



*IL1R1* (NM\_000877.4) - cDNA - 2026-06-14

AGTTC~~CC~~GGC CGCGAGGGCG GGC~~GC~~AGCTT GTGGCCGGCG GCCGGAGCCG -309  
ACTCGGAGCG CGCGGCGCCG GCCGGGAGGA GCCGGAGAGC GGCCGGGGCCG -259  
GGCGGTGGGG GCGCCGGCCT GCCCCGCGCG CCC~~CAG~~GGAG CGGCAGGAAT -209  
GTGACAATCG CGCGCCCGCG CACCGAAGCA CTCCTCGCTC GGCTCCTAGG -159  
GCTCTCGCCC CTCTGAGCTG AGCCGGGTTC CGCCCGGGGC TGGGATCCCA -109  
TCACCC~~TCCA~~ CGGCCGTCCG TCCAGGTAGA CGCACCC~~TCT~~ GAAGATGGTG -59  
ACTCCCTCCT GAGAAGCTGG ACCCCTTGGT AAAAGACAAG GCCTTCTCCA -9  
AGAAGAATAT GAAAGTGTTA CTCAGACTTA TTTGTTTCAT AGCTCTACTG 42  
ATTTCTTCTC TGGAGGCTGA TAAATGCAAG GAACGTGAAG AAAAAATAAT 92  
TTTAGTGTCA TCTGCAAATG AAATTGATGT TCGTCCCTGT CCTCTTAACC 142  
CAAATGAACA CAAAGGCACT ATAAC~~TTGGT~~ ATAAAGATGA CAGCAAGACA 192  
CCTGTATCTA CAGAACAAGC CTCCAGGATT CATCAACACA AAGAGAAACT 242  
TTGGTTTGTT CCTGCTAAGG TGGAGGATTC AGGACATTAC TATTGCGTGG 292  
TAAGAAATTC ATCTTACTGC CTCAGAATTA AAATAAGTGC AAAATTTGTG 342  
GAGAATGAGC CTAAC~~TTATG~~ TTATAATGCA CAAGCCATAT TTAAGCAGAA 392 ~~K131E~~  
ACTACCCGTT GCAGGAGACG GAGGACTTGT GTGCCCTTAT ATGGAGTTTT 442  
TTAAAAATGA AAATAATGAG TTACCTAAAT TACAGTGGTA TAAGGATTGC 492  
AAACCTCTAC TTCTTGACAA TATACACTTT AGTGGAGTCA AAGATAGGCT 542  
CATCGTGATG AATGTGGCTG AAAAGCATAG AGGGAAC~~TAT~~ ACTTGT~~CATG~~ 592  
CATCCTACAC ATACTTGGGC AAGCAATATC CTATTACCCG GGTAATAGAA 642  
TTTATTACTC TAGAGGAAAA CAAACCCACA AGGCCTGTGA TTGTGAGCCC 692  
AGCTAATGAG ACAATGGAAG TAGACTTGGG ATCCCAGATA CAATTGATCT 742

GTAATGTCAC CGGCCAGTTG AGTGACATTG CTTACTGGAA GTGGAATGGG 792  
TCAGTAATTG ATGAAGATGA CCCAGTGCCTA GGGGAAGACT ATTACAGTGT 842  
GGAAAATCCT GCAAACAAAA GAAGGAGTAC CCTCATCACA GTGCTTAATA 892  
TATCGGAAAT TGAAAGTAGA TTTTATAAAC ATCCATTTAC CTGTTTTGCC 942  
AAGAATACAC ATGGTATAGA TGCAGCATAT ATCCAGTTAA TATATCCAGT 992  
CACTAATTTT CAGAAGCACA TGATTGGTAT ATGTGTCACG TTGACAGTCA 1042  
TAATTGTGTG TTTCTGTTTTT ATCTATAAAA TCTTCAAGAT TGACATTGTG 1092  
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TGGAAAGACC TATGACGCAT ATATACTGTA TCCAAAGACT GTTGGGGAAG 1192  
GGTCTACCTC TGACTGTGAT ATTTTTGTGT TTAAAGTCTT GCCTGAGGTC 1242  
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GAAGACTGAT TATCATTTTA GTCAGAGAAA CATCAGGCTT CAGCTGGCTG 1392  
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TGGTCAGGGG ACTTTACACA GGGACCACAG TCTGCAAAGA CAAGGTTCTG 1592  
GAAGAATGTC AGGTACCACA TGCCAGTCCA GCGACGGTCA CCTTCATCTA 1642  
AACACCAGTT ACTGTCACCA GCCACTAAGG AGAAACTGCA AAGAGAGGCT 1692  
CACGTGCCCTC TCGGGTAGCA TGGAGAAGTT GCCAAGAGTT CTTTAGGTGC \*32  
CTCCTGTCTT ATGGCGTTGC AGGCCAGGTT ATGCCTCATG CTGACTTGCA \*82  
GAGTTCATGG AATGTAAC TAATCATCCTT TATCCCTGAG GTCACCTGGA \*132  
ATCAGATTAT TAAGGGAATA AGCCATGACG TCAATAGCAG CCCAGGGCAC \*182  
TTCAGAGTAG AGGGCTTGGG AAGATCTTTT AAAAAGGCAG TAGGCCCGGT \*232  
GTGGTGGCTC ACGCCTATAA TCCCAGCACT TTGGGAGGCT GAAGTGGGTG \*282  
GATCACCAGA GGTGAGGAGT TCGAGACCAG CCCAGCCAAC ATGGCAAAAC \*332  
CCCATCTCTA CTA AAAAATAC AAAAATGAGC TAGGCATGGT GGCACACGCC \*382  
TGTAATCCCA GCTACACCTG AGGCTGAGGC AGGAGAAATG CTTGAACCGG \*432

GGAGACGGAG GTTGCAGTGA GCCGAGTTTG GGCCACTGCA CTCTAGCCTG \*482  
GCAACAGAGC AAGACTCCGT CTCAAAAAA GGGCAATAAA TGCCCTCTCT \*532  
GAATGTTTGA ACTGCCAAGA AAAGGCATGG AGACAGCGAA CTAGAAGAAA \*582  
GGGCAAGAAG GAAATAGCCA CCGTCTACAG ATGGCTTAGT TAAGTCATCC \*632  
ACAGCCCAAG GGCGGGGCTA TGCCTTGTCT GGGGACCCTG TAGAGTCACT \*682  
GACCCTGGAG CGGCTCTCCT GAGAGGTGCT GCAGGCAAAG TGAGACTGAC \*732  
ACCTCACTGA GGAAGGGAGA CATATTCTTG GAGAACTTTC CATCTGCTTG \*782  
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TTATTTTACA GAGCTTGAAA ACTCACTTCA ATGAACAAAAG GGATTCTCCA \*882  
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TGGAGGAACA GCTCCCTAGT GGCTTCCTCC GTCTGCAATG TCCCTTGACAC \*1082  
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CATGCTTTCC CACCAGCATG TCAC'TCCCAG ACCACCTCCC TGCCCTGTCC \*1232  
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TCTGGTGTTC TGAGTACTTG ATTTCAGGTC AATAACGGTC CCCCCTCACT \*1832

CCACACTGGC ACGTTTGTGA GAAGAAATGA CATTGCTA GGAAGTGACC \*1882  
GAGTCTAGGA ATGCTTTTAT TCAAGACACC AAATCCAAA CTCTAAATG \*1932  
TTGGAATTTT CAAAAATTGT GTTTAGATTT TATGAAAAAC TCTTCTACTT \*1982  
TCATCTATTC TTTCCCTAGA GGCAAACATT TCTTAAAATG TTTCATTTTC \*2032  
ATTAAAAATG AAAGCCAAAT TTATATGCCA CCGATTGCAG GACACAAGCA \*2082  
CAGTTTTAAG AGTTGTATGA ACATGGAGAG GACTTTTGGT TTTTATATTT \*2132  
CTCGTATTTA ATATGGGTGA ACACCAACTT TTATTTGGAA TAATAATTTT \*2182  
CCTCCTAAAC AAAAACACAT TGAGTTTAAAG TCTCTGACTC TTGCCTTTCC \*2232  
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CTGGAGCTGC TGTTCACAAC GACAGGGCCT AGCTTTCATT TGACACACAG \*2332  
ACTACAGCCA GAAGCCCATG GAGCAGGGAT GTCACGTCTT GAAAAGCCTA \*2382  
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TTAATGCCTT CCACATTAAT TAGATTTTCT TGCAGTTTTT TTATGGCATT \*2782  
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TATTTAATTA CCGGTTGTTA AAACGGTTT AGCACAATTT ATATTTTCCC \*2882  
TCTCTTGCC TCTTATTTG CAATAAAAGG TATTGAGCCA TTTTTTAAAT \*2932  
GACATTTTGT ATAAATTATG TTTGTACTAG TTGATGAAGG AGTTTTTTTT \*2982  
AACCTGTTTA TATAATTTG CAGCAGAAGC CAAATTTTTT GTATATTTAA \*3032  
GCACCAAATT CATGTACAGC ATGCATCACG GATCAATAGA CTGTACTTAT \*3082  
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