



NLRC4 (NM_021209.4) - cDNA - 2024-05-17

AAGCCCCTGG CTGTTTATAC TCCGGAGGGT GTCCCCGTGC GTCATCGGTG -226
GAGTGGACCA AAAGTGGTGA TCTGTTTGCC CTGTGTGACC TTGCCAGAA -176
CCCTGCTGAC TGAGAGAACA CATCTGCTGG AAGTCCTCTG GGATTCAAGG -126
TACAGGGAAT GAAGAGTAGT TTTACAGAAA AAAGAGGACA ATATTGGGAT -76
CACCTTTGAC CTTTCCATTT GGAAATAATA TTTTCTATTG TGTATAGAA -26
AGGTGGGAAG CTTTCATCCA GAACAATGAA TTTCATAAAG GACAATAGCC 25
GAGCCCTTAT TCAAAGAATG GGAATGACTG TTATAAAGCA AATCACAGAT 75
GACCTATTTG TATGGAATGT TCTGAATCGC GAAGAAGTAA ACATCATTTG 125
CTGCGAGAAG GTGGAGCAGG ATGCTGCTAG AGGGATCATT CACATGATTT 175
TGAAAAAGGG TTCAAGAGTCC TGTAACCTCT TTCTTAAATC CCTTAAGGAG 225 [E64Rfs*4](#) [L70F](#)
TGGAACATC CTCTATTTCA GGACTTGAAT GGACAAAGTC TTTTTCATCA 275
GACATCA~~G~~AA GGAGACTTGG ACGATTTGGC TCAGGATTTA AAGGACTTGT 325 [E95*](#)
ACCATACCCC ATCTTTTCTG AACTTTTATC CCCTTGGTGA AGATA~~I~~TGAC 375 [I124I](#)
ATTATTTTTA ACTTGAAAAG CA~~C~~CTTCACA GAACCTGTCC TGTGGAGGAA 425 [I133I](#)
GGACCAACAC CATCACCGCG TGGAGCAGCT GACCCTGAAT GGCCTCCTGC 475
AG~~G~~CTCTTCA GAGCCCC~~I~~GC ATCATTGAAG GGAAT~~C~~T~~G~~G CAAAGGCAAG 525 [A160I](#) [C165R](#) [S171F](#) [G172S](#)
TCC~~A~~C~~T~~CTGC TGCAG~~C~~GAAT TGCCATGCTC TGGGGCTCCG GAAAGTGCAA 575 [I177A](#) [I177S](#) [R181X](#)
GGCTCTGACC AAGTTCAAAT TCGTCTTCTT CCTCC~~G~~TCTC AGCA~~G~~GGCCC 625 [R204H](#) [R207K](#)
AGGGTGGACT TTTTGAAACC CTCTGTGATC AACTCCTGGA TATACCTGGC 675
ACAATCAGGA AGCAGACATT CATGGCCATG CTGCTGAAGC TGCGGCAGAG 725
GGTTCTTTTC CTTCTTGATG GCTACAATGA ATTCAAGCCC CAGAACTGCC 775
CAGAAATCGA AGCCCTGATA AAGGAAAACC ACCGCTTCAA GAACATGGTC 825

ATCGTCACCA CTACCACTGA GIGCCTGAGG CACATACGGC AGTTTGGTGC 875 C283G I287T G291S
 CCTGACTGCT GAGGTGGGGG ATATGACAGA AGACAGCGCC CAGGCTCTCA 925
 TCCGAGAAGT GCTGATCAAG GAGCTTGCTG AAGGCTTGTT GCTCCAAATT 975 R310*
 CAGAAATCCA GGTGCTTGAG GAATCTCATG AAGACCCTC TCTTTGTGGT 1025 T337S T337N L339P V341L V341A
 CATCACTTGT GCAATCCAGA TGGGTGAAAG TGAGTTCCAC TCTCACACAC 1075 I343N
 AAACAACGCT GTTCCATACC TTCTATGATC TGTTGATACA GAAAAACAAA 1125
 CACAAACATA AAGGTGTGGC TGCAAGTGAC TTCATTCGGA GCCTGGACCA 1175 I387T H392del
 CTGTGGAGAC CTAGCTCTGG AGGGTGTGTT CTCCCACAAG TTTGATTTCTG 1225
 AACTGCAGGA TGTGTCCAGC GTGAATGAGG ATGTCCTGCT GACAACTGGG 1275
 CTCCTCTGTA AATATACAGC TCAAAGGTTT AAGCCAAAGT ATAAATTCTT 1325
 TCACAAGTCA TTCCAGGAGT ACACAGCAGG ACGAAGACTC AGCAGTTTAT 1375 H443P H443Q S445P
 TGACGTCTCA TGAGCCAGAG GAGGTGACCA AGGGGAATGG TTACTTGCAAG 1425
 AAAATGGTTT CCATTTTCGGA CATTACATCC ACTTATAGCA GCCTGCTCCG 1475
 GTACACCTGT GGGTCATCTG TGGAAGCCAC CAGGGCTGTT ATGAAGCACC 1525
 TCGCAGCAGT GTATCAACAC GGCTGCCTTC TCGGACTTTC CATCGCCAAG 1575
 AGGCCTCTCT GGAGACAGGA ATCTTTGCAA AGTGTGAAAA ACACCACTGA 1625
 GCAAGAAATT CTGAAAGCCA TAAACATCAA TTCCTTTGTA GAGTGTGGCA 1675
 TCCATTTATA TCAAGAGAGT ACATCCAAAT CAGCCCTGAG CCAAGAATTT 1725
 GAAGCTTTCT TTCAAGGTAA AAGCTTATAT ATCAACTCAG GGAACATCCC 1775
 CGATTACTTA TTTGACTTCT TTGAACATTT GCCCAATTGT GCAAGTGCCC 1825
 TGGACTTCAT TAAACTGGAC TTTTATGGGG GAGCTATGGC TTCATGGGAA 1875
 AAGGCTGCAG AAGACACAGG TGGAATCCAC ATGGAAGAGG CCCCAGAAAC 1925
 CTACATTCCC AGCAGGGCTG TATCTTTGTT CTTCAACTGG AAGCAGGAAT 1975 W655S W655C Q657L
 TCAGGACTCT GGAGGTCACA CTCCGGGATT TCAGCAAGTT GAATAAGCAA 2025
 GATATCAGAT ATCTGGGGAA AATATTGAGC TCTGCCACAA GCCTCAGGCT 2075
 GCAAATAAAG AGATGTGCTG GTGTGGCTGG AAGCCTCAGT TTGGTCCTCA 2125
 GCACCTGTAA GAACATTTAT TCTCTCATGG TGGAAGCCAG TCCCCTCACC 2175
 ATAGAAGATG AGAGGCACAT CACATCTGTA ACAAACCTGA AAACCTTGAG 2225

TATTCATGAC CTACAGAATC AACGGCTGCC GGGTGGTCTG ACTGACAGCT 2275 P752L delexon5
 TGGGTAACCTT GAAGAACCTT ACAAAGCTCA TAATGGATAA CATAAAGATG 2325 M775I
 AATGAAGAAG ATGCTATAAA ACTAGCTGAA GGCCCTGAAAA ACCTGAAGAA 2375 G786V
 GATGTGTTTT TTTCAATTTGA CCCACTTGTC TGACATTGGA GAGGGAATGG 2425
 ATTACATAGT CAAGTCTCTG TCAAGTGAAC CCTGTGACCT TGAAGAAATT 2475
 CAATTAGTCT CCTGCTGCTT GTCTGCAAAT GCAGTGAAAA TCCTAGCTCA 2525
 GAATCTTCAC AATTTGGTCA AACTGAGCAT TCTTGATTTA TCAGAAAAATT 2575
 ACCTGAAAA AGATGGAAAT GAAGCTCTTC ATGAACTGAT CGACAGGATG 2625
 AACGTGCTAG AACAGCTCAC CGCACTGATG CTGCCCTGGG GCITGTGACGT 2675 Q880E C890R
 GCAAGGCAGC CTGAGCAGCC TGTTGAAACA TTTGGAGGAG GTCCCACAAC 2725
 TCGTCAAGCT TGGGTTGAAA AACTGGAGAC TCACAGATAC AGAGATTAGA 2775
 ATTTTAGGTG CATTTTTTGG AAAGAACCCT CTGAAAAACT TCCAGCAGTT 2825 A929S
 GAATTTGGCG GGAAATCGTG TGAGCAGTGA TGGATGGCTT GCCTTCATGG 2875
 GTGTATTTGA GAATCTTAAG CAATTAGTGT TTTTGGACTT TAGTACTAAA 2925
 GAATTTCTAC CTGATCCAGC ATTAGTCAGA AACTTAGCC AAGTGTTATC 2975
 CAAGTTAACT TTTCTGCAAG AAGCTAGGCT TGTGGGGTGG CAATTTGATG 3025
 ATGATGATCT CAGTGTTATT ACAGGTGCTT TTAAACTAGT AACTGCTTAA
 ATAAAGTGTA CTCGAAGCCA GTAA

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