



TNFRSF1A (NM_001065.4) - cDNA - 2024-12-04

ACTCTTCCCC TCCCACCTTC TCTCCCCTCC TCTCTGCTTT AATTTTCTCA -213
GAATTCTCTG GACTGAGGCT CCAGTTCTGG CCTTTGGGGT TCAAGATCAC -163
TGGGACCAGG CCGTGATCTC TATGCCCGAG TCTCAACCCT CAACTGTCAC -113
CCCAAGGCAC TTGGGACGTC CTGGACAGAC CGAGTCCCGG GAAGCCCCAG -63
CACTGCCGCT GCCACACTGC CCTGAGCCCA AATGGGGGAG TGAGAGGCCA -13
TAGCTGTCTG GCATGGGCCT CTCCACCGTG CCTGACCTGC TGCTGCCACT 38 c.15C>T c.36A>G
GGTGCTCCTG GAGCTGTTGG TGGGAATATA CCCCTCAGGG GTTATTGGAC 88 c.59T>C c.81T>C
TGGTCCCTCA CCTAGGGGAC AGGGAGAAGA GAGATAGTGT GTGTCCCCAA 138 D12E C15Y
GGAAATATA TCCACCCTCA AAATAATTCTG ATTGCTGTGTA CCAAGTGCCA 188 K19I Y20H Y20D Y20C H22Y H22R H22Q N25D S27S I28S C29R C29G C29F C29Y C29S C29W
C30R C30S C30Y C30F C33G C33Y
CAAAGGAACC TACTTGATACA ATGACTGTCC AGGCCCGGG CAGGATACGG 238 G36E T37I Y38C Y38S L39F D42DEL D42E C43R C43G C43Y C43S C43F P46L G47G Q48H T50M
T50K
ACTGCAGGGA GTGTGAGAGC GGCCTCCTTCA CCGCTTCAGA AAACCACTC 288 C52R C52G C52F C52Y C52S C52W R53G E54E C55R C55S C55Y S57_E64del E56D G58S G58Y
S59P F60L(264C>G) F60V F60S F60L(267A>G) T61P T61I T61N N65I N65K H66Y H66L H66P L67P
AGACACTGCC TCAGCTGCTC CAAATGCCGA AAGGAAATGG GTCAGGTGGA 338 H69fs C70R C70S C70G C70Y C73R C73Y C73W S74C C105Y Q82K p.Q111R V83L V83M
GATCTCTTCT TGCACAGTGG ACCGGGACAC CGTGTGTGGC TGCAGGAAGA 388 S86P S116del C88R C88G C88Y C88S T89A R92W R92P R92Q R121_D122insARHR D93H D93E T94T
V95M C96R C96Y C96F C96W C98R C98Y C98F
ACCAGTACCG GCATTATTGG AGTGAAAACC TTTTCCAGTG CTTCAATTGC 438 N101K Y103_R104DEL R104W R104Q H105P Y106C E109A F112I F112L F112C F112S C114R C114W
N116S
AGCCTCTGCC TCAATGGGAC CGTGCACCTC TCCTGCCAGG AGAACACAGAA 488 T124T V125M H126T K132R
CACCCTGTGC ACCTGCCATG CAGGTTTCTT TCTAAGAGAA AACGAGTGTG 538 V136M
TCTCCTGTAG TAACTGTAAG AAAAGCCTGG AGTGCACGAA GTTGTGCCA 588 K157K L167_G175del

CCCCAGATTG AGAATGTTAA GGGCACTGAG GACTCAGGCA CCACAGTGCT 638 I170N I170T V173D V173G G175C
GTTGCCCTG GTCATTTTCT TTGGTCTTTG CCTTTTATCC CTCTCTTCA 688 S197C S197S
TTGGTTTAAAT GTATCGCTAC CAACGGTGGG AGTCCAAGCT CTACTCCATT 738
GTTTGTGGGA AATCGACACC TGAAAAAGAG GGGGAGCTTG AAGGAACTAC 788
TACTAAGCCC CTGGCCCCAA ACCCAAGCTT CAGTCCCCT CCAGGCTTCA 838
CCCCACCCT GGGCTTCAGT CCCGTGCCCA GTTCCACCTI CACCTCCAGC 888 S261I F264 T2269del
TCCACCTATA CCCCGGTGA CTGTCCCAAC TTTGCGGCTC CCCGCAGAGA 938 P272H R283K
GGTGGCACCA CCTATCAGG GGGCTGACCC CATCCTTGCG ACAGCCCTCG 988 A330T
CCTCCGACCC CATCCCCAAC CCCCTTCAGA AGTGGGAGGA CAGCGCCAC 1038
AAGCCACAGA GCCTAGACAC TGATGACCCC GCGACGCTGT ACGCCGTGGT 1088 S321I L330L Y331X
GGAGAACGTG CCCCCGTTGC GCTGGAAGGA ATTCTGTGCGG CGCCTAGGGC 1138 R341R
TGAGCGACCA CGAGATCGAT CGGCTGGAGC TGCAGAACGG GCCTGCCTG 1188 R365H
CGCGAGGCGC AATACAGCAT GCTGGCGACC TGGAGGCGGC GCACGCCGCG 1238
GCGCGAGGCC ACGCTGGAGC TGCTGGGACG CGTGCTCCGC GACATGGACC 1288 R426L D398E
TGCTGGGCTG CCTGGAGGAC ATCGAGGAGG CGCTTTGCGG CCCC GCCG 1338 L412I
CTCCCGCCCC CGCCAGTCT TCTCAGATGA GGTGCGCCCC CTGCGGGCAG *20 P419L
CTCTAAGGAC CGTCTGCGA GATCGCCTTC CAACCCCACT TTTTCTTGGG *70 c.*64T>C
AAGGAGGGGT CCTGCAGGGG CAAGCAGGAG CTAGCAGCCG CCTACTTGGT *120
GCTAACCCCT CGATGTACAT AGCTTTTCTC AGCTGCCTGC GCGCCGCCGA *170
CAGTCAGCGC TGTGCGCGG GAGAGAGGTG CGCCGTGGGC TCAAGAGCCT *220
GAGTGGGTGG TTTGCGAGGA TGAGGGACGC TATGCCTCAT GCCCGTTTGG *270
GGTGTCTCA CCAGCAAGGC TGCTCGGGGG CCCCTGGTTC GTCCCTGAGC *320
CTTTTTTACA GTGCATAAGC AGTTTTTTTT GTTTTTGTTT TGTTTTGTTT *370
TGTTTTTAAA TCAATCATGT TACTACTAATA GAAACTTGGC ACTCCTGTGC *420
CCTCTGCCTG GACAAGCACA TAGCAAGCTG AACTGTCCTA AGGCAGGGGC *470
GAGCACGGAA CAATGGGGCC TTCAGCTGGA GCTGTGGACT TTTGTACATA *520
CACTAAAATT CTGAAGTTAA A

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