

Nucleotide sequence of the cynomolgus monkey apolipoprotein E cDNA

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Submitted December 8, 1988

EMBL accession no. X13887

The complete nucleotide sequence of the cynomolgus monkey apolipoprotein E and the deduced amino acid sequence have been determined. The nucleotide sequence is 1178 nucleotides in length and has a single open reading frame which encodes a protein of 317 amino acids. The cynomolgus apolipoprotein E nucleotide and the deduced amino acid sequence are quite similar to that reported for human (1,2) with approximately 93% homology.

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CTACTCAGCCCCAGCGGAGGTGAAGGACGTCTTCCCCAGGAGCCGACTGCCAAGGCAGCTGCCAATCACAGGCAGGA
AGATGAAGGTTCTGTGGGCTCGCTGGTCACATTCTGGCAGGATGCCAGGCCAAGGTGGAGCAACCGGTGGAGCCA
M K V L W A A L L V T F L A G C Q A K V E Q P V E P
GAGACGGAACCCGAGCTCGCCAGCAGGCTGAGGCCAGAGCGCCAGCCCTGGGAGCTGGCACTGGTCGCTTTGGGA
E T E P E L R Q Q A E G Q S G Q P W E L A L G R F W D
TTACCTCGCCTGGGTGAGACACTGTCTGAGCAGGTGCAAGGAGCTGCTCAGCCCCCAGGTACCCAGGAACGTGACGA
Y L R W V Q T L S E Q V Q E E L L S P Q V T Q E L T
CGCTGATGGACGAGACCATGAAGGAGTTGAAGGCCTACAAATCGGAACACTGGAGGAACAGCTGAGCCCGTGGCGAGGAG
T L M D E T M K E L K A Y K S E L E E Q L S P V A E E
ACCGGGGACGGCTGCTCAAGGAGCTGCAGGGCGCAGGCCCGCTGGGTGCGCACATGGAGGACGTGCGCAGCCGCT
T R A R L S K E L Q A A Q A R L G A D M E D V R S R L
GGTCAGTACCGCAGCGAGGTGCAAGGCCATGCTGGGCCAGAGTACCGAGGAGCTGCCGGCGCCTGCCCTCCCACCTGC
V Q Y R S E V Q A M I G Q S T E E L R A R L A S H L
GCAAGCTGCGCAAGCGGCTCTCCGCGATGCTGATGACCTGCAAGGCCCTGGCAGTGTATCAGGCCGGGCCGAG
R K L R K R L L R D A D D L Q K R L A V Y Q A G A R E
GGCGCCGAGCGGGGGTCAAGGCCATCCCGAGGCCCTGGGACCCCTGGTGAGCAGGCCCGTGGGGCCACTGT
G A E R G V S A I R E R L G P L V E Q G R V R A A T V
GGGCTCCCTGGCCAGCCAGCGCTTCAGGAGCGGGGCCAGGCCCTGGTGAGCGCTGCCACGGATGGAGGAGATGG
G S L A S Q P L Q E R A Q A L G E R L R A R M E E M
GCAGCCGGACCCGCGACCGCCTGGACGGAGGTGAAGGAGCAGGTGGCGAGGTGCGGCCAAGCTGGAGGAACAGGCCAG
G S R T R D R L D E V K E Q V A E V R A K L E E Q A Q
CAGATAAGCCTGCAGGCCAGGCCCTTCAGGCCCTGGACGGCCCTCAAGAGCTGGTCAGGCCCTGGTAAGATATGCAGGCCA
Q I S L Q A E A F Q A R L K S W F E P L V E D M Q R Q
GTGGGCTGGGCTGGTGGAGAAGGTGAGCGCTGCCCTGGGCGCCAGCACGCCCTGTGCCCATGACAATCACTGAACGC
W A G L V E K V Q A A V G A S T A P V P I D N H
CCAGGCCCTACAGCCATGCGACCGACTCCACCCATGCCCTCTCCGCCAGGCCCTGTCCCCG
CCCCAGCCGTCCAGGGGTGGGCCCTAGTTAATAAGATTGCCAAGTTCACCG

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