



TRNT1 (NM_182916.3) - cDNA - 2024-05-19

AGCCCGGAAG TGC GCGTGGC GGC GGTGGCG GCTGCGGCAA CAGCGGGGCC -29
GATGTGTAGT TGGTGACTGC CTCTCCAGAT GCTGAGGTGC CTGTATCATT 22
GGCACAGGCC AGTGCTGAAC CGTAGGTGGA GTAGGCTGTG CCTCCGAAG 72
CAGTATCTAT TCACAATGAA GTTGCAGTCT CCCGAATTCC AGTCACTTTT 122 M30V
CACAGAGGA CTGAAGAGTC TGACAGAAAT ATTTGTCAAA GAGAATCACG 172 c.126_128delAGA T49*
AATTAAGAAT AGCAGGAGGA GCAGTGAGGG ATTTATTAAA TGGAGTAAAG 222
CCTCAGGATA TAGATTTTGC CACCACTGCT ACCCCTACTC AAATGAAGGA 272
GATGTTTCAG TCGGCTGGGA TTGGGATGAT AAACAACAGA GGAGAAAAGC 322 R99W
ACGGAACAAT TACTGCCAGG CTTTCATGAAG AAAATTTTGA GATTACTACA 372 E121D
CTACGGATTG ATGTCACCAC TGATGGAAGA CATGCTGAGG TAGAATTTAC 422 D128G
AACTGACTGG CAGAAAGATG CGGAACGCAG AGATCTCACT ATAAATTCTA 472 A148V R150C T154L I155T M158V
TGTTTTTAGG TTTTGATGGC ACTTIAATTTG ACTACTTTAA TGGTTATGAA 522 L166S F167Tfs*9 Y173E
GATTIAAAAA ATAAGAAAGT TAGATTTGTT GGACATGCTA AACAGAGAAT 572 p.Leu176X R190I
ACAAGAGGAT TATCTTAGAA TTTTAAGATA CTTCAAGTTT TATGGGAGAA 622 Q192* R203K
TTGTAGACAA ACCTGGTGAC CATGATCCTG AGACTTTGGA AGCAATTGCA 672 I223I
GAAAATGCAA AAGGCTTGGC TGGAATATCA GGAGAAAGGA TTTGGGTGGA 722
ACTGAAAAAA ATTCTTGTTG GTAACCATGT AAATCATTTG ATTCACCTTA 772
TCTATGATCT TGATGTGGCT CCTTATATAG GTTTACCTGC TAATGCAAGT 822
TTAGAAGAAT TTGACAAAGT CAGTAAAAAT GTTGATGGTT TTTCACCAAA 872 E77X
GCCAGTGACT CTTTTGGCCT CATTATTCAA AGTACAAGAT GATGTCACAA 922
AATTGGATTT GAGGTGAAG ATCGCAAAAG AGGAGAAAAA CCTTGGCTTA 972 p.Leu313Ser A316V p.(Glu318Argfs*11) c.948-949delAAinsGG
TTTAAGTTA AAAATAGGAA AGATTTAATT AAAGCAACAG ATAGTTCAGA 1022 I326I

CCCATTGAAA CCCTATCAAG ACTTCATTAT AGATTCTAGG GAACCTGATG 1072 D348V c.1054_1056+10del
CAACTACTCG TGTATGTGAA CTACTGAAGT ACCAAGGAGA GCACTGTCTC 1122
CTAAAGGAAA TGCAGCAGTG GTCCATTCCCT CCATTTCCTG TAAGTGGCCA 1172 W381*
TGACATCAGA AAAGTGGGCA TTTCTTCAGG AAAAGAAATT GGGGCTCTAT 1222 G405R
TACAACAGTT GCGAGAACAG TGGAAAAAA GTGGTTACCA AATGGAAAAA 1272 K416E S418fs c.1246del
GATGAAC TTC TGAGTTACAT AAAGAAGACC TAAAACTGAT GGCTACTAAA *17
AAGCAGAGCA TTTCTGGTAA GACTAAATTT TCTCCCCTCC CTCTTAATGA *67
GGTTTTAGAG ACTACACCAG AATAAAAGAC AGTTTAGGGG ACCTCTGTAG *117
AACAAACAAGG GTCTTATTTT GTGAATTATA TATTTCAAGA ACTAAACAGA *167
GATCCACCTT TCTGGATCTG ATTTATATCA CTGAAATGTA CAGTTCTTTT *217
GGAATAGTTT CACCTGAGAA AACATAGTTG GCTATTATCT ATCTTAACCT *267
GTTCAGGCTT TTAATAAAAA CTGTTTTTGC ATAGGGTAGT ACTAAGATCT *317
TAAAAAGTGG TAACTGTCTT GAAGAAAAAA CGTTTATTGT TTGTTTGCAA *367
TTGAAATAAC AGGGTTACCT TAACAATGAC TGTCTATGAT GTGTCAGTTC *417
TTATCTGAAT TCCAAAATAA ACCTGTGCTT AAAAAAGAAA TAATTGACCA *467
AGTAAGTTTG CATAAAATGT GAATACTAAA TGTGTCCCCA GTTGCTGGCA *517
TTCATATGTA CAGGATTTGT TCTAGCAAGC TATGCTTCAG TATGTGGTTG *567
ATATTTTTCT GTCACAATGA TTTCTTTATG CATGCAGAGC CTGGGGAAGT *617
CATGGGATTA ACTTGAGGGT CACTATTGAG CCTATTAATT AATTATTGTT *667
TTAATAAAAC AAACATTGGT ATTGGAAGAT AAATATGTTT ATGTGGTATC *717
TGACAATGTG TATTAGGTGT CATATACAAT GGTAAATATGC CTGTCTTTAA *767
AGTGTTATTT TATTAATTAA AAGGATATGG CTATTATTAT ATATTCTCTA *817
AAGATTTGAG TCCTAAATGC TTTCATCAGG TAAATAAAAT GTATAATACA *867

AA

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