



NLRP7 (NM_001127255.2) - cDNA - 2026-05-25

CCCTTTTAT CCGTTTCCTT TGCAGGAAAC ACAGGCTGGA AGCAAGACCT -52
GACCTGAGGG AGTTCCTCAG CCTTAACCTA AGGTCTCATA CTCGGAGCAC -2
TATGACATCG CCCAGCTAG AGTGGACTCT GCAGACCCTT CTGGAGCAGC 49
TGAACGAGGA TGAATTAAAG AGTTTCAAAT CCCTTTTATG GGCTTTTCCC 99
CTCGAAGACG TGCTACAGAA GACCCCATGG TCTGAGGTGG AAGAGGCTGA 149
TGGCAAGAAA CTGGCAGAAA TTCTGGTCAA CACCTCCTCA GAAAATTGGA 199 T61TfsX7 W66* W66C
TAAGGAATGC GACTGTGAAC ATCTTGGAAG AGATGAATCT CACGGAATTG 249 N73N
TETAAGATGG CAAAGGCTGA GATGATGGAG GACGGACAGG TGCAAGAAAT 299 C84Y E99X
AGATAATCCT GAGCTGGGAG ATGCAGAAGA AGACTCGGAG TTAGCAAAGC 349 E113GfsX7 K116X
CAGGTGAAAA GGAAGGATGG AGAAATTCOA TGGAGAAACA GTCTTTGGTC 399 Q130Q L132Gfs*12
TGGAAGAACA CCTTTTGGCA AGGAGACATT GACAATTTCC ATGACGACGT 449 Q140* D149G
CACTCTGAGA AACCAACGGT TCATTCATT CTTGAATCCC AGAACACCCA 499 R156Q
GGAAGCTAAC ACCTTACACG GTGGTGTGC ACGGCCCCGC AGGCCTGGGG 549 V182M
AAAACCACGC TGGCCAAAA GTGTATGCTG GACTGGACAG ACTGCAACCT 599 T185del M192L W195X
CAGCCCGACG CTCAGATACG CGTTCTACCCT CAGCTGCAAG GAGCTCAGCC 649 S201S L210F
GCATGGGCC CTGTCAGTTT GCAGAGCTGA TCTCCAAAGA CTGCTCCTGAA 699 C221Lfs*13 W231X
TTGTCAGGATG ACATTCCAAG CATCCTAGCC CAAGCACAGA GAATCCTGTT 749 L234S A243V F250C
CGTGGTTCGAT GGCCTTGATG AGCTGAAAGT CCCACCTGGG GCGCTGATCC 799 F250L
AGGACATCTG CGGGGACTGG GAGAAGAAGA AGCCGCTGCC CGTCCCTCTG 849 K277Q V279L
GGGAGTTTGC TGAAGAGGAA GATGTTACCC AGGGCAGCCT TGCTGGTCAC 899
CACGCGGCC AGGGCACTGA GGGACCTCCA GCTCCTGGCG CAGCAGCCGA 949 A305A Q310R Q310Hfs*38 L311H L311I Y318CfsX7 Q314fs
TCTACGTAAG GGTGGAGGGC TTCCTGGAGG AGGACAGGAG GGCCTATTC 999 V319I R329K

CTGAGACACT TTGGAGACGA GGACCAAGCC ATGCGTGCCT TTGAGCTAAT 1049 D339D E340K E340Qfs*11
GAGGAGCAAC GCGGCCCTGT TCCAGCTGGG CTCGGCCCCC GCGGTGTGCT 1099 F357L
GGATTGTGTG CACGACTCTG AAGCTGCAGA TGGAGAAAGG GGAGGACC CG 1149 I368I I368M K379K K379N G380R P383L
GTCCCCACCT GCCTCACCCG CACGGGGCTG TTCTCTGCGTT TCCCTCTGCAG 1199 R390Afs*26 R390H T391A F394F L398R C399Y
CCGTTTCCCG CAGGGCGCAC AGCTGCGGGG CGCGCTGCGG ACGCTGAGCC 1249 L412P R413W R413Q
TCCTGGCCGC GCAGGGCCTG TGGGCGCAGA TGTCCGTGT TCCACCGAGAG 1299 M427T F430L F430F R432X
GACCTGGAAA GGCCTCGGGT GCAGGAGTCC GACCTCCGTC TGTTCCTGGA 1349 D434D L435L
CGGAGACATC CTCCGCCAGG ACAGAATCTC CAAAGGCTGC TACTCCTTCA 1399 L454L R458Sfs*69 Y464*
TCCACCTCAG CTTCCAGCAG TTTCTCAC TGCTGTCTA CGCCCTGGAG 1449 O472* A481T
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ACAGAAGCTG CTTTCCGGAG AAGAAAGACT CAAGAACC CC GACCTGATTC 1549 G506* E508DfsX27 K511R N512fs P513S P513L
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TTGGAGGCCA CTTTIGGCTG CCGGATGTCA CCGGACATCA AACAGGAATT 1649 F538L M542Tfs
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TACTTCTGAA GTGATGCATT GTTCTTTCAG CCTGAAGCAT TGTCAAGACT 1849 H613Rfs*8 S608F
TGCAGAAACT CTCACTGCAG GTAGCAAAGG GGGTGTTCCT GGAGAATTAC 1899 K619Nfs*18 S621Tfs*16
ATGGATTTTG AACTGGACAT TGAATTTGAA AGGTGCACTT ACCTAACCAT 1949 E637*
TCCGAAC TGG GCTCGGCAGG ATCTTCGCTC TCTTCGCCCTC TGGACAGATT 1999 P651S W653* D657V R659L L661F
TC TGTCTCTCT CTTCAGCTCA AACAGCAACC TCAAGTTTCT GGAAGTGAAA 2049 C668R S673X S675T L677PfsX6
CAAAGCTTCC TGAGTGACTC TTCTGTGCGG ATTCTTTGTG ACCACGTAAC 2099 R693W R693G R693P R693Q H698H V699I
CCGTAGCACC TGTATCTGCA AGAAAGTGGG GATTA AAAAC GTCACCCCTG 2149 R701C C704Y V714I P716A P716S P716LfsX21
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GGTTGGGAGG TCACTGTGCC ACCCCGAGC AGTGGGCTGA ATTCTTCTAT 2349 P775P E776Gfs*14 W778X
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GCTCCTGGAT GAGGGTGCCA TGTGCTGTA CAAGACCATG ACACGCCCAA 2449 [R815H](#)
 AACACTTCCT GCAGATGTIG TCGTTGGAAA ACITGTCGTCT TACAGAAGCC 2499 [L820Cfs*29](#) [L823X](#) [c.2482_2483delTG](#) [A833T](#)
 AGTTGCAAGG ACCTTGCTGC TGTCTTGGTT GTCAGCAAGA AGCTGACACA 2549
 CCTGTGCTTG GCCAAGAACC CCAITGGGGA TACAGGGGTG AAGTTTCTGT 2599 [I858Hfsx11](#) [I858T](#)
 GTGAGGGCTT GAGTTACCCT GATTGTAAAC TGCAGACCTT GGTGTTACAG 2649 [Y872*](#) [L880S](#)
 CAATGCAGCA TAACCAAGCT TGGCTGTAGA TAITCTCTCAG AGGCGCTCCA 2699 [Y894Y](#)
 AGAAGCCTGC AGCCTCACAA ACCCTGGACTT GAGTATCAAC CAGATAGCTC 2749 [A902A](#) [N907N](#) [N913S](#)
 GTGGATTGTG GATTCCTGT CAGGCATTAG AGAATCCAAA CITGTAACCTA 2799 [W920*](#) [W920*\(2760\)](#) [A925A](#) [L926*](#) [C931X](#)
 AAACACCTAC GCCTCTGGAG CTGCTCCCTC ATGCCTTTCT ATTGTCAGCA 2849
 TCTTGGATCT GCTCTCCTCA GCAATCAGAA GCTTGAAACT CTGGACCTGG 2899 [L964P](#)
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 TAATTTGGAA ATCAAGAAGC TGTGAGGGA AGTGAAAGAA AAGAATCCCA 3049
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 GACTTTTTTT GCTGAGCAGC CTGGGATCGC TCTACGAATT ACACAGGAAG *35 [c.*5C>T](#) [c.*20C>T](#)
 CGGGATTCGG GTCTCTAAGA TGTCTTATGA ATGCAGGTCA GAGGGTCACA *85 [c.*44G>A](#)
 TGTTAACACT AGAGTCTGTC GAGAGGTAGG ATTTGACACT GGTTTTCTCA *135
 CTATTTTTGG GAGATTCTGC ACGAGTCACG CACCCCTTC ACATGACGCT *185 [c.*180G>C](#)
 ATGTACTTTC TCACAGGGAT AATAAAGTTA GAGCACTCTC

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