

Nucleotide sequence of a rat heart cDNA encoding the isotype β of the catalytic subunit of protein phosphatase 2A

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The screening of a rat heart cDNA library yielded a cDNA encoding the catalytic subunit of β isotype of protein phosphatase 2A. The predicted 309 amino acids protein is identical or almost identical to other type β mammalian phosphatases 2A. An unusual conservation between different mammalian species is observed when non coding sequences are considered.

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-44          TGCAGGTCGGCGGCCGGGGAGGGCGGGGGACGGCTGCCGCC
  1 ATGGACGACAAGGGCGTTCAACCAAGGAGCTGGACCGATGGGGAGCAGCTGAACGAGTGTAAAGCAGCTG
    M D D K A F T K E L D Q W V E Q L N E C K Q L
  70 AACGAGAACCAAGTGCAGCGCTGTGAGAAGGCTAAGGAAATTAAACAAAAGAACATCAAATGTACAA
    N E N Q V R T L C E K A K E I L T K E S N V Q
139 GAGGTCTCGCTGCTCTGTTACCGCTGTGGAGATGTGCATGGCCATTCCATGACCTTATGGAACTCTTC
    E V R C P V T V C G D V H G Q F H D L M E L F
208 AGAATGGTGGAAATCACCAAGACACCAACTATCTATTCTATGGTGACTATGTAGACAGAGGGTATTAT
    R I G G K S P D T N Y L F M G D Y V D R G Y Y
277 TCTGTGGAGACGGTGAECTCTCTGTAGCATTAAGGTGGCTATCCAGAGCGTATCACATAATTGCGA
    S V E T V T L L V A L K V R Y P E R I T I L R
346 GGAATCATGAAAGCCGGCAGATCACACAAGTATGGCTTTATGATGAATGCCATACGAAAGTATGGG
    G N H E S R Q I T Q V Y G F D E C L R K Y G
415 AACGCGAACGCTGTGAAATACTTACAGATCTCTTGATTATCTCCACTTACAGCTTAGATAGTGGAA
    N A N V W K Y F T D L F D Y L P L T A L V D G
484 CAGATATTCTGCCTCCACGGTGGCTCTCCATCCATAGATACACTGGATCACATAAGAGCCCTGGAT
    Q I C F L H G G L S P S I D T L D H I R A L D
553 CGCTTACAGGAAGTCCACATGAGGGCCCAATGTGTGATCTCTATGGTCAGATCCAGATGACCGTGGT
    R L Q E V P H B G P M C D L L W S D P D D R G
622 GGCTGGGGCATTCCTCCACGTGGCTGCTACACATTGGCAAGACATTCTGAACATTTAACCAT
    G W G I S P R G A G Y T F G Q D I S E T F N H
691 GCCAACGGCCCTCACACTGGTGTCCCGTGCTCACAGCTGTAAATGGAAGGATATAATTGGTGCCATGAT
    A N G L T L V S R A H Q L V M E G Y N W C H D
760 CGGAATGTGGTACCATTTAGTGCACCCAATTAAGTGTGTGCTACCGCTGTGGGAACCCAGGCTGTCTATCATG
    R N V V T I F S A P N Y C Y R C G N Q A A I M
829 GAATTAGACGACACTTTAAATACTCTTTCTCAAGTTGACCCAGCACCTCGTGTGGAGAGCCTCAT
    E L D D T L K Y S F L Q F D P A P R R G E P H
898 GTGACCCGGCGACCCCGAGACTCTCTATAAATTCTCCCCAGGACCTGTCTTGATGTTGAAGTA
    V T R R T P D Y F L
967 TACCTGCTTTAAAAATATATACATATATATATTTAAAAACACAGTTATCTGTGTCTCTGT
1036 ACAAAATGTGCTATGTCTTGACGTTAAAAACACATCATGGACCAAAACGTGCACATAATGGTGAGCCA
1105 TCAGCACGGGTGAACTTGAGTCACCTGCTCTGGCCACGGAGTCACCCAGGAGCCCTGGCCCTG
1174 CTGCTGTAGTAGCCGCTCTCGTGAATGGTTAAAGGAAAGGGTCACTGGTGCTTCATCTCTTCGCG
1243 CCTACTTGGAAATTAGTACAGTTAACGGCATGGATTATAGAGTTGGAGTTTATTTAAGAATT
1312 GACAAGCTGACTTCACTTAAATTCTACATACCCCTTATTTGTGAAATGTATGACTAACTGAAGAAGAG
1381 ATTCTGGAGATGTGTCTACAACTAAGATTTCTCAAGTTCTGACTGAATTACTGTGGAA
1450 TGTTGACCTGCACATTCTGTATATTGTCTGACAGTGTGCTACTGAACAAATAA
1519 ACTTCCAATTAGAGAGAAAAAAA

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