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

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AGTCCACTGC CTTGCTGCAG TCACAGAATG GAAATCTGCA GAGGCCTCCG 23
                             Met GluIleCysA rgGlyLeuAr 8

CAGTCACCTA ATCACTCTCC TCCTCTTCCT GTTCCATTCA GAGACGATCT 73   S21*
gSerHisLeu IleThrLeuL euLeuPheLe uPheHisSer GluThrIleC 25

GCCGACCCTC TGGGAGAAAA TCCAGCAAGA TGCAAGCCTT CAGAATCTGG 123  R26*
ysArgProSe rGlyArgLys SerSerLysM etGlnAlaPh eArgIleTrp 41

GATGTTAACC AGAAGACCTT CTATCTGAGG AACAACCAAC TAGTTGCTGG 173  Q45* T47Tfs*4 N52KfsX25 N18Kfs*4 Q54X A60A
AspValAsnG lnLysThrPh eTyrLeuArg AsnAsnGlnL euValAlaGl 58

ATACTTGCAA GGACCAAATG TCAATTTAGA AGAAAAGATA GATGTGGTAC 223  p.Asp72_Ile76del
yTyrLeuGln GlyProAsnV alAsnLeuGl uGluLysIle AspValValP 75

CCATTGAGCC TCATGCTCTG TTCTTGGGAA TCCATGGAGG GAAGATGTGC 273  E77X C91F
roIleGluPr oHisAlaLeu PheLeuGlyI leHisGlyGl yLysMetCys 91

CTGTCCTGTG TCAAGTCTGG TGATGAGACC AGACTCCAGC TGGAGGCAGT 323
LeuSerCysV alLysSerGl yAspGluThr ArgLeuGlnL euGluAlaVa 108

TAACATCACT GACCTGAGCG AGAACAGAAA GCAGGACAAG CGCTTCGCCT 373  Q119X A124T
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lAsnIleThr AspLeuSerG luAsnArgLy sGlnAspLys ArgPheAlaP 125

TCATCCGCTC AGACAGTGGC CCCACCACCA GTTTTGAGTC TGCCGCCTGC 423  S130S  c.396delC  A106T
heIleArgSe rAspSerGly ProThrThrS erPheGluSe rAlaAlaCys 141

CCCGGTTGGT TCCTCTGCAC AGCGATGGAA GCTGACCAGC CCGTCAGCCT 473
ProGlyTrpP heLeuCysTh rAlaMetGlu AlaAspGlnP roValSerLe 158

CACCAATATG CCTGACGAAG GCGTCATGGT CACCAAATTC TACTTCCAGG 523
uThrAsnMet ProAspGluG lyValMetVa lThrLysPhe TyrPheGlnG 175

AGGACGAGTA GTACTGCCCA GGCCTGCCTG TTCCCATTCT TGCATGGCAA *39
luAspGluSt op

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
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TTCCTTTTAA AACACTTCCA TAATCTGGAC TCCTCTGTCC AGGCACTGCT *739
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ACATGACTCG TATATGTCTC AGGTCCCTGC AGGGCCAAGC ACCTAGCCTC *1039
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TGCC
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