



TRNT1 (NM_182916.3) - cDNA - 2025-02-13

AGCCCGGAAG TGCGCGTGGC GGCGGTGGCG GCTGCGGCAA CAGCGGGGCC -29
GATGTGTAGT TGGTGACTGC CTCTCCAGAT GCTGAGGTGC CTGTATCATT 22
GGCCACAGGCC AGTGCTGAAC CGTAGGTGGA GTAGGCTGTG CCTTCCGAAG 72
CAGTATCTAT TCACAATGAA GTTGCAGTCT CCCGAATTCC AGTCACTTTT 122 M30V
CACAGAAGGA CTGAAGAGTC TGACAGAATT ATTTGTCAA GAGAATCACG 172 c.126_128delAGA T49*
AATTAAGAAT AGCAGGAGGA GCAGTGAGGG ATTTATTTAA TGGAGTAAAG 222
CCTCAGGATA TAGATTTTGC CACCACTGCT ACCCCTACTC AAATGAAGGA 272 T82S P86S
GATGTTTCAG TCGGCTGGGA TTCGGATGAT AAACAACAGA GGAGAAAAGC 322 R99W
ACGGAACAAT TACTGCCAGG CTTCATGAAG AAAATTTTGA GATTACTACA 372 E121D
CTACGGATTG ATGTCACCAC TGATGGAAGA CATGCTGAGG TAGAATTTAC 422 D128G
AACTGACTGG CAGAAAGATG CGGAACGCAG AGATCTCACT ATTAAATCTA 472 A148V R150C T154L I155T M158V
TGTTTTTTAGG TTTTGATGGC ACTTTTAATTTG ACTTACTTTAA TGGTTATGAA 522 L166S F167Tfs*9 Y173F
GATTTAAAA ATAAGAAAGT TAGATTTGTT GGACATGCTA AACAGAGAAT 572 p.Leu176X R190I
ACAAGAGGAT TATCTTAGAA TTTTAAGATA CTTCAGGTTT TATGGGAGAA 622 Q192* R203K
TTGTAGACAA ACCTGGTGAC CATGATCCTG AGACTTTGGA AGCAATTGCA 672 I223T
GAAAATGCAA AAGGCTTGGC TGGAATATCA GGAGAAAGGA TTTGGGTGGA 722
ACTGAAAAAA ATTCTTGTG GTAACCATGT AAATCATTTG ATTCACCTTA 772
TCTATGATCT TGATGTGGCT CCTTATATAG GTTTACCTGC TAATGCAAGT 822
TTTAGAAGAAT TTGACAAAGT CAGTAAAAAT GTTGATGGTT TTTCACCAA 872 E77X
GCCAGTGACT CTTTTGGCCT CATTATTCAA AGTACAAGAT GATGTCACAA 922
AATTGATT GAGGTGAAG ATCGCAAAAG AGGAGAAAA CCTTGGCTTA 972 p.Leu313Ser A316V p.(Glu318Argfs*11) c.948-949delAAinsGG
TTTATAGTTA AAAATAGGAA AGATTTAATT AAGCAACAG ATAGTTCAGA 1022 I326T

CCCATTGAAA CCCTATCAAG ACTTCATTAT AGATTCTAGG GAACCTGATG 1072 D348V c.1054_1056+10del
CAACTACTCG TGTATGTGAA CTACTGAAGT ACCAAGGAGA GCACTGTCTC 1122
CTAAAGGAAA TGCAGCAGTG GTCCATTCCCT CCATTTCCCTG 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 TCTCCCTCC CTCTTAATGA *67
GGTTTTAGAG ACTACACCAG AATAAAAGAC AGTTTAGGGG ACCTCTGTAG *117
AACAAACAAGG GTCCTATTTT GTGAATTATA TATTTCAAGA ACTAAACAGA *167
GATCCACCTT TCTGGATCTG ATTTATATCA CTGAAATGTA CAGTTCCTTT *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 GGTAATATGC CTGTCTTTAA *767
AGTGTATTTT TATTAATTA AAGGATATGG CTATTATTAT ATATTCTCTA *817
AAGATTTGAG TCCTAAATGC TTTTCATCAGG TAAATAAAAT GTATAATACA *867

AA

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