Sequence of a cDNA specifying subunit VIIa of human cytochrome c oxidase

Gian Maria Fabrizi¹, Rosario Rizzuto¹, Hirofumi Nakase¹, Shuji Mita¹, Margaret I.Lomax³, Lawrence I.Grossman⁴ and Eric A.Schon^{1.2*}

Departments of ¹Neurology and ²Genetics and Development, Columbia University College of Physicians & Surgeons, 630 West 168th Street, New York, NY 10032, ³Department of Anatomy and Cell Biology, University of Michigan Medical School, Ann Arbor, MI 48109 and ⁴Department of Molecular Biology and Genetics and the Center for Molecular Biology, Wayne State University School of Medicine, Detroit, MI 48201, USA Submitted July 24, 1989 EMBL accession no. X15822

Cytochrome \underline{c} oxidase (COX; EC 1.9.3.1), the last component of the mitochondrial respiratory chain, catalyzes the transfer of electrons from reduced cytochrome \underline{c} to molecular oxygen. In mammals, the apoprotein is composed of three large catalytic subunits, encoded by the mitochondrial genome, and by ten smaller, nuclear-encoded subunits, which may play a regulatory role (1); subunits VIa, VIIa, and VIII have been shown to have heart- and liver-specific isoforms in cows and pigs (2,3).

Using a bovine liver COX VIIa cDNA (4) as a probe, we isolated a full-length cDNA (clone pCOX7.22; sequence below) from a human adult endothelial cell cDNA library (a gift of M. Chao and D. Littman), specifying the liver-specific isoform of human COX VIIa. The deduced amino sequence of the mature polypeptide is 83% identical to beef liver COX VIIa (3,4) but is only 57% identical to beef heart COX VIIa (5). Human COX VIIa contains a 23-aa cleavable presequence for importation into mitochondria (underlined).

[GAATTOGG]AGTAACAGOCAAG (13)

ATGCTGCGGAATCTGCTGGCTCTTCGTCAGATTGGGCAGAGGACGACGACGACGACTAAGCACTGCTTCC (73) $M \ L \ R \ N \ L \ L \ A \ L \ R \ Q \ I \ G \ Q \ R \ T \ I \ S \ T \ A \ S \ [-4]$	
OGCAGGCATTITIAAAAATAAAGTICOOGGAGAAGCAAAAACIGTICCAGGAGGATGATGAA (133) <u>R R H</u> F K N K V P E K Q K L F Q E D D E [17]	
ATTOCACIGIATCIAAAGGIGGGGIAGCIGAIGCOCICCIGIATAGAGCCACCAIGAIT (193) I P L Y L K G G V A D A L L Y R A T M I [37]	
CTTACAGTTGGIGGAACAGCATATGCCATATATGAGCTGGCTGGCTTCATTTCCCAAG (253) L T V G G T A Y A I Y E L A V A S F P K [57]	
AAGCAGGAGTGACTTCAGTCATCCCAGCAATCGCTTGGTTCAGTTTCATTCA	
TGGACCAGTAATCTGATAAATAACCGAGCTCTTCTTTGGGGATCAATATTTATT	I

*To whom correspondence should be addressed

Notes: (1) Kadenbach et al. (1987) Curr. Top. Bioenerg. 15, 113; (2) Kadenbach et al. (1983) TIBS 8, 398; (3) Yanamura et al. (1988) Biochemistry 27, 4909; (4) Seelan et al. (1989) Nucl. Acids Res., in press; (5) Meinecke and Buse (1986) Biol. Chem. Hoppe-Seyler 367, 67. Supported by grants from NIH (NS11766 [E.A.S.], RR05384 [L.I.G.], and CM37086 [M.I.L.]); the Muscular Dystrophy Association; the Aaron Diamond Foundation; the Center for Molecular Biology, Wayne State University; and Dr. and Mrs. Libero Danesi.