

TNFRSF1A (NM_001065.4) - cDNA - 2026-11-11

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

GGAAAATATA TCCACCCTCA AAATAATTCG ATTTGCTGTA 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 TACTTGTACA ATGACTGTCC AGGCCCGGGG CAGGATACGG 238 G36E T37I Y38C Y38S L39F D42DEL D42E C43R C43G C43Y C43S C43F P46L G47G Q48H T50M T50K

ACTGCAGGGA GTGTGAGAGC GGCTCCTTCA CCGCTTCAGA AAACCACCTC 288 C52R C52G C52F C52Y C52S C52W R53G E54E C55R C55S C55Y S57_E64del E56D G58S G58V 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 AGAAACAGAA 488 T124I V125M H126T K132R

CACCGTGTGC ACCTGCCATG CAGGTTTCTT TCTAAGAGAA AACGAGTGTG 538 V136M

TCTCCTGTAG TAACTGTAAG AAAAGCCTGG AGTGCACGAA GTTGTGCCTA 588 K157K L167_G175del

CCCCAGATTG AGAATGTTAA GGGCACTGAG GACTCAGGCA CCACAGTGCT 638 I170N I170T V173D V173G G175C

GTTGCCCCTG GTCATTTTCT TTGGTCTTTG CCTTTTATCC CTCCTCTTCA 688 S197C S197S

TTGGTTTAAT GTATCGCTAC CAACGGTGGA AGTCCAAGCT CTACTCCATT 738

GTTTGTGGGA AATCGACACC TGAAAAAGAG GGGGAGCTTG AAGGAACTAC 788

TACTAAGCCC CTGGCCCCAA ACCCAAGCTT CAGTCCCACT CCAGGCTTCA 838

CCCCCACCCT GGGCTTCAGT CCCGTGCCCA GTTCCACCTT CACCTCCAGC 888 S261I F264 T2269del

TCCACCTATA CCCCCGGTGA CTGTCCCAAC TTTGCGGCTC CCCGCAGAGA 938 P272H R283K

GGTGGCACCA CCCTATCAGG GGGCTGACCC CATCCTTGCG ACAGCCCTCG 988 A330T

CCTCCGACCC CATCCCCAAC CCCCTTCAGA AGTGGGAGGA CAGCGCCCAC 1038

AAGCCACAGA GCCTAGACAC TGATGACCCC GCGACGCTGT ACGCCGTGGT 1088 S321I L330L Y331X

GGAGAACGTG CCCCCGTTGC GCTGGAAGGA ATTCGTGCGG CGCCTAGGGC 1138 R341R

TGAGCGACCA CGAGATCGAT CGGCTGGAGC TGCAGAACGG GCGCTGCCTG 1188 R365H

CGCGAGGCGC AATACAGCAT GCTGGCGACC TGGAGGCGGC GCACGCCGCG 1238

GCGCGAGGCC ACGCTGGAGC TGCTGGGACG CGTGCTCCGC GACATGGACC 1288 R426L D398E

TGCTGGGCTG CCTGGAGGAC ATCGAGGAGG CGCTTTGCGG CCCCGCCGCC 1338 L412I

CTCCCGCCCG CGCCCAGTCT TCTCAGATGA GGCTGCGCCC CTGCGGGCAG *20 P419L

CTCTAAGGAC CGTCCTGCGA GATCGCCTTC CAACCCCACT TTTTTCTGGA *70 c.*64T>C

AAGGAGGGGT CCTGCAGGGG CAAGCAGGAG CTAGCAGCCG CCTACTTGGT *120

GCTAACCCCT CGATGTACAT AGCTTTTCTC AGCTGCCTGC GCGCCGCCGA *170

CAGTCAGCGC TGTGCGCGCG GAGAGAGGTG CGCCGTGGGC TCAAGAGCCT *220

GAGTGGGTGG TTTGCGAGGA TGAGGGACGC TATGCCTCAT GCCCGTTTTG *270

GGTGTCCTCA CCAGCAAGGC TGCTCGGGGG CCCCTGGTTC GTCCCTGAGC *320

CTTTTTCACA GTGCATAAGC AGTTTTTTTT GTTTTTGTTT TGTTTTGTTT *370

TGTTTTTAAA TCAATCATGT TACACTAATA GAAACTTGGC ACTCCTGTGC *420

CCTCTGCCTG GACAAGCACA TAGCAAGCTG AACTGTCCTA AGGCAGGGGC *470

GAGCACGGAA CAATGGGGCC TTCAGCTGGA GCTGTGGACT TTTGTACATA *520

CACTAAAATT CTGAAGTTAA A

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