



NOD2 (NM_022162.3) - cDNA - 2024-12-04

ACTTACTTGT GGCCTGTCCC CTCGTGAATG TGTCTCATGT CCCAGTGGG -247
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TCCGGCTTTT CCTTTGGGAA TTTCCCTTGA AGGTGGGGTT GGTAGACAGA -97
TCCAGGCTCA CCAGTCCTGT GCCACTGGGC TTTTGGCGTT CTGCACAAGG -47 c.-53C>T
CCTACCCGCA GATGCCATGC CTGCTCCCC AGCCTAATGG GCTTTGATGG 4
GGGAAGAGGG TGGTTCAGCC TCTCACGATG AGGAGGAAAG AGCAAGTGTC 54
CTCCTCGGAC ATTCTCCGGG TTGTGAAATG TGCTCGCAGG AGGCTTTTCA 104
GGCACAGAGG GAGCCAGCTGG TCGAGCTGCT GGTCTCAGGG TCCCTGGAAG 154 R38M E43Y
GCTTCGAGAG TGTCCTGGAC TGGCTGCTGT CCTGGGAGGT CCTCTCCTGG 204
GAGGACTACG AGGGCTTCCA CCTCCTGGGC CAGCCTCTCT CCCACTTGGC 254 L81Y
CAGGCGCCTT CTGGACACCG TCTGGAATAA GGGTACTTGG GCCTGTCAGA 304 T91A
AGTCATCGC GGCTGCCCAA GAAGCCCAGG CCGACAGCCA GTCCCCAAG 354 A105A D113N
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GAGTCACCGG CCAGCCATTG TCAGGAGGCT CCACAGCCAT GTGGAGAACA 454 R138Q A140T
TGCTGGACCT GGCATGGGAG CGGGGTTTCG TCAGCCAGTA TGAATGTGAT 504 D154N L155Q W157R
GAAATCAGGT TGCCGATCTT CACACCGTCC CAGAGGGCAA GAAGGCTGCT 554 S178S
TGATCTTGCC ACGGTGAAAG CGAATGGATT GGCTGCCTTC CTTCTACAAC 604 T189M T189T F198L
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TGCAAGAAGT ATATGGCCAA GCTGAGGACC ACGGTGTCTG CTCAGTCTCG 704 K225M Q233X R235C
CTTCCTCAGT ACCTATGATG GAGCAGAGAC CGCTCTGCCTG GAGGACATAT 754 T245M L248R
ACACAGAGAA TGTCTGGAG GTCTGGGCAG ATGTGGGCAT GGCTGGACCC 804 P268S/SNP5

CCGCAGAAGA GCCCAGCCAC CCTGGGCCTG GAGGAGCTCT TCAGCACCCC 854
TGGCCACCTC AATGACGATG CGGACACTGT GCTGGTGGTG GGTGAGGCGG 904 H287Y N289S D291N A292V T294S V298V A301V
GCAGTGGCAA GAGCACGCTC CTGCAGCGGC TGCAC TTGCT GTGGGCTGCA 954 R311W
GGGCAAGACT TCCAGGAATT TCTCTTTGTC TTCCCATTC A GCTGCCGGCA 1004 R334W R334O
GCTGCAGTGC ATGGCCAAAC CACTCTCTGT GCGGACTCTA CTCTTTGAGC 1054 L348V L349F
ACTGCTGTTG GCCTGATGTT GGTCAAGAAG ACATCTTCCA GTTACTCCTT 1104 H352R W355X D357A I363F
GACCACCCCTG ACCGTGTCTT GTTAACCTTT GATGGCTTTG ACGAGTTCAA 1154 H343Y R373C D382N D382E E383K E383G
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GCTGATGTGC CACCAGCTTT GCTCAGACAC CTCTTCAATT GTGGCAGGCC 1904
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CGGAGGGAAA GGACAGCAGC GTGGCAGCTT TGCTGCAGAA GGCCGAGCCG 2004 E667K P668L
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TTGACATTTT GCAGTGTGGG CCCCCTGAG TGTGCTGCCC TGGCCTTTGT 2354 [E778K](#)
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GAAAATCCTG AAGTTGTCCA ATAACATGCAT CACCTACCTA GGGGAGAAAG 3004 [c.2998G>A](#)
CCCTCCTGCA GGCCCTTGAA AGGAATGACA CCATCCTGGA AGTCTGGCTC 3054 [1007FS/SNP13](#)
CGAGGGAACA CTTTCTCTCT AGAGGAGGTT GACAAGCTCG GCTGCAGGGA 3104 [R1019X](#) [R1019G](#) [R1019L](#)
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GCAGCCTCTT CAAAATGAGC CCTGTCTGTC CTAAGGCTGA ACTTGTTTTC *131
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CAAGATTCAA TCCCAGGATG TACAAGGACA GCCCTCCTC CATAGTATGG *281
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ATGGATGTGC TTGTTAACTG AGTGCCTTTT GGTGGAGAGG CCCGCCTCT *381 [c.3498G>A](#)
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ATAATTCAGG AAGCAGCTTT CCCCATGTCT CGACTCATCC ATCCAGGCCA *481

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