IL36RN (NM_012275.3) - cDNA + Protein - 2026-11-14

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AGGCAGACTC CACAGCTCCC GCCAGGAGAA AGGAACATTC TGAGGGGAGT -22
CTACACCCTG TGGAGCTCAA GATGGTCCTG AGTGGGGCGC TGTGCTTCCG 29    V2F R10X
                      MetValLeu SerGlyAlaL euCysPheAr 10

AATGAAGGAC TCGGCATTGA AGGTGCTTTA TCTGCATAAT AACCAGCTTC 79    S14X L21P
gMetLysAsp SerAlaLeuL ysValLeuTy rLeuHisAsn AsnGlnLeuL 27

TAGCTGGAGG GCTGCATGCA GGGAAGGTCA TTAAAGGTGA AGAGATCAGC 129   L27P H32R K35R G39C I42N
euAlaGlyGl yLeuHisAla GlyLysValI leLysGlyGl uGluIleSer 43

GTGGTCCCCA ATCGGTGGCT GGATGCCAGC CTGTCCCCCG TCATCCTGGG 179   V44M P46S N47S R48W G48Q V57I
ValValProA snArgTrpLe uAspAlaSer LeuSerProV alIleLeuGl 60

TGTCCAGGGT GGAAGCCAGT GCCTGTCATG TGGGGTGGGG CAGGAGCCGA 229   C67F P76L
yValGlnGly GlySerGlnC ysLeuSerCy sGlyValGly GlnGluProT 77

CTCTAACACT AGAGCCAGTG AACATCATGG AGCTCTATCT TGGTGCCAAG 279   P82L M86I
hrLeuThrLe uGluProVal AsnIleMetG luLeuTyrLe uGlyAlaLys 93

GAATCCAAGA GCTTCACCTT CTACCGGCGG GACATGGGGC TCACCTCCAG 329   E94X T99_F100del R102W R102Q R103Q
GluSerLysS erPheThrPh eTyrArgArg AspMetGlyL euThrSerSe 110

CTTCGAGTCG GCTGCCTACC CGGGCTGGTT CCTGTGCACG GTGCCTGAAG 379   E112K S113L S113X P117L T123R T123M
rPheGluSer AlaAlaTyrP roGlyTrpPh eLeuCysThr ValProGluA 127

CCGATCAGCC TGTCAGACTC ACCCAGCTTC CCGAGAATGG TGGCTGGAAT 429   G141Mfs*29
laAspGlnPr oValArgLeu ThrGlnLeuP roGluAsnGl yGlyTrpAsn 143

GCCCCCATCA 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 CCCCCACTGG ATGGTGCTAC *211
TGCTGTGGAA TCTTGTAAAA ACCATGTGGG GTAAACTGGG AATAACATGA *261
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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 ATAGAAAAAA GTCTGGAAGA GTTTACTTCA ATTGTAGCAA *811

TGTCAGGGTG GTGGCAGTAT AGGTGATTTT TCTTTTAATT CTGTTAATTT *861

ACCTGTATTT CCTAATTTTT CTACAATGAA GATGAATTCC TTGTATAAAA *911

ATAAGAAAAG AAATTAATCT TGAGGTAAGC AGAGTAGACA TCATCTCTGA *961

TTGTCCTCAG CCTCCACTTC CCCAGAGTAA ATTCAAATTG AATCGAGCTC *1011

TGCTGCTCTG GTTGGTTGTA GTAGTGATCA GGAAACAGAT CTCAGCAAAG *1061

CCACTGAGGA GGAGGCTGTG CTGAGTTTGT GTGGCTGGAA TCTCTGGGTA *1111

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CAGCCCCTGG GATTCCAAGG CATTGGATCC AGTCTCTAAG AAGGCTGCTG *1211

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CACCAAGGAT TGTGGCAACC ATCAGAAGCT TGGAAGAGGC AAAGAAGAAT *1511

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CAGACTTCCA GCCTCCTGAA CGAAGAAAGA ATAAATTTCG GCTGTTTTAA *1611

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ACAATGTACC AAAGTTGTCT TTGTGACCAA TAGAATATGG CAGAAGTGAT *1761

GGCATGCCAC TTCCAAGATT AGGTTATAAA AGACACTGCA GCTTCTACTT *1811

GAGCCCTCTC TCTCTGCCAC CCACCGCCCC CAATCTATCT TGGCTCACTC *1861

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TACGTGGTAA AAAATGAAGT CTCCTGCCCA CAGCCACATT AGTGAACCTA *1961

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GTTTTGGTCT AACTTGTTAT GCAGCAATAG ATAAATAATA TGCAGAGAAA *2061

GAGAAA

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