



TNFAIP3 (NM_001270508.2) - cDNA - 2026-03-03

GCAGTCTGCA GTCTTCGTGG CGGGCCAAGC GAGCTTGGAG CCCGCGGGGG -250
CGGAGCGGTG AGAGCGGCCG CCAAGAGAGA TCACACCCCC AGCCGACCCT -200
GCCAGCGAGC GAGCCCGACC CCAGGCGTCC ATGGAGCGTC GCCTCCGCCC -150
GGTCCCTGCC CCGACCCCCG CCTGCGGCGC GCTCCTGCCT TGACCAGGAC -100
TTGGGACTTT GCGAAAGGAT CGCGGGGCCG GGAGAGGTAA CCGCCGCGCC -50
TCCCGGAGAG GTAACCGCCG CGCCTCCCGG AGAGGTGTTG GAGAGCACAA 1
TGGCTGAACA AGTCCTTCCT CAGGCTT**T**GT ATTTGAGCAA TATGCGG**A**AA 51 **L10*** **K17***
GCTGTGAAGA TAC**G**GGAGAG AACTCCAGAA GACATTTTTA AACCTACTAA 101 **R22Q**
TGGGATCATT CATCATTTTA AAACCATGCA C**C**GATACACA CTGGAAATGT 151 **R45X**
TCAGAACTTG CCAGTTTTGT CCTCAGTTTC GGGAGATCAT CCACAAAGCC 201
CTCATCGACA GAAACATCCA GGCCACCCTG GAAAGCCAGA AGAAA**C**TCAA 251 **L83F**
CTGGTGT**C**GA GAAGTCCGG**A** AGCTTGTGGC GCTGAAAACG AAC**G**GTGACG 301 **p.W85GfsX11 R87X K91***
GCA**A**TTGCCT CATGCATGCC **A**CTTCTCAGT ACATGTGGGG CGTTCAGGAC 351 **N102S T108A**
ACAGACTTGG TACTGAGGAA GGCGCTGT**T**C AGCA**C**GCTCA AGGAAACAGA 401 **F127C T129M**
CAC**A**CGCAAC TTTAAATT**C**C GCTGGCAACT GGAGTCT**T**C **AA**A**T**CTCAGG 451 **D134fs R141C c.436-437deTC K148Nfs*68**
AATTTGTT**G**A AA**C**GGGGCTT TGCTATGATA CTCGGA**A**CTG **G**AATGATGAA 501 **E154* T155M W164***
TGGGACAATC TTATCAAAAT **G**GCTTCCACA GACA**C**ACCCA TGGCC**C**GAAG 551 **A175P T179K R183***
TGGACTT**C**AG TACAAC**T**CAC TG**G**AAGAAAT ACACATATTT GTCCTT**T**GCA 601 **Q187X E192K C596_598 del_A**
ACATCC**T**CAG AAGGCCA**A**TC ATTGT**C**ATTT CAGACAAAAT G**C**TAAGA**A**GT 651 **L203R I207L S217R**
TTGGAATCAG GTTCCAATT**T** CGCCC**C**TT**T**G AAAGTGGGTG GAATTTACTT 701 **p.F224Sfs*4 P226Lfs*2 p.L227***
GCCT**C**TC**C**AC TGGCCTGCC AGGAAT**G**CTA CAGATACCCC ATTGTTCTCG 751 **L236P C243Y**
GCTA**T**GACAG CCATCATTTT GTACCC**T**TGG TGACCCTGAA GGACAGTGG**G** 801 **D253_F257DEL p.P268Lfs*19**

CCTGAAATCC GAGCTGTTCC AC~~T~~TGTAAAC AGAGACCGGG GAAGATTTGA 851 [p.R271*](#) [P274Hfs](#) [L275P](#)
AGACTTAAAA GTTCAC~~T~~TTT TG~~A~~CAGATCC TGAAAATGAG ATGAAGGAGA 901 [T292P](#)
AGCTCTTA~~A~~A AGAGTA~~C~~TTA ATG~~G~~TGA~~T~~AG AAATCCCCGT CCAAGGCTGG 951 [p.L303fs](#) [p.Y306*](#) [V309dup](#) [I310T](#)
GACCATGGCA CAACTCATC~~T~~ CATCAATGCC GCAAAGT~~T~~GG AT~~G~~AA~~G~~CTAA 1001 [p.\(L324Ofs*7\)](#) [Exon7-8deletion](#) [E32*](#) [A333fs2](#)
CTT~~A~~CCAAAA ~~G~~AAATCAATC TGGTAGATGA TTACTTTTAA CTTGTTTCAGC 1051 [P336fs](#) [E338*](#)
ATGAGTACAA GAAA~~T~~GGCAG GAAAACAGCG AGCAGGGGAG GAGAGAGGGG 1101 [W356R](#)
CACGCC~~C~~AGA ATCCCATGGA ACCTTCCGTG CCCAGC~~T~~TT CTCTCATGGA 1151 [Q370Rfs*16](#) [p.\(N371Sfs*17\)](#) [D384G](#)
TGTA~~A~~AATGT GAAACGCCCA ACTGCC~~C~~TT CTTCATGTCT GTGAACACCC 1201
AGCCTTTATG CCATGAGTGC TCAGAGAGGC GGCAAAGAA TCA~~A~~AACAAA 1251 [Q415fs](#)
CTCCCAAAGC TGA~~A~~CTCAA GCCGGGCCCT GAGGGGCTCC CTGGCATG~~G~~C 1301 [A434*](#) [A434V](#)
GCTCGGGGCC TCTCGG~~G~~GAG AAGCCTATGA GCCCTTGGCG TG~~G~~AACCCTG 1351 [c.1316_1317del](#) [W448C](#) [N449Tfs*28](#)
AGGAGTCCAC TGGGGG~~G~~CCT CATTCCGCC CACCGACAGC ACCCAGCCCT 1401 [P457Afs*16](#)
TTTCTGTTCA GTGAGACCAC TGCCAT~~G~~AAG TG~~C~~AGGAGCC CCGGCTGCC 1451 [M476I](#) [C478*](#)
CTTCACACTG AATG~~T~~GCAGC ACAACGGATT TTGTGAACGT TGCCACAAC~~G~~ 1501 [V489Afs*7](#) [A501Ffs*193](#)
CC~~C~~GG~~C~~AACT TCACGCCAGC CACGCCCCAG ACCACACAAG GCACTTGGAT 1551 [R502W](#) [Q503*](#)
CCC~~G~~GGAAGT GCCAAGCCTG C~~C~~TCCAGGAT GTTACCAGGA CATT~~T~~AATGG 1601 [L525V](#)
GATCTGCAGT ACTTGCTTCA AAAGGACTAC AGCAGAG~~G~~CC TCCTCCAGCC 1651 [A547T](#)
TCAGCACCAG CCTCCCTCCT TCCTGTCACC AGCGTTCCAA GTCAGATCCC 1701
TCGCGGCTCG TCCGGAGCCC CTCCC~~C~~GCAT TCTTGCCACA GAGCTGGAAA 1751 [H577Afs*95](#)
CGACGCC~~C~~T GCTGGCTGCC TGTCT~~C~~AAGC TGCACGGACT CC~~T~~GGGGACA 1801 [A588Vfs*80](#) [Q593X](#) [G599Wfs*73](#)
GG~~A~~CGGG~~G~~AC GAGCAAGTGC AGAAAAGCCG GCTGCGTGTA TTTTGGGACT 1851 [T602S](#) [G603R](#) [p.T604Rfs*93](#)
CCAGAAAACA AGGGCTTTTG CACACTGTGT TTCATCGAGT ACAGAGAAAA 1901
CAAA~~C~~ATTTT GCTGCTGCCT CAGGGAAAGT CAGTCCC~~A~~CA GCGTCCAGGT 1951 [p.His636fsTer1](#) [T647P](#)
TCCAGAACAC CATTCCGTGC CTGGGGAGGG AATGCGGCAC CCTTGGAAGC 2001
ACCATGTTTG AAGGATACTG CCAGAAGTGT TTCA~~T~~TGAAG CTCAGAATCA 2051 [I679T](#)
GAGATTTTCA TGGGCCAAAA GGACAGAAGA GCAACTGAGA TCGAGCCAGC 2101
GCAGAGATGT GCCTCGAACC ACAC~~A~~AAGCA CCTCAAGGCC CAAGTGCGCC 2151 [Q709R](#)
CGGGCTCCT GCAAGAACAT CC~~T~~GGCCTGC CGCAGCGAGG AGCTCTGCAT 2201 [L725P](#)

GGAGTGTCAG CATCCCAACC AGAGGATGGG CCCTGGGGCC CACCGGGGTG 2251 Q737sfs*79
AGCCTGCCCC CGAAGACCCC CCCAAGCAGC GTTGCCGGGC CCCCGCCTGT 2301
GATCATTTTG GCAATGCCAA GTGCAACGGC TACTGCAACG AATGCTTTCA 2351
GTTCAAGCAG ATGTATGGCT **AA**CCGGAAAC AGGTGGGTCA CCTCCTGCAA *28 K786T
GAAGTGGGGC CTCGAGCTGT CAGTCATCAT GGTGCTATCC TCTGAACCCC *78
TCAGCTGCCA CTGCAACAGT GGGCTTAAGG GTGTCTGAGC AGGAGAGGAA *128
AGATAAGCTC TTCGTGGTGC CCACGATGCT CAGGTTTGGT AACCCGGGAG *178
TGTTCCCAGG TGGCCTTAGA AAGCAAAGCT TGTAAC TGGC AAGGGATGAT *228
GTCAGATTCA GCCCAAGGTT CCTCCTCTCC TACCAAGCAG GAGGCCAGGA *278
ACTTCTTTGG ACTTGGAAGG TGTGCGGGGA CTGGCCGAGG CCCCTGCACC *328
CTGCGCATCA GGACTGCTTC ATCGTCTTGG CTGAGAAAGG GAAAAGACAC *378
ACAAGTCGCG TGGGTTGGAG AAGCCAGAGC CATTCCACCT CCCCTCCCCC *428
AGCATCTCTC AGAGATGTGA AGCCAGATCC TCATGGCAGC GAGGCCCTCT *478
GCAAGAAGCT CAAGGAAGCT CAGGGAAAAT GGACGTATTC AGAGAGTGTT *528
TGTAGTTCAT GGTTTTTCCC TACCTGCCCG GTTCCTTTCC TGAGGACCCG *578
GCAGAAATGC AGAACCATCC ATGGACTGTG ATTCTGAGGC TGCTGAGACT *628
GAACATGTTT ACATTGACAG AAAACAAGC TGCTCTTTAT AATATGCACC *678
TTTTAAAAAA TTAGAATATT TTACTGGGAA GACGTGTAAC TCTTTGGGTT *728
ATTACTGTCT TTACTTCTAA AGAAGTTAGC TTGAACTGAG GAGTAAAAGT *778
GTGTACATAT ATAATATACC CTTACATTAT GTATGAGGGA TTTTTTTAAA *828
TTATATTGAA ATGCTGCCCT AGAAGTACAA TAGGAAGGCT AAATAATAAT *878
AACCTGTTTT CTGGTTGTTG TTGGGGCATG AGCTTGTTGTA TACACTGCTT *928
GCATAAATC AACCAGCTGC CTTTTTAAAG GGAGCTCTAG TCCTTTTTGT *978
GTAATCACT TTATTTATTT TATTACAAAC TTCAAGATTA TTTAAGTGAA *1028
GATATTTCTT CAGCTCTGGG GAAAATGCCA CAGTGTTCCT CTGAGAGAAC *1078
ATCCTTGCTT TGAGTCAGGC TGTGGGCAAG TTCCTGACCA CAGGGAGTAA *1128
ATTGGCCTCT TTGATACACT TTTGCTTGCC TCCCAGGAA AGAAGGAATT *1178
GCATCCAAGG TATACATACA TATTCATCGA TGTTTCGTGC TTCTCCTTAT *1228

GAAACTCCAG CTATGTAATA AAAAATACTATA CTCTGTGTTC TGTTAATGCC *1278
TCTGAGTGTC CTACCTCCTT GGAGATGAGA TAGGGAAGGA GCAGGGATGA *1328
GACTGGCAAT GGTCACAGGG AAAGATGTGG CCTTTTGTGA TGGTTTTATT *1378
TTCTGTTAAC ACTGTGTCCT GGGGGGGCTG GGAAGTCCCC TGCATCCCAT *1428
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TATTTGTGAT CCATAACTCT ACACAGCCTT TACTCATACT ATTAGGCACA *1678
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AAAAAAAGTA ATTATTTATA CTTATTATAA AAAGTATTTG AAATTTGCAC *1878
ATTTAATTGT CCCTAATAGA AAGCCACCTA TTCTTTGTTG GATTTCTTCA *1928
AGTTTTTCTA AATAAATGTA ACTTTTCACA AGAGTCAACA TTAAAAATA *1978
AATTATTTAA GAACA

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