



RELA (NM_021975.4) - cDNA - 2024-11-22

ATTTCGCCT CTGGCGAATG GCTCGTCTGT AGTGCACGCC GCGGGCCCAG -35
CTGCGACCCC GGCCCCGCCC CCGGGACCCC GGCCATGGAC GAACTGTTCC 16
CCCTCATCTT CCCGGCAGAG CCAGCCCAGG CCTCTGGCCC CTATGTGGAG 66
ATCATTGAGC AGCCCAAGCA GCGGGGCATG CGCTTCCGCT ACAAGTGCGA 116
GGGGCGCTCC GCGGGCAGCA TCCCAGGCGA GAGGAGCACA GATACCACCA 166
AGACCCACCC CACCATCAAG ATCAATGGCT ACACAGGACC AGGGACAGTG 216
CGCATCTCCC TGGTCACCAA GGACCCCTCT CACCGGCCCT ACCCCCACGA 266
GCTTGTAGGA AAGGACTGCC GGGATGGCTT CTATGAGGCT GAGCTCTGCC 316
CGGACCGCTG CATCCACAGT TTCCAGAACC TGGGAATCCA GTGTGTGAAG 366
AAGCGGGACC TGGAGCAGGC TATCAGTCAG CGCATCCAGA CCAACAACAA 416
CCCCTTCAA GTTCCATATAG AAGAGCAGCG TGGGGACTAC GACCTGAATG 466
CTGTGCGGCT CTGCTTCCAG GTGACAGTGC GGGACCCATC AGGCAGGCCC 516
CTCCGCTGC CGCCTGTCTT TTCTCATCCC ATCTTTGACA ATCGTGCCCC 566
CAACACTGCC GAGCTCAAGA TCTGCCGAGT GAACCGAAAC TCTGGCAGCT 616
GCCTCGGTGG GGATGAGATC TTCCTACTGT GTGACAAGGT GCAGAAAGAG 666
GACATTGAGG TGTATTTAC GGGACCAGGC TGGGAGGCC GAGGCTCCTT 716
TTCGCAAGCT GATGTGCACG GACAAGTGGC CATTGTGTTT CGGACCCCTC 766 R246*
CCTACGCAGA CCCAGCCTG CAGGCTCCTG TCCGTGTCTC CATGCAGCTG 816
CGGCGGCCTT CCGACCGGGA GCTCAGTGAG CCCATGGAAT TCCAGTACCT 866
GCCAGATACA GACGATCGTC ACCGGATTGA GGAGAAACGT AAAAGGACAT 916
ATGAGACCTT CAAGAGCATC ATGAAGAAGA GTCCTTTCAG CGGACCCACC 966
GACCCCGGC CTCCACCTCG ACGCATTGCT GTGCCTTCCC GCAGCTCAGC 1016 R329*

TTCTGTCCCC AAGCCAGCAC CCCAGCCCTA TCCCTTTACG TCATCCCTGA 1066 p.Y349*
 GCACCATCAA CTATGATGAG TTTCCCACCA TGGTGTTTCC TTCTGGGCAG 1116
 ATCAGCCAGG CCTCGGCCCTT GGCCCCGGCC CCTCCCCAAG TCCTGCCCA 1166 Q385* Q389*
 GGCTCCAGCC CCTGCCCTG CTCCAGCCAT GGTATCAGCT CTGGCCCAGG 1216
 CCCCAGCCCC TGTCCCAGTC CTAGCCCCAG GCCCTCCTCA GGCTGTGGCC 1266 V410Sfs*4
 CCACCTGCCC CCAAGCCCAC CCAGGCTGGG GAAGGAACGC TGTCAGAGGC 1316 E438Rfs*9
 CCTGCTGCAG CTGCAGTTTG ATGATGAAGA CCTGGGGGCC TTGCTTGGCA 1366
 ACAGCACAGA CCCAGCTGTG TTCACAGACC TGGCATCCGT CGACAACTC 1416 T464Rfs*26 p.E473Rfs*18
 GAGTTTCAGC AGCTGCTGAA CCAGGGCATA CCTGTGGCCC CCCACACAAC 1466 H487Tfs*7
 TGAGCCCATG CTGATGGAGT ACCCTGAGGC TATAACTCGC CTAGTGACAG 1516
 GGGCCCAGAG GCCCCCCGAC CCAGCTCCTG CTCCACTGGG GGCCCCGGGG 1566
 CTCCCCAATG GCCTCCTTTC AGGAGATGAA GACTTCTCCT CCATTGCGGA 1616
 CATGGACTTC TCAGCCCTGC TGAGTCAGAT CAGCTCCTAA GGGGGTGACG *10
 CCTGCCCTCC CCAGAGCACT GGGTTGCAGG GGATTGAAGC CCTCCAAAAG *60
 CACTTACGGA TTCTGGTGGG GTGTGTTCCTA ACTGCCCCCA ACTTTGTGGA *110
 TGTCTTCCTT GGAGGGGGGA GCCATATTTT ATTCTTTTAT TGTCAGTATC *160
 TGTATCTCTC TCTCTTTTTG GAGGTGCTTA AGCAGAAGCA TTAACCTCTC *210
 TGGAAAGGGG GGAGCTGGGG AAACCTCAAAC TTTTCCCCTG TCCTGATGGT *260
 CAGCTCCCTT CTCTGTAGGG AACTCTGGGG TCCCCATCC CCATCCTCCA *310
 GCTTCTGGTA CTCTCCTAGA GACAGAAGCA GGCTGGAGGT AAGGCCTTTG *360
 AGCCACAAA GCCTTATCAA GTGTCTTCCA TCATGGATTC ATTACAGCTT *410
 AATCAAATA ACGCCCCAGA TACCAGCCCC TGTATGGCAC TGGCATTGTC *460
 CCTGTGCCTA ACACCAGCGT TTGAGGGGCT GGCCTTCCTG CCCTACAGAG *510
 GTCTCTGCCG GCTCTTTCCT TGCTCAACCA TGGCTGAAGG AAACCAAGTGC *560
 AACAGCACTG GCTCTCTCCA GGATCCAGAA GGGGTTTGGT CTGGGACTTC *610
 CTTGCTCTCC CTCTTCTCAA GTGCCTTAAT AGTAGGGTAA GTTGTTAAGA *660
 GTGGGGGAGA GCAGGCTGGC AGCTCTCCAG TCAGGAGGCA TAGTTTTTAC *710
 TGAACAATCA AAGCACTTGG ACTCTTGCTC TTTCTACTCT GAACTAATAA *760

ATCTGTTGCC AAGCTGG

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