

IL1RN (NM_173842.3) - cDNA - 2026-11-24

AGTCCACTGC CTTGCTGCAG TCACAGAATG GAAATCTGCA GAGGCCTCCG 23

CAGTCACCTA ATCACTCTCC TCCTCTTCCT GTTCCATTCA GAGACGATCT 73 S21*

GCCGACCCTC TGGGAGAAAA TCCAGCAAGA TGCAAGCCTT CAGAATCTGG 123 R26*

GATGTTAACC AGAAGACCTT CTATCTGAGG AACAACCAAC 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 A124T

TCATCCGCTC AGACAGTGGC CCCACCACCA GTTTTGAGTC TGCCGCCTGC 423 S130S c.396delC A106T

CCCGGTTGGT TCCTCTGCAC AGCGATGGAA GCTGACCAGC CCGTCAGCCT 473

CACCAATATG CCTGACGAAG GCGTCATGGT CACCAAATTC TACTTCCAGG 523

AGGACGAGTA GTACTGCCCA GGCCTGCCTG TTCCCATTCT TGCATGGCAA *39

GGACTGCAGG GACTGCCAGT CCCCCTGCCC CAGGGCTCCC GGCTATGGGG *89

GCACTGAGGA CCAGCCATTG AGGGGTGGAC CCTCAGAAGG CGTCACAACA *139 c.*138C>G

ACCTGGTCAC AGGACTCTGC CTCCTCTTCA 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 GATTTCATGA TTTTTTTTTT *489

TCAGTCCCCG TGAAGGAGAG CCCTTCATTT GGAGATTATG TTCTTTCGGG *539

GAGAGGCTGA GGACTTAAAA TATTCCTGCA TTTGTGAAAT GATGGTGAAA *589

GTAAGTGGTA GCTTTTCCCT TCTTTTTCTT CTTTTTTTGT GATGTCCCAA *639

CTTGTAAAAA TTAAAAGTTA 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 GGCACTTGGA 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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