

Nucleotide sequence of cDNA encoding subunit Va from rat heart cytochrome *c* oxidaseMartin Droste, Eric Schon¹ and Bernhard Kadenbach*Fachbereich Chemie der Philipps-Universität, Hans-Meerwein-Straße, D-3550 Marburg, FRG and ¹Columbia University, College of Physicians and Surgeons, New York, NY 10032, USA

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Mammalian cytochrome *c* oxidase (COX) is composed of 3 catalytic, on mitochondrial DNA encoded, and of 10 nucleus-encoded subunits (1). The nucleus encoded subunits occur in tissue- and developmental-specific forms (2) and appear to modulate the functional activity of the enzyme (3). Recently the cDNA for human COX subunit Va was described (4). This gene probe was used for screening a rat heart cDNA library in λgt11 to isolate the corresponding gene of the rat. The cDNA sequence was determined after subcloning in phage M13mp8 by standard procedures. The deduced amino acid sequence of the mature protein is 100% identical to bovine heart subunit Va (5) and 96% homologous to human subunit Va (4). The presequence of rat subunit Va contains 37, that of human 41 amino acids. The identical amino acids with human subunit Va are underlined, the cleavage site of the presequence is indicated by an arrowhead.

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M L A A A A L R R C T A A A A A A A R
GTGGTCGCCGTATGCTCGCCGCTGCCCTCCGTGCTGTACCGCAGCCGCGGCCGCCGA 60

G L L H P V S A P S P A A A A V C S I R C
GGCCTCTGCACCCCGTCTCGGCTCCAGCCCCGCTGCCGCTGCTGTTCCATTGCTGC 120

Y S H G S H E T D E E F D A R W V T Y F
TATTCTCATGGTCACATGAGACAGATGAGGAGTTGATGCTCGCTGGTGACATACTTC 180

N K P D I D A W E L R K G M N T L V G Y
AACAAAGCCAGACATTGATGCCTGGAGTTGCGTAAGGGATGAATACTTGTGGCTAT 240

D L V P E P K I I D A A L R A C R R L N
GATCTGGTCCCTGAGCCAAAATCATTGATGCTGCTTGAGGGCATGTAGACGGTTAAAT 300

D F A S A V R I L E V V K D K A G P H K
GATTTGCTAGTGCCTGTCGATCTGGAGGTTGTTAAGGACAAGCAGGACCTCATAAG 360

E I Y P Y V I Q E L R P T L N E L G I S
GAAATCTACCCCTATGTCATCCAGGAACCTAGACCAACTTAAATGAATTGGGAATCTCC 420

T P E E L G L D K V #
ACTCCAGAGGAACCTGGGCCTTGACAAAGTGTAAACCTCCCTCGATGGCTTCCAAGGAC 480
TTAACCTGCTATTGCTACTTGATTGAAACAGTTGCTGGAAATGTTTATTGAAACAAATT 540
TTCCCTTGAGTATCAAACCATGTAACCTGGACTTTAAAGGGAAATGAGTTGACCCCG 600
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 644

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