

Molecular cloning of mouse cathepsin D

Michael J. Grusby*, Steve C. Mitchell and Laurie H. Glimcher¹

Harvard School of Public Health, Department of Cancer Biology, Boston, MA 02115 and ¹Harvard Medical School, Department of Medicine, Boston, MA 02115, USA

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In order to identify those gene products involved in T cell-mediated cytotoxicity, we have recently described the generation and screening of a cDNA library prepared from the murine cytotoxic T lymphocyte CT.4R (1). We report here the complete nucleotide and deduced amino acid sequences of one clone isolated from this library. The cDNA contains 1979 bp and contains a single long open reading frame encoding a protein of 410 residues. This cDNA shows 81% amino acid identity with the previously reported sequence for human cathepsin D (2, 3). Cathepsin D is a lysosomal protease thought to be present in all mammalian cells. Northern blot analysis of various mouse tissues demonstrated that the clone reported herein is expressed ubiquitously (data not shown). Interestingly, cathepsin D has recently been shown by immunoelectron microscopy to be present in elevated amounts extralysosomally in cytotoxic T cells (4),

suggesting an additional role for this molecule in these cells. The availability of this cDNA will allow further examination of this hypothesis.

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1  CGTCATCCTGCCTATAAGCCGGCGACCTCTGGCTTTAAGCTTTGCTCTCTTCGGGCGCCGCGACCATGAAGACTCCCGGC
      M K T P G
82  GTCTTGCTGCTCATTCTCGGCCCTCGGCTTCGTCCTCCTTCGCGATTATCAGAATCCCTCTGCGCAAGTTCACATCTATC
      V L L L I L G L L A S S S F A I I R I P L R K F T S I
163  CGTCGGACTATGACGGAGGTGGCGGCTCTGTGGAAGACCTGATCCTCAAAGGCCCATCACAAGTACTCCATGCAGTCA
      R R T M T E V G G S V E D L I L K G P I T K Y S M Q S
244  TCGCCTAAGACCACGGAGCCAGTGTCAAGTTACTCAAAACTACCTGGATGCCAGTACTATGGCGATATCGGCATCGGGA
      S P K T T E P V S E L L K N Y L D A Q Y Y G D I G I G
325  ACCCCGCGCAGTGTTCACAGTCGTCTTGATCTGGCTCCTTAACCTGTGGGTCCCTCCATTGCAAGATACTT
      T P P Q C F T V V F D T G S S N L W V P S I H C K I L
406  GACATAGCCTGCTGGGTCCACCATAAGTACAACAGTGACAAGTCCAGCACCTATGTGAAGAACGGCAGTCTTTGACATC
      D I A C W V H H K Y N S D K S S T Y V K N G T S F D I
487  CACTACGGCTCAGGAAGCCTCTCTGGGTACCTGAGCCAGGACACTGTATCGGTCCATGTAAGTCTGACCAGTCAAAGCA
      H Y G S G S L S G Y L S Q D T V S V P C K S D Q S K A
568  AGAGGTATCAAGGTGGAGAACAGATCTTTGGAGAAGCCACCAAGCAGCCTGGAATCGTATTGTTGACAGCAAGTTTGT
      R G I K V E K Q I F G E A T K Q P G I V F V A A K F D
649  GGCATCTTGGGCATGGGCTACCTCATATCTGTAAACACAGTGTCTCCGGTCTTTGACAACCTGATGCAACAGACTG
      G I L G M G Y P H I S V N N V L P V F D N L M Q Q K L
730  GTGGACAAGAATCTTCTCCTTACCTGAACAGGACCCAGAAGGGCAACCCGAGGAGAATAATGCTTGGTGGCACT
      V D K N I F S F Y L N R D P E G Q P G G E L M L G G T
811  GACTCCAAGTACTACCACGGGAGCTGTCTACCTGAACGTCACTCGAAAGCCCTACTGGCAGGTGCACATGGACCAGTTG
      D S K Y Y H G E L S Y L N V T R K A Y W Q V H M D Q L
892  GAGGTGGGCAATGAGCTGACCCTGTGCAAGGGAGGCTGTGAGGCTATTGTGGACACAGGACATCTCTTGGTGGGCGCT
      E V G N E L T L C K G G C E A I V D T G T S L L V G P
973  GTGGAAGAGGTGAAGGAGCTGCAGAAGGCCATCGGGGACAGTGCCTCTTATCCAGGGTGAATATGATTCCTTGTGAGAAG
      V E E V K E L Q K A I G A V P L I Q G E Y M I P C E K
1054  GTGTCCAGCCTGCCACGGTCTACCTGAAGCTAGGAGGCAAAACTATGAACTACACCAGACAAGTATATACTCAAGTA
      V S S L P T V Y L K L G G K N Y E L H P D K Y I L K V
1135  TCGCAGGGTGGAAAGACAATCTGCCTGAGTGGCTTACGGGAATGGACATACCCCTCCAGTGGGCGCTCTGGATCTCTG
      S Q G G K T I C L S G F M G M D I P P P S G P L W I L
1216  GCGCATGCTTCATTGGTTCTACTACACTGTGTTGACAGAGACAACAATAGGGTCCGCTTTGCCAATGCTGTGCTACTC
      G D V F I G S Y Y T V F D R D N N R V G F A N A V L
1297  TAACTTGCTCCTTCTCCACTGTGAGGAACTGGATCAGAGTCCAGTAGAGGAAGCCAGCCAGCCCATCCCTCCACCTGCC
1378  CCACTCACATAGTACACTCGCTTAGTGTGCTGGACCTTGGGGGAGACCTGGCTGGAGCTTGTCCAGCTTCTGTGTT
1459  CTGTGGTTCCTCACCTGGGTTAGATGCTGCCCCTGCTGCTGCTGAAGGAGGCAAGGCCACCCAGTACACGAGGCTG
1540  CCTCAAAGGCCCTACTGGTTAATAGCTGCTGAGATGATGCTGTCTGCTGCCGCCCTTTGCTGTGTGGGCACTACTC
1621  TGAAGCAGGCAAAATGGTCTTAGGATCCCTCCAGAAACCTGCTGTGACCAGCCATCACCAGCTTGGGGATGGCACC
1702  ACGTTCTACTGCCCTCCAACTCTGGCCTGGCAAGGCCTGAAGGTGAGCAGGAAGGAGCAAGAGGACAGAAGCAAACTA
1783  TGAACCTGGGGGTTACCTAGGGCTTACCCTGGCCCTCTGGGAAGGCATGCCCTAGCCTGGGGTAGAGGTAGGATGACTG
1864  ACTGGTTTTGGCTGGCCTCTGCTGCCCTCATCTGGGCTAGGCAGCTGGGAGCCAAAGTTGAAATACAATAAAGTTGTTT
1945  TGGCCTTGAAAAAATAAAAAAAAAAAAAAAAAAAAAA

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* To whom correspondence should be addressed