



*IL1RN* (NM\_173842.3) - cDNA - 2025-04-02

AGTCCACTGC CTTGCTGCAG TCACAGA**ATG** GAAATCTGCA GAGGCCTCCG 23  
CAGTCACCTA ATCACTCTCC TCCTCTTCCT GTTCCATT**CA** GAGACGATCT 73 [S21\\*](#)  
GCC**G**ACCC**TC** TGGGAGAAAA TCCAGCAAGA TGCAAGCCTT CAGAATCTGG 123 [R26\\*](#)  
GATGTTAACC **C** AGAAGAC**C**TT CTATCTGAGG AA**CAACCAAC** TAGTTGC**TGG** 173 [Q45\\*](#) [T47Tfs\\*4](#) [N52KfsX25](#) [N18Kfs\\*4](#) [Q54X](#) [A60A](#)  
ATACTTGCAA GGACCAAATG TCAATTTAGA AGAAAAGAT**A** GATGTGGTAC 223 [p.Asp72\\_Ile76del](#)  
CCATT**G**AGCC TCATGCTCTG TTCTTGGGAA TCCATGGAGG GAAGATGT**GC** 273 [E77X](#) [C91F](#)  
CTGTCTGTG TCAAGTCTGG TGATGAGACC AGACTCCAGC TGGAGGCAGT 323  
TAACATCACT GACCTGAGCG AGAACAGAAA G**C**AGGACAAG CGCTTC**G**CCT 373 [Q119X](#) [A124T](#)  
TCATCCGCTC AGACAG**TGGC** CC**C**ACCACCA GTTTTGAGTC TGCC**G**CCTGC 423 [S130S](#) [c.396delC](#) [A106T](#)  
CCCCGTTGGT TCCTCTGCAC AGCGATGGAA GCTGACCAGC CCGTCAGCCT 473  
CACCAATATG CCTGACGAAG GCGTCATGGT CACCAAATTC TACTTCCAGG 523  
AGGACGAG**TA** **G**TACTGCCCA GGCCTGCC**CTG** TTCCCAT**TCT** TGCATGGCAA \*39  
GGACTGCAGG GACTGCCAGT CCCCCTGCC CAGGGCTCCC GGCTATGGGG \*89  
GCACTGAGGA CCAGCCATTG AGGGGTGGAC CCTCAGAAGG CGTCACAA**CA** \*139 [c.\\*138C>G](#)  
ACCTGGTCAC AGGACTCTGC CT**C**CTCTTCA ACTGACCAGC CTCCATGCTG \*189 [c.\\*162C>T](#)  
CCTCCAGAAT GGTCTTTCTA ATGTGTGAAT CAGAGCACAG CAGCCCCTGC \*239  
ACAAAGCCCT TCCATGTCGC CTCTGCATTC AGGATCAAAC CCCGACCACC \*289  
TGCCCAACCT GCTCTCCTCT TGCCACTGCC TCTTCCTCCC TCATTCCACC \*339  
TTCCCATGCC CTGGATCCAT CAGGCCACTT GATGACCCCC AACCAAGTGG \*389  
CTCCACACC CTGTTTTACA AAAAAGAAAA GACCAGTCCA TGAGGGAGGT \*439  
TTTTAAGGGT TTGTGGAAAA TGAAAATTAG GATTT**CATGA** TTTTTTTTTT \*489  
TCAGTCCCCG TGAAGGAGAG CCCTTCATTT GGAGATTATG TTCTTTCGGG \*539

GAGAGGCTGA GGACTTAAAA TATTCCTGCA TTTGTGAAAT GATGGTGAAA \*589  
GTAAGTGGTA GCTTTTCCCT TCTTTTCTT CTTTTTTTGT GATGTCCCAA \*639  
CTTGTAAGAAA TTAAAAGTTA TGGTACTATG TTAGCCCCAT AATTTTTTTT \*689  
TTCCTTTTAA AACACTTCCA TAATCTGGAC TCCTCTGTCC AGGCACTGCT \*739  
GCCCAGCCTC CAAGCTCCAT CTCCACTCCA GATTTTTTAC AGCTGCCTGC \*789  
AGTACTTTAC CTCCTATCAG AAGTTTCTCA GCTCCCAAGG CTCTGAGCAA \*839  
ATGTGGCTCC TGGGGTCTT TCTTCTCT GCTGAAGGAA TAAATTGCTC \*889  
CTTGACATTG TAGAGCTTCT GGCAC TTGGA GACTTGTATG AAAGATGGCT \*939  
GTGCCTCTGC CTGTCTCCCC CACCGGGCTG GGAGCTCTGC AGAGCAGGAA \*989  
ACATGACTCG TATATGTCTC AGGTCCCTGC AGGGCCAAGC ACCTAGCCTC \*1039  
GCTCTTGGCA GGTACTCAGC GAATGAATGC TGTATATGTT GGGTGCAAAG \*1089  
TTCCCTACTT CCTGTGACTT CAGCTCTGTT TTACAATAAA ATCTTGAAAA \*1139  
TGCC

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