



IL1RN (NM_173842.3) - cDNA - 2024-11-24

AGTCCACTGC CTTGCTGCAG TCACAGA**ATG** GAAATCTGCA GAGGCCTCCG 23
CAGTCACCTA ATCACTCTCC TCCTCTTCCT GTTCCATT**CA** GAGACGATCT 73 [S21*](#)
GCC**G**ACCCTC 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**G**C 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 GGCCTGCCCTG TTCCCATCTT 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 GATTTTCATGA 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 TGGGGTTCT TTCTTCTCT 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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