

Nucleotide sequence of a full-length cDNA for arylamine N-acetyltransferase from rabbit liver

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A full-length cDNA clone encoding rabbit liver arylamine N-acetyltransferase (NAT; EC 2.3.1.5) was isolated from a λ gt11 expression library by concomitant screening with an oligonucleotide deduced from partial amino acid sequences (1) and with antibodies raised against the purified protein. The nucleotide sequence of the clone, designated *rnat*, consists of 1102 nucleotides and contains a single open reading frame of 870 nucleotides encoding a 290 amino acid protein (Mr 33,655 Da) with 14 of 16 tryptic peptide sequences recently determined from purified rabbit liver NAT (1). The *rnat* cDNA shows 61% and 46% sequence homology at the nucleotide and deduced amino acid level, respectively, to those of NAT from chicken liver (2).

CTGATTGCAGGTCTGGAAACCTGAGTTGGCTGGGAAGATAAAATCCAACTCACCTGAATTAAAATGAAAGATCAACCCCTTTC	90
AACTTACTAAGGGTATCATGGATATTGAAGCATATTATCAAAGATTGGTTACAAAAACCCAAAGGAACTAACATTGGACTTGGAATCCTTA	180
M D I E A Y Y Q R I G Y K P R N K L D L E S L T	
CAGACATTTTCAGCACAGATCCGAACGTGCTCTATGAGAACCTTAGCATTCACTGTGGGAGTCATGGAGTTGGACTTAGAGGCCA	270
D I F Q H Q I R T V P Y E N L S I H C G E S M E L D L E A I	
TTTTGATCAAATTGTGAGGAGGAATAGAGGTGGCTGGTGTCCAGGTCAACTATCTTTGTACTGGGCTTTGACTACCACCGGTTTG	360
F D Q I V R R N R G G W C L Q V N Y E L L Y W A L T T T G F E	
AGACCCACAATGTTAGGAGGTTTGTGTTATGGCAGTAATAATGACAAATACAGCCTGGCATGATTACCTCATAGTACAGGTGACCATCA	450
T T M L G G F V Y G S N N D K Y T S G M I H L V Q V T I N	
ACGGCAGGAACATACATTGTGCGATGCTGGTTGGACGCTCTACCGAGATGTGGCAGCTGTGGAGTTAATTCTGGGAAAGATCAGCCTC	540
G R N Y I V D A G F G R S Y Q M W Q P V E L I S G K D Q P Q	
AA GTGCCTCCATATTCCGCTTGAGAGAAGAGGGAGAACCTGGACTTGGACCAAATCAGAACAGCACATGTTCCAGACCAAGAAT	630
V P S I F R L R E E G E T W Y L D Q I R R Q Q H V P D Q E F	
TTCTTAATTCTGAACTCTGGAAAAAGAAAATACACAAAATCTACTGCTTTACTCTCAACCTCGAACAAATTGAAGAATTGAGTC	720
G L N S E L L E K K I Y Q K L Y C F T L Q P R T I E E F F E S A	
CAACACACATCTCAAGAACCCCCATCATCTGTGTTAGAACAAATCTACTGCTCTGGCAGACCCAGAACGGTGTACTGTTGG	810
N T Y L Q E S P S S V F L D K S I C S L Q T P E G V H C L V	
TGGGCCTCACCCCTCACCTCCAGGACCTACAATTATAAGGAAAATACAGATCTGGTAGAGTTAAAGTCTGACTGAGGAAGATGGAGG	900
G L T L T S R T Y N Y K E N T D L V E F K V L T E E E V E G	
GACTGGCTGAAGAACTATATAATTCTCTGGGAAAAAAAGGTTGTGTCACAAATGCTATTGTCTTCACTATTAGAGTACGGAGG	990
V L K T I F N I S L G K K L V S K N G H L S F T I	
AAAAAAATACCTCTATCTTAGACAAGTACTTAAGCTTTTATATAAAATGATCAACACATGAAAGAAATAGGATAATAAA	1080
TTCATATGTTTATCAAAAAA 1102	

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