

A mouse cDNA sequence for epididymal androgen-regulated proteins related to glutathione peroxidase

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A cDNA was isolated from a murine epididymal cDNA library in λ gt11 by differential hybridization. This cDNA codes for 24kDa androgen-regulated epididymal secretory proteins (1). The sequence is presented below, along with the predicted amino acid sequence. Comparison of the present data with entries in protein sequence data bases revealed about 67% homology with the mouse glutathione peroxidase (2) at the amino acid level. This cDNA emphasizes previous observations suggesting that the major protective system against peroxide damage in mouse sperm could be a glutathione peroxidase-like system (3).

REFERENCES

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TCTCTTAATGGGAAGGAACACATTCCATTCAAGCAGTATCGAGGAAAGCACGTCCTCTTTGTCAATGTGGCTACC      75
S L N G K E H I P F K Q Y R G K H V L F V N V A T      25
TATTGGCGTCTGACAATCCAGTACCCCTGAGCTGAATGCACTCCAGGAGGATCTGAAGCCATTTGGCTTGGTTATA      150
Y C G L T I Q Y P E L N A L Q E D L K P F G L V I      50
TTGGGCTTTCCCTGCAACCAATTTGGAAAGCAAGAACCAGGAGACAATTTAGAGATTCTTCCCTGGGCTCAAGTAT      225
L G F P C N Q F G K Q E P G D N L E I L P G L K Y      75
GTTCTCCAGGAAAGGGTTTTTACCTAACTTCCAGCTTTTTGCAAAGGGGATGTAAATGGTGAACGAGCAG      300
V R P G K G F L P N F Q L F A K G D V N G E N E Q      100
AAAATCTTCACTTCTTGAAGCGTTCTTGTCTCACCCTCAGAGACTGTGGTCATGAGCAAACATACCTCCTGG      375
K I F T F L K R S C P H P S E T V V M S K H T S W      125
GAGCCAATAAAAGTCCATGACATCCGCTGGAACCTTGAAGAAGTTCTGGTGGGACCCGATGGCGTCCCTGTCTATG      450
E P I K V H D I R W N F E K F L V G P D G V P V M      150
CGCTGGTCCACCAGGCTCCTGTCTCAGCACTGTCAAGTCTGACATCATGGCGTACCTGAGCCATTTCAAACCCATA      525
R W F H Q A P V S T V K S D I M A Y L S H F K T I      175
TAGGAAGGCCAAGCTTCTGACCTTCTCCTTCCCCCTTAAAGACTGCTCTGAAAAAAGACTCCATCTTCTCA      600
End
GCACACTCTTCACTGAAATGGACTCTACCTCCCAAGTCACCCCTAAATTGCCTAAGTTCTTCCCCTGCACAAGTA      675
GATTTGTGTCTGGGAAGCTGTAGATGTTTTTCCCTTGTAGATTATGAGTTGAAGAGAGAAAATAAAATAAAAA      750
GAAAAAGCTAAATCCAGAGACCTCAGAGGTTGGCTGAGTATGTTAGTACTACCTATAATGTGCGCACTCAGCA      825
GACATTACAGACATTTGACAGAGTAAGCGCAGGAGGAACATGAATGGCAGGCCAGCCTAAGCTACAAGATATCA      900
TGTGTCCAAAAAATAAATAAATCCACGACCACCAACAACCCGATTGAACTACTCTAATTCACCAAAGGATA      975
TGGGGATAGCTTGGTTGAAGGCTGTATCTGAAGGAAGAGTCTTGGCCATTGAGTCTTTCTTCCCAGCCTG      1050
AAGGTGGAGAAAGAGCAATGGAGGCTGGTCAGACAATCTAGTTTGTCTCTGAAACTGTGTCTCTCTGAGACA      1125
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GCCAATCTGAGGGAGACTCTGAGCAGATTGACTGGCACAGGAGGAGGGCATCTCCCTGATGCCAGGATCGGGGAC      1275
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CTTTCATCATCTGAAGAGTAGGACCCCCACATGCAGACCCCAACCTGGGATTCTTCAACTTCGAAACTAGACAT      1425
ATTATTTCCAATAAAATGTTTTCTGAAGCAAAAAAATAAATAA      1471

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