



IL36RN (NM_012275.3) - cDNA + Protein - 2026-06-18

AGGCAGACTC CACAGCTCCC GCCAGGAGAA AGGAACATTC TGAGGGGAGT -22
CTACACCCTG TGGAGCTCAA **GATGG**TCCTG AGTGGGGCGC TGTGCTTC**CG** 29 **V2F R10X**
MetValLeu SerGlyAlaL euCysPheAr 10

AATGAAGGAC **T**CGGCATTGA AGGTGCTTTA **TC**TGCATAAT AACCAGCTTC 79 **S14X L21P**
gMetLysAsp SerAlaLeuL ysValLeuTy rLeuHisAsn AsnGlnLeuL 27

TAGCTGGAGG GCTGC**A**TGCA GGG**A**AGGTCA **TTAA****A**GGTGA AGAG**A**TCAGC 129 **L27P H32R K35R G39C I42N**
euAlaGlyGl yLeuHisAla GlyLysValI leLysGlyGl uGluIleSer 43

GTGGT**C**CCA **A**TCGGTGGCT GGATGCCAGC CTGT**C**CCC**G** TCATCCTGGG 179 **V44M P46S N47S R48W G48Q V57I**
ValValProA snArgTrpLe uAspAlaSer LeuSerProV alIleLeuGl 60

TGTCCAGGGT GGAAGCCAGT **G**CCGTGCATG TGGGGTGGGG CAGGAG**C**GA 229 **C67F P76L**
yValGlnGly GlySerGlnC ysLeuSerCy sGlyValGly GlnGluProT 77

CTCTAACACT AGAG**C**AGTG AACATCAT**G** AGCTCTATCT TGGTGCCAAG 279 **P82L M86I**
hrLeuThrLe uGluProVal AsnIleMetG luLeuTyrLe uGlyAlaLys 93

GAATCCAAGA GCTTC**A**CCTT CTAC**C**GG**C**G GACATGGGGC TCACCTCCAG 329 **E94X T99_F100del R102W R102Q R103Q**
GluSerLysS erPheThrPh eTyrArgArg AspMetGlyL euThrSerSe 110

CTTCGAGTCG GCTGCCTACC CGGGCTGGTT CCTGTGCACG GTGCCTGAAG 379 E112K S113L S113X T123R T123M

rPheGluSer AlaAlaTyrP roGlyTrpPh eLeuCysThr ValProGluA 127

CCGATCAGCC TGTCAGACTC ACCCAGCTTC CCGAGAATGG TGGCTGGAAT 429 G141Mfs*29

laAspGlnPr oValArgLeu ThrGlnLeuP roGluAsnGl yGlyTrpAsn 143

GCCCCATCA CAGACTTCTA CTTCCAGCAG TGTGACTAGG GCAACGTGCC *11 I146V

AlaProIleT hrAspPheTy rPheGlnGln CysAspStop

CCCCAGAACT CCCTGGGCAG AGCCAGCTCG GGTGAGGGGT GAGTGGAGGA *61 *43G>A

GACCCATGGC GGACAATCAC TCTCTCTGCT CTCAGGACCC CCACGTCTGA *111

CTTAGTGGGC ACCTGACCAC TTTGTCTTCT GGTTCCCAGT TTGGATAAAT *161

TCTGAGATTT GGAGCTCAGT CCACGGTCCT CCCCCTGG ATGGTGCTAC *211

TGCTGTGGAA TCTTGTAATA ACCATGTGGG GTAAACTGGG AATAACATGA *261

AAAGATTCTT GTGGAGGTGG GGTGGGGGAG TGGTGGGAAT CATTCCTGCT *311

TAATGGTAAC TGACCAGTGT TACCCTGAGC CCCGCAGGCC AACCCATCCC *361

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ACCACTATGC AGGAGAGGGA GGTGGTCATA GAGTCAGGGA TCTATGGCCC *461

TTGGCCCAGC CCCACCTCCT TCCCTTTAAT CCTGCCACTG TCATATGCTA *511

CCTTTCCTAT CTCTTCCCTC ATCATCTTGT TGTGGGCATG AGGAGGTGCT *561

GATGTCAGAA GAAATGGCTC GAGCTCAGAA GATAAAAGAT AAGTAGGGTA *611

TGCTGATCCT CTTTTAAAAA CCCAAGATAC AATCAAAATC CCAGATGCTG *661

GTCTCTATTC CCATGAAAAA GTGCTCATGA CATATTGAGA AGACCTACTT *711

ACAAAGTGGC ATATATTGCA ATTTATTTTA ATTAAAAGAT ACCTATTTAT *761

ATATTTCTTT ATAGAAAAA GTCTGGAAGA GTTTACTTCA ATTGTAGCAA *811

TGTCAGGGTG GTGGCAGTAT AGGTGATTTT TCTTTTAATT CTGTTAATTT *861

ACCTGTATTT CCTAATTTTT CTACAATGAA GATGAATTCC TTGTATAAAA *911

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CAGCCCCCTGG GATTCCAAGG CATTGGATCC AGTCTCTAAG AAGGCTGCTG *1211
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GAGCCCTCTC TCTCTGCCAC CCACCGCCCC CAATCTATCT TGGCTCACTC *1861
GCTCTGGGGG AAGCTAGCTG CCATGCTATG AGCAGGCCTA TAAAGAGACT *1911
TACGTGGTAA AAAATGAAGT CTCTGCCCCA CAGCCACATT AGTGAACCTA *1961
GAAGCAGAGA CTCTGTGAGA TAATCGATGT TTGTTGTTTT AAGTTGCTCA *2011
GTTTTGGTCT AACTTGTAT GCAGCAATAG ATAAATAATA TGCAGAGAAA *2061

GAGAAA

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