



IL1RN (NM_173842.3) - cDNA - 2025-07-16

AGTCCACTGC CTTGCTGCAG TCACAGA**ATG** GAAATCTGCA GAGGCCTCCG 23
CAGTCACCTA ATCACTCTCC TCCTCTTCCT GTTCCATT**CA** GAGACGATCT 73 S21*
GC**C**GACCCTC TGGGAGAAAA TCCAGCAAGA TGCAAGCCTT CAGAATCTGG 123 R26*
GATGTTAAC**C** AGAACAG**CTT** CTATCTGAGG AA**CAAC** AAC TAGTTG**CTGG** 173 Q45* T47Tfs*4 N52KfsX25 N18Kfs*4 Q54X A60A
ATACTTGCAA GGACCAAATG TCAATTTAGA AGAAAAG**ATG** GATGTGGTAC 223 p.Asp72_Ile76del
CCATT**GAGCC** TCATGCTCTG TTCTTGGGAA TCCATGGAGG GAAGAT**GTGC** 273 E77X C91F
CTGTCCTGTG TCAAGTCTGG TGATGAGACC AGACTCCAGC TGGAGGCAGT 323
TAACATCACT GACCTGAGCG AGAACAGAAA GCAGGACAAG CGCTTC**GCCT** 373 Q119X A124T
TCATCCGCTC AGACAG**TGGC** CC**CACCACCA** GTTTGAGTC TGCC**GCCTGC** 423 S130S c.396delC A106T
CCCGGTTGGT TCCTCTGCAC AGCGATGGAA GCTGACCAGC CCGTCAGCCT 473
CACCAATATG CCTGACGAAG GCGTCATGGT CACCAAATTG TACTTCCAGG 523
AGGACGAG**TA** GTACTGCCCA GGCCTGCCTG TTCCCATTCT TGCATGGCAA *39
GGACTGCAGG GACTGCCAGT CCCCTGCC CAGGGCTCCC GGCTATGGGG *89
GCACTGAGGA CCAGCCATTG AGGGGTGGAC CCTCAGAAGG CGTCACAA**CA** *139 c.*138C>G
ACCTGGTCAC AGGACTCTGC CT**C**CTCTTCA ACTGACCAGC CTCCATG**CTG** *189 c.*162C>T
CCTCCAGAAT GGTCTTCTA ATGTGTGAAT CAGAGCACAG CAGCCCC**TC** *239
ACAAAGCCCT TCCATGTCGC CTCTGCATTG AGGATCAAAC CCCGACCACC *289
TGCCCCAACCT GCTCTCCTCT TGCCACTGCC TCTTCCCTCC TCATTCCACC *339
TTCCCATGCC CTGGATCCAT CAGGCCACTT GATGACCCCC AACCAAGTGG *389
CTCCCCACACC CTGTTTACA AAAAAGAAAA GACCAGTCCA TGAGGGAGGT *439
TTTTAAGGGT TTGTGGAAAA TGAAAATTAG GATTCATGA TTTTTTTTTT *489
TCAGTCCCCG TGAAGGAGAG CCCTTCATTG GGAGATTATG TTCTTCGGG *539

GAGAGGCTGA GGACTTAAAA TATTCCTGCA TTTGTGAAAT GATGGTGAAA *589
GTAAGTGGTA GCTTTCCCT TCTTTTCTT CTTTTTTGT GATGTCCCAA *639
CTTGTAAAAA TTAAAAGTTA TGGTACTATG TTAGCCCCAT AATTTTTTTT *689
TCCCTTTAA AACACTTCCA TAATCTGGAC TCCTCTGTCC AGGCACTGCT *739
GCCCAGCCTC CAAGCTCCAT CTCCACTCCA GATTTTTAC 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
GCTCTGGCA GGTACTCAGC GAATGAATGC TGTATATGTT GGGTGCAAAG *1089
TTCCCTACTT CCTGTGACTT CAGCTCTGTT TTACAATAAA ATCTTGAAAA *1139

TGCC

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