

Infevers - IL36RN (NM_173170.1) - cDNA + Protein - 2019-08-22

GGAGAGTCCC	ACCTCTAACA	TCTCCTGTAG	GCCTGGCAAT	GGCAGGCAGG	-84	
AAAGACAGAG	GAAGGAAGGA	GGGAGAAGGG	AAGGAGTGAA	GGAAGGAGTG	-34	
AAAAAGGGGA	GTCTACACCC	TGTGGAGCTC	AAGATGGTCC	TGAGTGGGGC	17	<u>V2F</u>
			MetValL	euSerGlyAl	6	
GCTGTGCTTC	CGAATGAAGG	ACTCGGCATT	GAAGGTGCTT	TATCTGCATA	67	<u>R10X S14X L21P</u>
aLeuCysPhe	ArgMetLysA	spSerAlaLe	uLysValLeu	TyrLeuHisA	23	
ATAACCAGCT	TCTAGCTGGA	GGGCTGCATG	CAGGGAAGGT	CATTAAAGGT	117	<u>L27P H32R K35R</u>
snAsnGlnLe	uLeuAlaGly	GlyLeuHisA	laGlyLysVa	lIleLysGly	39	
GAAGAGATCA	GCGTGGTCC	CAATCGGTGG	CTGGATGCCA	GCCTGTCCCC	167	<u>I42N P46S N47S R48W</u>
GluGluIleS	erValValPr	oAsnArgTrp	LeuAspAlaS	erLeuSerPr	56	
CGTCATCCTG	GGTGTCCAGG	GTGGAAGCCA	GTGCCTGTCA	TGTGGGGTGG	217	<u>C67E</u>
oValIleLeu	GlyValGlnG	lyGlySerGl	nCysLeuSer	CysGlyValG	73	
GGCAGGAGCC	GACTCTAACA	CTAGAGCCAG	TGAACATCAT	GGAGCTCTAT	267	<u>P76L</u>
lyGlnGluPr	oThrLeuThr	LeuGluProV	alAsnIleMe	tGluLeuTyr	89	
CTTGGTGCCA	AGGAATCCAA	GAGCTTCACC	TTCTACCGGC	GGGACATGGG	317	<u>E94X R102W R102Q</u>
LeuGlyAlaL	ysGluSerLy	sSerPheThr	PheTyrArgA	rgAspMetGl	106	
GCTCACCTCC	AGCTTCGAGT	CGGCTGCCTA	CCCGGGCTGG	TTCCTGTGCA	367	<u>E112K S113L</u>
yLeuThrSer	SerPheGluS	erAlaAlaTy	rProGlyTrp	PheLeuCysT	123	
CGGTGCCTGA	AGCCGATCAG	CCTGTCAGAC	TCACCCAGCT	TCCCGAGAAT	417	<u>T123R T123M</u>
hrValProGl	uAlaAspGln	ProValArgL	euThrGlnLe	uProGluAsn	139	
GGTGGCTGGA	ATGCCCCCAT	CACAGACTTC	TACTTCCAGC	AGTGTGACTA	467	<u>G141Mfs*29 I146V</u>
GlyGlyTrpA	snAlaProIl	eThrAspPhe	TyrPheGlnG	lnCysAspSt	156	
GGGCAACGTG	CCCCCAGAA	CTCCCTGGGC	AGAGCCAGCT	CGGGTGAGGG	*49	<u>*43G>A</u>
GTGAGTGGAG	GAGACCCATG	GCGGACAATC	ACTCTCTCTG	CTCTCAGGAC	*99	
CCCCACGTCT	GACTTAGTGG	GCACCTGACC	ACTTTGTCTT	CTGGTTCCCA	*149	
GTTTGGATAA	ATTCTGAGAT	TTGGAGCTCA	GTCCACGGTC	CTCCCCCACT	*199	
GGATGGTGCT	ACTGCTGTGG	AATCTTGTA	AAACCATGTG	GGGTAAACTG	*249	

GGAATAACAT GAAAAGATTT CTGTGGAGGT GGGGTGGGG AGTGGTGGGA *299
ATCATTCCCTG CTTAATGGTA ACTGACCAGT GTTACCCTGA GCCCCG CAGG *349
CCAACCCATC CCCAGTTGAG CCTTATAGGG TCAGTAGCTC TCCACATGAA *399
GACCTGTAC TCACCACTAT GCAGGAGAGG GAGGTGGTCA TAGAGTCAGG *449
GATCTATGGC CCTTGGCCCA GCCCCACCTC CTTCCCTTTA ATCCTGCCAC *499
TGTCATATGC TACCTTTCT ATCTCTTCCC TCATCATCTT GTTGTGGGCA *549
TGAGGAGGTG CTGATGTCAG AAGAAATGGC TCGAGCTCAG AAGATAAAAAG *599
ATAAGTAGGG TATGCTGATC CTCTTTTAAA AACCCAAGAT ACAATCAAAA *649
TCCCAGATGC TGGTCTCTAT TCCCATGAAA AAGTGCTCAT GACATATTGA *699
GAAGACCTAC TTACAAAGTG GCATATATTG CAATTTATTT TAATTA AAAAG *749
ATACCTATTT ATATATTTCT TTATAGAAAA AAGTCTGGAA GAGTTTACTT *799
CAATTGTAGC AATGTCAGGG TGGTGGCAGT ATAGGTGATT TTTCTTTTAA *849
TTCTGTTAAT TTACCTGTAT TTCCTAATTT TTCTACAATG AAGATGAATT *899
CCTTGTATAA AAATAAGAAA AGAAATTAAT CTTGAGGTAA GCAGAGTAGA *949
CATCATCTCT GATTGTCCCTC AGCCTCCACT TCCCCAGAGT AAATTC AAAT *999
TGAATCGAGC TCTGCTGCTC TGGTTGGTTG TAGTAGTGAT CAGGAAACAG *1049
ATCTCAGCAA AGCCACTGAG GAGGAGGCTG TGCTGAGTTT GTGTGGCTGG *1099
AATCTCTGGG TAAGGAACTT AAAGAACAAA AATCATCTGG TAATTCTTTC *1149
CTAGAAGGAT CACAGCCCCT GGGATTCCA GGCATTGGAT CCAGTCTCTA *1199
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TAGTTAGTTA AGACAAGGTC ATGCTGGATG AAGGTAGACC TAAATTCAAT *1349
ATGACTGGTT TCCTTGTATG AAAAGGAGAG GACACAGAGA CAGAGGAGAT *1399
GCGGGGAAGA CTATGTAAAG ATGAAGGCAG AGATCGGAGT TTTGCAGCCA *1449
CAAGCTAAGA AACACCAAGG ATTGTGGCAA CCATCAGAAG CTTGGAAGAG *1499
GCAAAGAAGA ATTCTTCCCT AGAGGCTTTA GAGGGATAAC GGCTCTGCTG *1549
AAACCTTAAT CTCAGACTTC CAGCCTCCTG AACGAAGAAA GAATAAATTT *1599
CGGCTGTTTT AAGCCACCAA GGATAATTGG TTACAGCAGC TCTAGGAAAC *1649
TAATACAGCT GCTAAAATGA TCCCTGTCTC CTCGTGTTTA CATTCTGTGT *1699
GTGTCCCCTC CCACAATGTA CCAAAGTTGT CTTTGTGACC AATAGAATAT *1749
GGCAGAAGTG ATGGCATGCC ACTTCCAAGA TTAGGTTATA AAAGACACTG *1799
CAGCTTCTAC TTGAGCCCTC TCTCTCTGCC ACCCACC GCC CCCAATCTAT *1849
CTTGGCTCAC TCGCTCTGGG GGAAGCTAGC TGCCATGCTA TGAGCAGGCC *1899
TATAAAGAGA CTTACGTGGT AAAAAATGAA GTCTCCTGCC CACAGCCACA *1949
TTAGTGAACC TAGAAGCAGA GACTCTGTGA GATAATCGAT GTTTGTTGTT *1999
TTAAGTTGCT CAGTTTTGGT CTAAC TTGTT ATGCAGCAAT AGATAAATAA *2049
TATGCAGAGA AAGAGAAACA AA