



IL1RN (NM_173842.3) - cDNA - 2024-05-19

AGTCCACTGC CTTGCTGCAG TCACAGAAATG GAAATCTGCA GAGGCCTCCG 23
CAGTCACCTA ATCACTCTCC TCCTCTTCCT GTTCCATTCA GAGACGATCT 73 [S21*](#)
GCCGACCCTC TGGGAGAAAA TCCAGCAAGA TGCAAGCCTT CAGAATCTGG 123 [R26*](#)
GATGTTAACC AGAAGACCTT CTATCTGAGG AACAAACAAC TAGTTGCTGG 173 [Q45*](#) [T47Tfs*4](#) [N52KfsX25](#) [N18Kfs*4](#) [Q54X](#) [A60A](#)
ATACTTGCAA GGACCAAATG TCAATTTAGA AGAAAAGATA GATGTGGTAC 223 [p.Asp72_Ile76del](#)
CCATTGAGCC TCATGCTCTG TTCTTGGGAA TCCATGGAGG GAAGATGTGC 273 [E77X](#) [C91F](#)
CTGTCCTGTG TCAAGTCTGG TGATGAGACC AGACTCCAGC TGGAGGCAGT 323
TAACATCACT GACCTGAGCG AGAACAGAAA GCAGGACAAG CGCTTCGCCT 373 [Q119X](#) [A124I](#)
TCATCCGCTC AGACAGTGGC CCACCACCA GTTTTGAGTC TGCCGCCTGC 423 [S130S](#) [c.396delC](#) [A106I](#)
CCCGGTTGGT TCCTCTGCAC AGCGATGGAA GCTGACCAGC CCGTCAGCCT 473
CACCAATATG CCTGACGAAG GCGTCATGGT CACCAAATTC TACTTCCAGG 523
AGGACGAGTA GACTGCCCA GGCCTGCCTG TTCCCATTCT TGCATGGCAA *39
GGACTGCAGG GACTGCCAGT CCCCTGCCC CAGGGCTCCC GGCTATGGGG *89
GCACTGAGGA CCAGCCATTG AGGGGTGGAC CCTCAGAAGG CGTCACAACA *139 [c.*138C>G](#)
ACCTGGTCAC AGGACTCTGC CTCTCTTCA 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
CTCCCACACC 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
CTTGTA AAAA TTAAGTTA TGGTACTATG TTAGCCCCAT AATTTTTTTT *689
TTCCTTTTAA AACACTTCCA TAATCTGGAC TCCTCTGTCC AGGCACTGCT *739
GCCCAGCCTC CAAGCTCCAT CTCCACTCCA GATTTTTTAC AGCTGCCTGC *789
AGTACTTTAC CTCCTATCAG AAGTTTCTCA GCTCCCAAGG CTCTGAGCAA *839
ATGTGGCTCC TGGGGGTTCT TTCTTCCTCT GCTGAAGGAA TAAATTGCTC *889
CTTGACATTG TAGAGCTTCT GGCAC TTGGA GACTTGTATG AAAGATGGCT *939
GTGCCTCTGC CTGTCTCCCC CACCGGGCTG GGAGCTCTGC AGAGCAGGAA *989
ACATGACTCG TATATGTCTC AGGTCCCTGC AGGGCCAAGC ACCTAGCCTC *1039
GCTCTTGGA GGTACTCAGC GAATGAATGC TGTATATGTT GGGTGCAAAG *1089
TTCCCTACTT CCTGTGACTT CAGCTCTGTT TTACAATAAA ATCTTGAAAA *1139
TGCC

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