

Sequence of the human adenovirus type 3 protease

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Adenoviruses code for a chymotrypsin-like endoprotease specific for particular Gly-Ala bonds in several viral coded precursor proteins (1,2). To determine the conserved domains of this enzyme we are sequencing the gene in prototype viruses from different subgroups (3,4). Here we report a 1273 nucleotide sequence which includes the 208 amino acid (23,775 daltons) protease gene (*, translation start and stop) of Ad3, a member of subgroup B. Both strands were sequenced in M13 clones, the 1-strand being presented.

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1   ACCCAGCCAATTATCCATACCCGCTCATCGGAACTACTGCCGTTAAGAGTGTACCCAGAAAAAGTTCTCTGTG
74  CGACAGGACCATGTGGCGATTCCCTTCTCCAGCAACTTCATGTCCATGGGGGCCCTTACCGACCTGGGACAG
147 AACATGCTCTATGCCAACTCAGCCATGCGCTGGACATGACTTTTGAGGTGGATCCCATGGATGAGCCACCCCT
220 GCTTTATCTTCTTTTCGAAGTCTTCGACGTGGTACAGTGCACCAGCCACACCGGGCGTTCATCGAGGCCGTC
293 TACCTGCGCACACCGTTCTCGGCCGGCAACGCCACCACATAAGAAGCCTCTTGCTTCTTGAAGCAGCAGCTG
366 CAGCCATGACATGCGGGTCCGGAACGGCTCGAGCGAGCAAGAGCTCAAAGCCATCGTCCGAGACCTGGGCTG
439 CCGACCTATTTCCTGGGAACCTTTGACAAGCGTTTCCCGGGGTTTCATGGCCCCGACAAGCTCGCCTGGCCG
512 ATAGTCAAACTGCCGGACCGGAGACGGGGGAGAGCACTGGCTGGCTTTTGGTTGAAACCCGGCTACAACA
585 CCTGCTACCTTTTGTATCCTTTGGGTTCTCGGATGAGCGGCTCAAACAGATTTACCAGTTTGTAGTACGAGGG
658 GCTCCTGCGTGCAGTGCCTTGTACCAAAGACCGCTGCATCACCTGGAGAAGTCTACCCAAAGCGTGCAG
731 GGTCCGCGCTCAGCCGCTGTGGACTTTTTGCTGTATGTTCCCTTCATGCCTTTGTGCACTGGCCCGACCGCC
804 CCATGGACGGAAACCCACCATGAAGTTGGTACTGGGGTGTCCAACAGCATGCTCCAATCACCCCAAGTCCA
877 GCCACCTGCGCCGCAACAGGAGGTGCTATACCGCTTCTCTAAACCCCACTCATCTTACTTTCTGTTCTCAC
950 CGCGCGGCATGAAAGGGCCACCGCGTTTGACCGTATGGATATGCAATAAGTTCATGTAACCGTGTTCAT
1023 AAACAGCACTTATTTTTACATGCACTGAGGCTCTGGTTTTGGTCACTTTGTTTCATCACTTACTCAGAAGTGC
1096 AATGGGTTCTGGCGGGAGTCAGAGTGACCCGGGGCAGGGATACGTTGCGGAACGTAAACCTGTTCTGCCACT
1169 TGAACCTGGGGATTACCAGCTTGGAACTGGAATCTCGGAAAGGTGTCTTGCACAACCTTTGTGGTCAAGTTG
1242 CATAGCCCAAGCAGGTACAGGACAGAGATCT

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