



TRAP1 (NM_016292.3) - cDNA - 2024-12-22

GCTCTGGGAG TACGACATGG CGCGCGAGCT GCGGGCGCTG CTGCTGTGGG 34
GCCGCCGCCT GCGGCCTTTG CTGCGGGCGC CGGCGCTGGC GGCCGTGCCG 84
GGAGGAAAAC CAATTCTGTG TCCTCGGAGG ACCACAGCCC AGTTGGGCCC 134
CAGGCGAAAC CCAGCCTGGA GCTTGCAGGC AGGACGACTG TTCAGCACGC 184
AGACCGCCGA GGACAAGGAG GAACCCCTGC ACTCGATTAT CAGCAGCACA 234
GAGAGCGTGC AGGGTTCCAC TTCCAAACAT GAGTTCCAGG CCGAGACAAA 284
GAAGCTTTTG GACATTGTTG CCCGGTCCCT GTACTCAGAA AAAGAGGTGT 334
TTATACGGGA GCTGATCTCC AATGCCAGCG ATGCCTTGGG AAAACTGCGT 384 R128H
CACAAACTGG TGTCTGACGG CCAAGCACTG CCAGAAATGG AGATTCACTT 434
GCAGACCAAT GCCGAGAAAG GCACCATCAC CATCCAGGAT ACTGGTATCG 484
GGATGACACA GGAAGAGCTG GTGTCCAACC TGGGGACGAT TGCCAGATCG 534
GGGTCAAAGG CCTTCC TGGA TGCTCTGCAG AACCAGGCTG AGGCCAGCAG 584
CAAGATCATC GGCCAGTTTG GAGTGGGTTT CTACTCAGCT TTCATGGTGG 634
CTGACAGAGT GGAGGTCTAT TCCCGCTCGG CAGCCCCGGG GAGCCTGGGT 684
TACCAGTGGC TTTCAGATGG TTCTGGAGTG TTTGAAATCG CCGAAGCTTC 734
GGGAGTTAGA ACCGGGACAA AAATCATCAT CCACCTGAAA TCCGACTGCA 784
AGGAGTTTTC CAGCGAGGCC CGGGTGCAG ATGTGGTAAAC GAAGTACAGC 834
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GCAGGCCATC TGGATGATGG ACCCCAAGGA TGTCCTGAG TGGCAACATG 934
AGGAGTTCTA CCGCTACGTC GCGCAGGCTC ACGACAAGCC CCGCTACACC 984 R316H
CTGCACTATA AGACGGACGC ACCGCTCAAC ATCCGCAGCA TCTTCTACGT 1034
GCCCGACATG AAACCGTCCA TGTTTGATGT GAGCCGGGAG CTGGGCTCCA 1084

CGGTTGCACT GTACAGCCGC AAAGTCCTCA TCCAGACCAA GGCCACGGAC 1134
ATCCTGCCCA AGTGGCTGCG CTTTCATCCGA GGTGTGGTGG ACAGTGAGGA 1184
CATTCCCCTG AACCTCAGCC GGGAGCTGCT GCAGGAGAGC GCACTCATCA 1234
GGAAACTCCG GGACGTTTTA CAGCAGAGGC TGATCAAATT CTTCATTGAC 1284
CAGAGTAAAA AAGATGCTGA GAAGTATGCA AAGTTTTTTG AAGATTACGG 1334 Y444N
CCTGTTTCATG CGGGAGGGCA TTGTGACCGC CACCGAGCAG GAGGTCAAGG 1384
AGGACATAGC AAAGCTGCTG CGCTACGAGT CCTCGGCGCT GCCCTCCGGG 1434
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GAAGCTGATC TCTGTGGAGA CGGACATAGT CGTGGATCAC TACAAGGAGG 1684
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CAACGTGAAG GTGACCCCTCC GACTGGACAC CCACCCTGCC ATGGTCACCG 1834
TGCTGGAGAT GGGGGCTGCC CGCCACTTCC TGCGCATGCA GCAGCTGGCC 1884
AAGACCCAGG AGGAGCGCGC ACAGCTCCTG CAGCCCACGC TGGAGATCAA 1934
CCCCAGGCAC GCGCTCATCA AGAAGCTGAA TCAGCTGCGC GCAAGCGAGC 1984
CTGGCCTGGC TCAGCTGCTG GTGGATCAGA TATACGAGAA CGCCATGATT 2034
GCTGCTGGAC TTGTTGACGA CCCTAGGGCC ATGGTGGGCC GCTTGAATGA 2084
GCTGCTTGTC AAGGCCCTGG AGCGACACTG ACAGCCAGGG GGCCAGAAGG *19
ACTGACACCA CAGATGACAG CCCACCTCC TTGAGCTTTA TTTACCTAAA *69
TTTAAAGGTA TTTCTTAACC CGA

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