



ADAM17 (NM_003183.6) - cDNA - 2026-05-02

AGCGGCGGCC GGAAGCTGGC TGAGCCGGCC TTTGGTAACG CCACCTGCAC -176
TTCTGGGGGC GTCGAGCCTG GCGGTAGAAT CTTCCCAGTA GGCGGCGCGG -126
GAGGGAAAAG AGGATTGAGG GGCTAGGCCG GGCGGATCCC GTCCTCCCC -76
GATGTGAGCA GTTTTCCGAA ACCCCGTCAG GCGAAGGCTG CCCAGAGAGG -26
TGGAGTCGGT AGCGGGGCCG GGAACATGAG GCAGTCTCTC CTATTCCTGA 25
CCAGCGTGGT TCCTTTCGTG CTGGCGCCGC GACCTCCGGA TGACCCGGGC 75
TTCGGCCCCC ACCAGAGACT CGAGAAGCTT GATTCTTTGC TCTCAGACTA 125
CGATATTCTC TCTTTATCTA ATATCCAGCA GCATTCGGTA AGAAAAAGAG 175
ATCTACAGAC TTCAACACAT GTAGAAACAC TACTAACTTT TTCAGCTTTG 225
AAAAGGCATT TTAAATTATA CCTGACATCA AGTACTGAAC GTTTTTCACA 275
AAATTTCAAG GTCGTGGTGG TGGATGGTAA AAACGAAAGC GAGTACACTG 325
TAAAATGGCA GGACTTCTTC ACTGGACACG TGGTTGGTGA GCCTGACTCT 375
AGGGTTCTAG CCCACATAAG AGATGATGAT GTTATAATCA GAATCAACAC 425
AGATGGGGCC GAATATAACA TAGAGCCACT TTGGAGATTT GTTAATGATA 475
CCAAAGACAA AAGAATGTTA GTTTATAAAT CTGAAGATAT CAAGAAATGTT 525
TCACGTTTGC AGTCTCCAAA AGTGTGTGGT TATTTAAAAG TGGATAATGA 575
AGAGTTGCTC CCAAAGGGT TAGTAGACAG AGAACCACCT GAAGAGCTTG 625 D201Efs*11
TTCATCGAGT GAAAAGAAGA GCTGACCCAG ATCCCATGAA GAACACGTGT 675
AAATTATTTG TGGTAGCAGA TCATCGCTTC TACAGATACA TGGGCAGAGG 725
GGAAGAGAGT ACAACTACAA ATTACTTAAT AGAGCTAATT GACAGAGTTG 775
ATGACATCTA TCGGAACACT TCATGGGATA ATGCAGGTTT TAAAGGCTAT 825
GGAATACAGA TAGAGCAGAT TCGCATCTCT AAGTCTCCAC AAGAGGTAAA 875 I284T

ACCTGGTGAA AAGCACTACA ACATGGCAAA AAGTTACCCA AATGAAGAAA 925
AGGATGCTTG GGATGTGAAG ATGTTGCTAG AGCAATTTAG CTTTGATATA 975
GCTGAGGAAG CATCTAAAGT TTGCTTGGCA CACCTTTTCA CATACCAAGA 1025
TTTTGATATG GGAACTCTTG GATTAGCTTA TGTTGGCTCT CCCAGAGCAA 1075
ACAGCCATGG AGGTGTTTTGT CCAAAGGCTT ATTATAGCCC AGTTGGGAAG 1125
AAAAATATCT ATTTGAATAG TGGTTTGACG AGCACAAAGA ATTATGGTAA 1175
AACCATCCTT ACAAAGGAAG CTGACCCTGT TACAACTCAT GAATTGGGAC 1225
ATAATTTTGG AGCAGAACAT GATCCGGATG GTCTAGCAGA ATGTGCCCCG 1275
AATGAGGACC AGGGAGGGAA ATATGTCATG TATCCCATAG CTGTGAGTGG 1325
CGATCACGAG AACAATAAGA TGTTTTCAA CTGCAGTAAA CAATCAATCT 1375
ATAAGACCAT TGAAAGTAAG GCCCAGGAGT GTTTTCAAGA ACGCAGCAAT 1425
AAAGTTTGTG GGAACTCGAG GGTGGATGAA GGAGAAGAGT GTGATCCTGG 1475
CATCATGTAT CTGAACAACG ACACCTGCTG CAACAGCGAC TGCACGTTGA 1525
AGGAAGGTGT CCAGTGCAGT GACAGGAACA GTCCCTGCTG TAAAACTGT 1575
CAGTTTGAGA CTGCCCAGAA GAAGTGCCAG GAGGCGATTA ATGCTACTTG 1625
CAAAGGCGTG TCCTACTGCA CAGGTAATAG CAGTGAGTGC CCGCCTCCAG 1675
GAAATGCTGA AGATGACACT GTTTGCTTGG ATCTTGCCAA GTGTAAGGAT 1725
GGGAAATGCA TCCCTTTCTG CGAGAGGGAA CAGCAGCTGG AGTCCTGTGC 1775
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GCCGCTGTGT GCCCTATGTC GATGCTGAAC AAAAGAACTT ATTTTTGAGG 1875
AAAGGAAAGC CCTGTACAGT AGGATTTTGT GACATGAATG GCAAATGTGA 1925
GAAACGAGTA CAGGATGTAA TTGAACGATT TTGGGATTTT ATTGACCAGC 1975
TGAGCATCAA TACTTTTGGG AAGTTTTTGG CAGACAACAT CGTTGGGTCT 2025
GTCCTGGTTT TCTCCTTGAT ATTTTGGATT CCTTTCAGCA TTCTTGTTCA 2075
TTGTGTGGAT AAGAAATTGG ATAAACAGTA TGAATCTCTG TCTCTGTTTC 2125
ACCCCAGTAA CGTCGAAATG CTGAGCAGCA TGGATTCTGC ATCGGTTTCGC 2175
ATTATCAAAC CCTTTCCTGC GCCCAGACT CCAGGCCGCC TGCAGCCTGC 2225
CCCTGTGATC CCTTCGGCGC CAGCAGCTCC AAAACTGGAC CACCAGAGAA 2275

TGGACACCAT CCAGGAAGAC CCCAGCACAG ACTCACATAT GGACGAGGAT 2325
GGGTTTGAGA AGGACCCCTT CCCAAATAGC AGCACAGCTG CCAAGTCATT 2375
TGAGGATCTC ACGGACCATC CGGTCACCAG AAGTGAAAAG GCTGCCCTCCT 2425
TTAAACTGCA GCGTCAGAAT CGTGTGACA GCAAAGAAAC AGAGTGC**TAA**
TTTAGTTCTC AGCTCTTCTG ACTTAAGTGT GCAAAATATT TTTATAGATT *50
TGACCTACAA ATCAATCACA GCTTGTATTT TGTGAAGACT GGAAGTGAC *100
TTAGCAGATG CTGGTCATGT GTTTGAACCTT CCTGCAGGTA AACAGTTCCTT *150
GTGTGGTTTG GCCCTTCTCC TTTTGAAAAG GTAAGGTGAA GGTGAATCTA *200
GCTTATTTTG AGGCTTTCAG GTTTTAGTTT TTAATAATATC TTTTGACCTG *250
TGGTGCAAAA GCAGAAAATA CAGCTGGATT GGGTTATGAA TATTTACGTT *300
TTTGTAATTT AATCTTTTAT ATTGATAACA GCACTGACTA GGGAAATGAT *350
CAGTTTTTTT TTATACACTG TAATGAACCG CTGAATATGA GGCATTTGGC *400
ATTTATTTGT GATGACAACCT GGAATAGTTT TTTTTTTTTT TTTTTTTTTT *450
TGCCTTCAAC TAAAAACAAA GGAGATAAAT CTAGTATACA TTGTCTCTAA *500
ATTGTGGGTC TATTTCTAGT TATTACCCAG AGTTTTTATG TAGCAGGGAA *550
AATATATATC TAAATTTAGA AATCATTTGG GTTAATATGG CTCTTCATAA *600
TTCTAAGACT AATGCTCTCT AGAAACCTAA CCACCTACCT TACAGTGAGG *650
GCTATACATG GTAGCCAGTT GAATTTATGG AATCTACCAA CTGTTTAGGG *700
CCCTGATTTG CTGGGCAGTT TTTCTGTATT TTATAAGTAT CTTCATGTAT *750
CCCTGTTACT GATAGGGATA CATGCTCTTA GAAAATTCAC TATTGGCTGG *800
GAGTGGTGGC TCATGCCGTGT AATCCCAGCA CTTGGAGAGG CTGAGGTTGC *850
GCCACTACAC TCCAGCCTGG GTGACAGAGT GAGACTCTGC CTCAAAAAAA *900
AAAAAATAAA AAAAAAATTC ACTATCTACA AACCTAGAAT ATTTAAAAATA *950
CAAAGATTGC CTGTTTTTCAA ACACTATTGA ATAAGAGGGT GAGATATTTT *1000
TTAACAACAA CAACAACAAA AAAACAGGT TGTTTTGAAT GTGATGAGCC *1050
AGCCAGGAGA TAGAATACTA CCTGCCCTTA GGGTTGGGGG CTGTCCCCAC *1100
AAGACTTGAT ACTTCAGAAA CCTTTTTTAT TGACCCACAA GCAGATATTT *1150
GAATTACTTC TTACTTTTATT GCTCCAGGAT TCTGGATGGG CTGCATTTAC *1200

TGTGTGAAGG ATAAAAATCA TTAGCCTGGA TTCTGATTTC TATAAATTGC *1250
CATTAAAAGC TTTTTTTCCC CTAAGAACTG AAATGTGCTC ACCAGCCAAA *1300
ACATTTTAAC TTGTAAATTT TGAGGGCAGT TAACCAAACC TGTGACTAAT *1350
CATATCTCCT CCTACCCCCC ATTTCCAAGG ACATTTGTTA CTCAGATACT *1400
TGTTATACTA ATACTTGAAC TTGTACCTTA TGGTATTTGC TATCTTTTAA *1450
CTAGTCATGA TATTCTTATA CTTTAGTTAC ACTTTTGGAA TTTGATACAA *1500
GGTTGAGTGG GGTGTGTGGG TGTATGTATG AGTGAAACAG TTCTCAAAAAG *1550
AATGTAAGAA AAACCATTTT TATAAAATTG TGACTTTTTTA AAAACATAGT *1600
CTTTGTCATT TATAGAATTA ACAAGCTGCT CAGGGTATAT TTTATAGCTG *1650
TAGCACTGAT ATCTGCATTA ATAAATACTG TCGAAACACA A

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