



NLRP7 (NM_001127255.1) - cDNA - 2024-05-19

GAAACACAGG CTGGAAGCAA GACCTGACCT GAGGGAGTTC TTCAGCCTTA **-27**
ACCTAAGGTC TCATAACTCGG AGCACT**ATGA** CATGCCCCA GCTAGAGTGG **24**
ACTCTGCAGA CCCTTCTGGA GCAGCTGAAC GAGGATGAAT TAAAGAGTTT **74**
CAAATCCCTT TTATGGGCTT TTCCCCTCGA AGACGTGCTA CAGAAGACCC **124**
CATGGTCTGA GGTGGAAGAG GCTGATGGCA AGAAACTGGC AGAAATTCTG **174**
GTCAACACC CCTCAGAAAA TTGGATAAGG AATGCGACTG TGAACATTT **224** T61TfsX7 W66* N73N
GGAAGAGATG AATCTCACGG AATTGTGTA GATGCCAAAG GCTGAGATGA **274** C84Y
TGGAGGACGG ACAGGTGCAA GAAATAGATA ATCCTGAGCT GGGAGATGCA **324** E99X
GAAGAAGACT CGGAGTTAGC AAGCCAGGT GAAAAGGAAG GATGGAGAAA **374** E113GfsX7 K116X
TTCAATGGAG AAACAGTCTT TGGTCTGGAA GAACACCTT TGGCAAGGAG **424** Q130Q L132Gfs*12 Q140*
ACATTGACAA TTTCCATGAC GACGTCACTC TGAGAAACCA ACGGTTCAT **474** D149G R156Q
CCATTCTTGA ATCCCAGAAC ACCCAGGAAG CTAACACCTT ACACGGTGGT **524**
GCTGCACGGC CCCGCAGGC TGGGGAAAAC CACGCTGGCC AAAAAGTGTA **574** V182M T185del M192L
TGCTGGACTG GACAGACTGC AACCTCAGC CGACGCTCAG ATACGCGTTC **624** W195X S201S
TACCTCAGCT GCAAGGAGCT CAGCCGCATG GGCCCCTGCA GTTTGCAGA **674** L210F
GCTGATCTCC AAAGACTGGC CTGAATTGCA GGATGACATT CCAAGCATCC **724** W231X L234S
TAGCCCAAGC ACAGAGAACT CTGTTCGTGG TCGATGGCCT TGATGAGCTG **774** F250C F250I
AAAGTCCCAC CTGGGGCGCT GATCCAGGAC ATCTGCGGGG ACTGGGAGAA **824**
GAAGAAGCCG GTGCCCGTCC TCCTGGGGAG TTTGCTGAAG AGGAAGATGT **874** K277Q V279I
TACCCAGGGC AGCCTTGCTG GTCACCACGC GGCCCAGGGC ACTGAGGGAC **924** A305A
CTCCAGCTCC TGGCGCAGCA GCCGATCTAC GTAAGGGTGG AGGGCTTCCCT **974** Q310R Q310Hfs*38 L311H L311I Y318CfsX7 V319I
GGAGGAGGAC AGGAGGGCCT ATTCCTGAG ACACTTGGA GACGAGGACC **1024** R329K D339D E340K E340Qfs*11

AAGCCATGCG TGCCTTGAG CTAATGAGGA GCAACGCGGC CCTGTTCCAG 1074
CTGGGCTCGG CCCCGCGGT GTGCTGGATT GTGTGCACGA CTCTGAAGCT 1124 I368I I368M
GCAGATGGAG AAGGGGAGG ACCCGGTCCC CACCTGCCTC ACCCGCACGG 1174 K379K K379N G380R P383L R390Afs*26 R390H T391A
GGCTGTT CCT GCGTTCCCTC TGCGGCCGGT TCCCGCAGGG CGCACAGCTG 1224 F394F L398R C399Y
CGGGGCGCGC TGCGGACGCT GAGCCTCCTG GCCGCGCAGG GCCTGTGGGC 1274 L412P R413W R413Q
GCAGATGTCC GTGTCACCC GAGAGGACT GGAAAGGCTC GGGGTGCAGG 1324 M427T F430L F430F R432X D434D L435L
AGTCCGACCT CCGTCTGTT CTGGACGGAG ACATCCT CCG CCAGGACAGA 1374 L454L R458Sfs*69
GTCTCAAAG GCTGCTACTC CTTCATCCAC CTCAGCTTCC AGCAGTTCT 1424 Y464*
CACTGCCCTG TTCTACGCCG TGGAGAAGGA GGAGGGGGAG GACAGGGACG 1474 A481T E486GfsX42 G487E
GCCACGCCCTG GGACATCGGG GACGTACAGA AGCTGCTTTC CGGAGAAGAA 1524 A494T I497I G498R G506* E508DfsX27
AGACTCAAGA ACCCGACCT GATTCAAGTA GGACACTTCT TATTGGCCT 1574 K511R P513S H520Tfs*46 c.1548 1566dup;1566 1567insAF15169.2:q.106 4191
CGCTAACGAG AAGAGAGCCA AGGAGTTGGA GGCCACTTT GGCTGCCGGA 1624 F538L
TGTCAACCGGA CATCAAACAG GAATTGCTGC AATGCAAAGC ACATCTTCAT 1674 M542Tfs
GCAAATAAGC CCTTATCCGT GACCGACCTG AAGGAGGTCT TGGGCTGCCT 1724 E570X c.1719 1720insT
GTATGAGTCT CAGGAGGAGG AGCTGGCGAA GGTGGTGGTG GCCCGTTCA 1774 L575L E580* P590P
AGGAAATTTC TATTCACCTG ACAAAACTT CTGAAGT GAT GCATTGTTCC 1824 H613Rfs*8 S608F
TTCAGCCTGA AGCATTGTCA AGACTTGCAG AAACCTCAC TGCAGGTAGC 1874 K619Nfs*18 S621Tfs*16
AAAGGGGGTG TTCCTGGAGA ATTACATGGA TTT TGAAGTG GACATTGAAT 1924 E637*
TTGAAAGGTG CACTTACCTA ACCATTCCGA ACTGGGCTCG GCAGGATCTT 1974 P651S W653* D657V
CGCTCTCTTC GCCTCTGGAC AGATTTCTGC TCTCTCTCA GCTCAAACAG 2024 R659L L661F C668R S673X S675T
CAACCTCAAG TTTCTGGAG TGAAACAAAG CTTCTGAGT GACTCTTCTG 2074 L677PfsX6
TGCGGATTCT TTGTGACCA CTAACCGTA GCACCTGTCA TCTGCAGAAA 2124 R693W R693G R693P R693Q H698H V699I R701C C704Y
GTGGAGATTA AAAACGTCAC CCTGACACC CGTACCGGG ACTTCTGTCT 2174 V714I P716A P716LfsX21 A719V R721W R721Q D722G
TGCTTCATT GGGAAAGAAGA CCCTCACCGA CCTGACCCCTG GCAGGGCACA 2224 H735P
TCGAGTGGGA ACGCACGATG ATGCTGATGC TGTGTGACCT GCTCAGAAAT 2274 I742I E743* L750V
CATAAAATGCA ACCTGCAGTA CCTGAGGTTG GGAGGTCACT GTGCCACCCC 2324 C761Y
GGAGCAGTGG GCTGAATTCT TCTATGTCT CAAAGCCAA CAGTCCCTGA 2374 P775P E776Gfs*14 W778X N788N
AGCACCTGCG TCTCTCAGCC AATGTGCTCC TGGATGAGGG TGCCATGTTG 2424 R795C

CTGTACAAGA CCATGACACG CCCAAACAC TTCCTGCAGA TGTTGTCGTT 2474 R815H L820Cfs*29 L823X
GGAAAACTGT CGTCTTACAG AAGCCAGTTG CAAGGACCTT GCTGCTGTCT 2524 c.2482_2483delTG A833T
TGGTTGTCAG CAAGAACCTG ACACACCTGT GCTTGGCCAA GAACCCCATT 2574 I858HfsX11 I858T
GGGGATACAG GGGTGAAGTT TCTGTGTGAG GGCTTGAGTT ACCCGTATTG 2624 Y872*
TAAACTGCAG ACCTTGGTGT TACAGCAATG CAGCATAACC AAGCTTGGCT 2674 L880S
GTAGATATCT CTCAGAGCG CTCCAAGAAG CCTGCAGCCT CACAACCTG 2724 Y894Y A902A N907N
GACTTGAGTA TCAACCAGAT AGCTCGTGG ATTGTGGATTC TCTGTCAGGC 2774 N913S W920* W920*(2760)
ATTAGAGAAAT CCAAACTGTA ACCTAAAACA CCTACGCCCT GGGAGCTGCT 2824 A925A L926* C931X
CCCTCATGCC TTTCTATTGT CAGCATCTT GATCTGCTCT CCTCAGCAAT 2874
CAGAAGCTTG AAACTGGA CCTGGGCCAG AATCATTGT GGAAGAGTGG 2924 L964P
CATAATTAAG CTCTTTGGGG TTCTAAGACA AAGAACTGG ATCTGAAGA 2974 G981Rfs
TACTCAGGTT GAAGACCTAT GAAACTAATT TGGAAATCAA GAAGCTGTTG 3024
GAGGAAGTGA AAGAAAAGAA TCCCAAGCTG ACTATTGATT GCAATGCTTC 3074 D1021V
CGGGGCAGC GCACCTCCGT GCTGTGACTT TTTTGCTGA GCAGCCTGGG *10 T1028A c.*5C>T
ATCGCTCA GAATTACACA GGAAGCGGG TTCGGGTCTC TAAGATGTCT *60 c.*20C>T c.*44G>A
TATGAATGCA GGTCAGAGGG TCACATGTTA ACACTAGAGT CTGTCGAGAG *110
GTAGGATTG ACACGGTTT TCTCACTATT TTTGGGAGAT TCTGCACGAG *160
TCACGCACCC CCTTCACATG ACGCTATGTA CTTTCTCAC A GGGATAATAA *210 c.*180G>C
AGTTAGAGCA CTCTCGTTGC A

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