



## IL36RN (NM\_173170.1) - cDNA + Protein - 2024-05-19

GGAGAGTCCC ACCTCTAAC A TCTCCTGTAG GCCTGGCAAT GGCAGGCAGG -84

AAAGACAGAG GAAGGAAGGA GGGAGAAGGG AAGGAGTGAA GGAAGGGAGTG -34

AAAAAGGGGA GTCTACACCC TGTGGAGCTC AAGATGGTCC TGAGTGGGGC 17 V2F

MetValL euSerGlyAl 6

GCTGTGCTTC CGAATGAAGG ACTCGGCATT GAAGGTGCTT TATTGCATA 67 R10X S14X L21P

aLeuCysPhe ArgMetLysA spSerAlaLe uLysValLeu TyrLeuHisA 23

ATAACCAGCT TCTAGCTGGA GGGCTGCATG CAGGGAGGT CATTAAGGT 117 L27P H32R K35R

snAsnGlnLe uLeuAlaGly GlyLeuHisA laGlyLysVa 1IleLysGly 39

GAAGAGATCA GCGTGGTCCC CAATCGGTGG CTGGATGCCA GCCTGTCCCC 167 I42N V44M P46S N47S R48W G48Q

GluGluIleS erValValPr oAsnArgTrp LeuAspAlaS erLeuSerPr 56

CGTCATCCTG GGTGTCCAGG GTGGAAGCCA GTGCCTGTCA TGTGGGGTGG 217 V57I C67F

ovalIleLeu GlyValGlnG lyGlySerG1 nCysLeuSer CysGlyValG 73

GGCAGGAGC GACTCTAAC A CTAGAGCAG TGAACATCAT GGAGCTCTAT 267 P76L P82L M86I

lyGlnGluPr oThrLeuThr LeuGluProV alAsnIleMe tGluLeuTyr 89

CTTGGTGCCA AGGAATCCAA GAGCTTCACC TTCTACCGGC GGGACATGGG 317 E94X T99\_F100del R102W R102Q R103Q

LeuGlyAlaL ysGluSerLy sSerPheThr PheTyrArgA rgAspMetG1 106

GCTCACCTCC AGCTTCGAGT CGGCTGCCTA CCCGGGCTGG TTCCTGTGCA 367 E112K S113L S113X  
yLeuThrSer SerPheGluS erAlaAlaTy rProGlyTrp PheLeuCysT 123

CGGTGCCTGA AGCCGATCAG CCTGTCAGAC TCACCCAGCT TCCCAGAAT 417 T123R T123M  
hrValProGl uAlaAspGln ProValArgL euThrGlnLe uProGluAsn 139

GGTGGCTGGA ATGCCCCCCAT CACAGACTTC TACTTCCAGC AGTGTGACTA 467 G141Mfs\*29 I146V  
GlyGlyTrpA snAlaProIl eThrAspPhe TyrPheGlnG lnCysAspSt 156

GGGCAACGTG CCCCCCAGAA CTCCCTGGGC AGAGCCAGCT CGGGGTGAGGG \*49 \*43G>A  
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CCCCACGTCT GACTTAGTGG GCACCTGACC ACTTTGTCTT CTGGTTCCCA \*149  
GTTTGGATAA ATTCTGAGAT TTGGAGCTCA GTCCACGGTC CTCCCCCACT \*199  
GGATGGTGC ACTGCTGTGG AATCTTGAA AAACCATGTG GGGTAAACTG \*249  
GGAATAACAT GAAAAGATT CTGTGGAGGT GGGGTGGGGG AGTGGTGGGA \*299  
ATCATTCCCTG CTTAACGGTA ACTGACCACT GTTACCCCTGA GCCCCGGCAGG \*349  
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GACCTGTCAC TCACCACTAT GCAGGAGAGG GAGGTGGTCA TAGAGTCAGG \*449  
GATCTATGGC CCTTGGCCCA GCCCCACCTC CTTCCCTTTA ATCCTGCCAC \*499  
TGTCAATATGC TACCTTCCT ATCTCTTCCC TCATCATCTT GTTGTGGCA \*549  
TGAGGAGGTG CTGATGTCAG AAGAAATGGC TCGAGCTCAG AAGATAAAAG \*599  
ATAAGTAGGG TATGCTGATC CTCTTTAAA AACCCAAGAT ACAATCAAA \*649  
TCCCAGATGC TGGTCTCTAT TCCCATGAAA AAGTGTCTCAT GACATATTGA \*699  
GAAGACCTAC TTACAAAGTG GCATATATTG CAATTTTATTT TAATTAAAAG \*749  
ATACCTATTT ATATATTCT TTATAGAAAA AAGTGTGGAA GAGTTTACTT \*799  
CAATTGTAGC AATGTCAGGG TGGTGGCAGT ATAGGTGATT TTTCTTTAA \*849  
TTCTGTTAAT TTACCTGTAT TTCTAATTT TTCTACAAATG AAGATGAATT \*899

CCTTGTATAA AAATAAGAAA AGAAATTAAT CTTGAGGTAA GCAGAGTAGA \*949  
CATCATCTCT GATTGTCCTC AGCCTCCACT TCCCCAGAGT AAATTCAAAT \*999  
TGAATCGAGC TCTGCTGCTC TGGTTGGTTG TAGTAGTGAT CAGGAAACAG \*1049  
ATCTCAGCAA AGCCACTGAG GAGGAGGCTG TGCTGAGTT GTGTGGCTGG \*1099  
AATCTCTGGG TAAGGAACCT AAAGAACAAA AATCATCTGG TAATTCTTC \*1149  
CTAGAAGGAT CACAGCCCCT GGGATTCCAA GGCATTGGAT CCAGTCTCTA \*1199  
AGAAGGCTGC TGTACTGGTT GAATTGTGTC CCCCTCAAAT TCACATCCTT \*1249  
CTTCCAATCT CAGTCGTGA GTTATTGG AGATAAGGTC TCTGCAGATG \*1299  
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GCGGGGAAGA CTATGTAAAG ATGAAGGCAG AGATCGGAGT TTTGCAGCCA \*1449  
CAAGCTAAGA AACACCAAGG ATTGTGGCAA CCATCAGAAG CTTGGAAGAG \*1499  
GCAAAGAAGA ATTCTCCCT AGAGGCTTTA GAGGGATAAC GGCTCTGCTG \*1549  
AACACCTTAAT CTCAGACTTC CAGCCTCCTG AACGAAGAAA GAATAAATTG \*1599  
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GTGTCCCCCTC CCACAATGTA CCAAAGTTGT CTTTGTGACC AATAGAATAT \*1749  
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CAGCTTCTAC TTGAGCCCTC TCTCTCTGCC ACCCACCGCC CCCAATCTAT \*1849  
CTTGGCTCAC TCGCTCTGGG GGAAGCTAGC TGCCATGCTA TGAGCAGGCC \*1899  
TATAAAGAGA CTTACGTGGT AAAAAATGAA GTCTCCTGCC CACAGCCACA \*1949  
TTAGTGAACC TAGAAGCAGA GACTCTGTGA GATAATCGAT GTTTGTTGTT \*1999  
TTAACGTTGCT CAGTTTGGT CTAACCTGTT ATGCAGCAAT AGATAAATAA \*2049  
TATGCAGAGA AAGAGAAA

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