D I I W I N T P (*)	113
CCTCAGCCCTGCGGGTCTCCGCCAGCCGGTCCCGCTGCAGGTCATGTGAGCCGCTGCTGGCCCTGGCCCCGG	1350
GAAACCCAACTCTGCCGTGTGGCCGGAGAGGCCCTCCAGGAGCAGTGAGCTGGAGAGAGGATTAGACACCGAGACGCCAGGAGTT	1440
GAGGTGGAGGAAAGCCCGAAAGTGTGCTGCCGACCCAGGGCTGAGTGTGCTTGCCACCTGCTGACCTCCCAGGCTGCCAGGAGTT	1530
TAGAATT	1538