



STAT4 (NM_003151.4) - cDNA - 2026-08-03

GACTTTGGAC TTGAGCACTG CCTGGGACCT GTGCTGAGAG AGCGCTAGCA 1
TG TCTCAGTG GAATCAAGTC CAACAGTTAG AAATCAAGTT TTTGGAGCAG 51
GTGGATCAAT TCTATGATGA CAACTTTCCC ATGGAAATTC GGCATCTGTT 101
GGCCCAATGG ATTGAAAATC AAGACTGGA GGCAGCTTCT AACAATGAAA 151
CCATGGCAAC GATTCTTCTT CAAAATTGT TAATACAACT GGATGAACAG 201
TTAGGTCGTG TTTCCAAAGA GAAAAACCTA CTCTTGATAC ACAATCTAAA 251
AAGAATTAGG AAGGTCCTTC AGGGAAAATT TCATGGAAAT CCAATGCATG 301
TAGCTGTGGT TATTTCAAAC TGTTTAAGGG AAGAGAGGAG AATATTGGCT 351
GCAGCCAACA TGCCTGTCCA GGGGCCTCTA GAGAAATCCT TACAAAGTTC 401
TTCAGTTTCA GAAAGACAGA GGAATGTGGA GCACAAAGTG GCTGCCATTA 451
AAAACAGTGT GCAGATGACA GAACAAGATA CCAAATACTT AGAAGATCTG 501
CAAGACGAAT TTGACTACAG GTATAAAACA ATTCAGACAA TGGATCAGAG 551
TGACAAGAAT AGTGCCATGG TGAATCAGGA AGTTTTGACA CTGCAGGAAA 601
TGCTTAACAG CCTCGATTTT AAGAGAAAGG AGGCTCTCAG TAAAATGACC 651
CAAATCATCC ATGAGACAGA CCTGTTAATG AACACCATGC TCATAGAAGA 701
GCTGCAAGAC TGGAAGCGGC GGCAGCAAAT CGCCTGCATC GGGGGTCCAC 751
TCCACAATGG GCTCGACCAG CTTCAGAACT GCTTTACACT ATTGGCAGAA 801
AGTCTTTTCC AACTGAGAAG GCAATTGGAG AACTAGAGG AGCAATCTAC 851
CAAAATGACA TATGAAGGTG ATCCCATTCC AATGCAAAGA ACTCACATGC 901
TAGAAAGAGT CACCTTCTTG ATCTACAACC TTTTCAAGAA CTCATTTGTG 951
GTTGAGCGAC AGCCATGTAT GCCAACCCAC CCTCAGAGGC CGTTGGTACT 1001
TAAAACCCTA ATTCAGTTCA CTGTAAACT AAGGCTACTA ATAAAATTGC 1051

CAGAACTAAA CTATCAGGTA AAGGTTAAGG CATCAATTGA CAAGAATGTT 1101
TCAACTCTAA GCAACCGAAG ATTTGTACTT TGTGGAAC TA GTCAAAGC 1151
CATGTCTATT GAAGAATCTT CCAATGGGAG TCTCTCAGTA GAATTTTCGAC 1201
ATTTGCAACC AAAGGAAATG AAGTCCAGTG CTGGAGGTAA AGGAAATGAG 1251
GGCTGTCACA TGGTGACTGA AGAACTTCAT TCCATAACGT TTGAAACACA 1301
GATCTGCCTC TATGGCCTGA CCATAGATTT GGAGACCAGC TCATTGCCTG 1351
TGGTGATGAT TTCCAATGTC AGTCAGTTAC CTAATGCTTG GGCATCCATC 1401
ATTTGGTACA ACGTGTCAAC CAACGATTCC CAGAACTTGG TTTTCTTTAA 1451
TAATCCTCCA CCTGCCACAT TGAGTCAACT ACTGGAGGTG ATGAGCTGGC 1501
AGTTTTCATC GTACGTTGGT CGTGGTCTTA ACTCAGATCA ACTCCATATG 1551
CTGGCAGAGA AGCTTACAGT CCAATCTAGC TACAGTGATG GTCACCTCAC 1601
CTGGGCCAAG TTCTGCAAGG AACATTTACC TGGTAAATCA TTTACCTTTT 1651
GGACATGGCT TGAAGCAATA TTGGATCTAA TTAAGAAACA CATTCTTCCC 1701
CTTTGGATTG ATGGGTATGT CATGGGCTTT GTTAGCAAAG AGAAGGAACG 1751
GCTGTTGCTA AAGGATAAAA TGCCTGGCAC CTTTTTATTA AGATTCAAGT 1801
AAAGCCATCT CGGAGGAATA ACTTTCACCT GGGTGGACCA TTCTGAAAGT 1851
GGGGAAGTGA GATTCCACTC TGTAGAACCC TACAATAAAG GCCGGTTGTC 1901
TGCTCTGCCA TTCGCTGACA TCCTGCGAGA CTACAAAGTT ATTATGGCTG 1951
AAAACATTCC TGAAAACCCT CTGAAGTACC TATATCCTGA CATTCCCCAA 2001
GACAAAGCCT TCGGTAAACA CTACAGCTCT CAGCCTTGCG AAGTTTCAAG 2051
ACCAACAGAA AGGGGTGACA AAGGTTATGT TCCTTCTGTT TTTATCCCCA 2101
TCTCAACAAT CCGAAGTGAT TCAACAGAGC CACATTCTCC ATCAGACCTT 2151
CTTCCCATGT CTCCAAGTGT GTATGCGGTG TTGAGAGAAA ACCTGAGTCC 2201
CACAACAATT GAAACTGCAA TGAAGTCTCC TTATTCTGCT GAAAGACAGG *4
ATAAACTCTG ACGCACCAAG AAAGGAAGCA AATGAAAAAG TTTAAAGACT *54
GTTCTTTGCC CAATAACCAC ATTTTATTTT TTCAGCTTTG TAAATACCAG *104
GTTCTAGGAA ATGTTTGACA TCTGAAGCTC TCTTCACACT CCCGTGGCAC *154
TCCTCAATTG GGAGTGTTGT GACTGAAATG CTTGAAACCA AAGCTTCAGA *204

TAAACTTGCA AGATAAGACA ACTTTAAGAA ACCAGTGTTA ATAACAATAT *254

TAACAGAAGA

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