

Infevers - PSMB9 (NM_002800.5) - cDNA + Protein - 2023-04-02

AGTGCCCCAG	GCGGCGAGGA	GAGCGGTGCC	TTGCAGGGAT	GCTGCGGGCG	12	
						Me tLeuArgAla 4
GGAGACCAA	CCGGGACTT	ACCCGGGCG	GGAGAAGTCC	ACACGGGAC	62	
GlyAlaProT	hrGlyAspLe	uProArgAla	GlyGluValH	isThrGlyTh	21	
CACCATCATG	GCAGTGGAGT	TTGACGGGGG	<u>CGTTGTGATG</u>	GGTTCTGATT	112	<u>V32I</u>
rThrIleMet	AlaValGluP	heAspGlyGl	yValValMet	GlySerAspS	38	
CCCAGTGTC	TGCAGGCGAG	GCGGTGGTGA	ACCGAGTGTT	TGACAAGCTG	162	
erArgValSe	rAlaGlyGlu	AlaValValA	snArgValPh	eAspLysLeu	54	
TCCCCGCTGC	ACGAGCGCAT	CTACTGTGCA	CTCTCTGGTT	CAGCTGCTGA	212	<u>R60C</u>
SerProLeuH	isGluArgIl	eTyrCysAla	LeuSerGlyS	erAlaAlaAs	71	
TGCCCAAGCC	GTGGCCGACA	TGGCCGCCTA	CCAGCTGGAG	CTCCATGGGA	262	
pAlaGlnAla	ValAlaAspM	etAlaAlaTy	rGlnLeuGlu	LeuHisGlyI	88	
TAGAACTGGA	GGAACCTCCA	CTTGTTTTGG	CTGCTGCAAA	TGTGGTGAGA	312	
leGluLeuGl	uGluProPro	LeuValLeuA	laAlaAlaAs	nValValArg	104	
AATATCAGCT	ATAAATATCG	AGAGGACTTG	TCTGCACATC	TCATGGTAGC	362	
AsnIleSerT	yrLysTyrAr	gGluAspLeu	SerAlaHisL	euMetValAl	121	
TGGCTGGGAC	CAACGTGAAG	GAGGTCAGGT	ATATGGAACC	CTGGGAGGAA	412	
aGlyTrpAsp	GlnArgGluG	lyGlyGlnVa	lTyrGlyThr	LeuGlyGlyM	138	
TGCTGACTCG	ACAGCCTTTT	GCCATTGGTG	GCTCCGGCAG	CACCTTTATC	462	
etLeuThrAr	gGlnProPhe	AlaIleGlyG	lySerGlySe	rThrPheIle	154	
TATGTTATG	TGGATGCAGC	ATATAAGCCA	<u>GGCATGTCTC</u>	CCGAGGAGTG	512	<u>G156D G165D</u>
TyrGlyTyrV	alAspAlaAl	aTyrLysPro	GlyMetSerP	roGluGluCy	171	
CAGGCGCTTC	ACCACAGACG	CTATTGCTCT	GGCCATGAGC	CGGGATGGCT	562	<u>R173C</u>
sArgArgPhe	ThrThrAspA	laIleAlaLe	uAlaMetSer	ArgAspGlyS	188	
CAAGCGGGGG	TGTCATCTAC	CTGGTCACTA	TTACAGCTGC	CGGTGTGGAC	612	

erSerGlyG1 yValIleTyr LeuValThrI leThrAlaAl aGlyValAsp 204

CATCGAGTCA TCTTGGGCAA TGAAGTCCA AAATTCTATG ATGAGTGAAC *2

HisArgValI leLeuGlyAs nGluLeuPro LysPheTyrA spGluStop

CTTCCCAGAG CTTCTCTTTC TTATTTTGTG ATAAACTCTC TAGGGCCAAA *52

ACCTGGTATG GTCATTGGGA AATGAGTGCT CAGGGAGATG GAGCTTAGGG *102

GAGGTGGGTG CTTCCCTCCT AGATGTCAGC ATACACTCTT TCTTCTTTTG *152

TCCCAGGTCT AAAACATCTT TCCTAGAGAA AACAAAAGGG ACTAAACTAG *202

AAATATAAAG AGCCCTATAC ATGACAGGTG ATCACGTACT GAATGATTTT *252

GAAGTAGTAC AAACAATAAA AATTCTCATT CCGCATCATC ATGCGGTCCA *302

TGATGATGAG GCCGCAA

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