

Infevers - PSTPIP1 (NM_003978.5) - cDNA + Protein - 2023-02-09

AAACCTTCCT GCGCAGGCCT CGGGCTGCCT GCCTGCCTGC CTGCCTGGCC -399
 CGGCCCGAGC TCCAGCCTGC CTCTTCCACT GGCCACTGCC TCCCACCCAG -349
 GGCTGGCATC CCTGCTCCCT GCCCTGGGTC CCAGACTGTG TCCTCCATCA -299
 CCGCAGGGTC GGTGAGGGC TGGGCTGGAC ACCAGGGCCC GCCCTCCAT -249 -282G>T
 CACTGAGCTC CACTCCTTCC TCATTTTGCT GCTGATTCTA GCCCCAAACA -199
 AAACAGGTTG AGCTTTTTCC TCCCCTCAGA AGCTCCTCTC TGGCTCGTGG -149 c.262G>A
 CTGCCTTCTG AGTGTTGCAG ACGGCGCCGG CCGGGAAGGG GGGCCTGGGC -99
 CAGCCCTGCC AGGACTGGGA CGCTGCTGCT GGCCTGGC CCTCCATCAG -49
 GCCAGCCTGT GGCAGGAGAG TGAGCTTTGC CGCGGCAGAC GCCTGAGGAT 2

Me 1

GATGCCCCAG CTGCAGTTCA AAGATGCCTT TTGGTGCAGG GACTTCACAG 52 M2T
 tMetProGln LeuGlnPheL ysAspAlaPh eTrpCysArg AspPheThrA 18

CCCACACGGG CTACGAGGTG CTGCTGCAGC GGCTTCTGGA TGGCAGGAAG 102 T20M K34T
 laHisThrGl yTyrGluVal LeuLeuGlnA rgLeuLeuAs pGlyArgLys 34

ATGTGCAAAG ACATGGAGGA GCTACTGAGG CAGAGGGCCC AGGCGGAGGA 152 A49V
 MetCysLysA spMetGluGl uLeuLeuArg GlnArgAlaG lnAlaGluGl 51

CGGTACGGG AAGGAGCTGG TGCAGATCGC ACGGAAGCA GGTGGCCAGA 202 E51D R52Q L57R R62W A64T
 uArgTyrGly LysGluLeuV alGlnIleAl aArgLysAla GlyGlyGlnT 68

CGGAGATCAA CTCCCTGAGG GCCTCCTTTG ACTCCTTGAA GCAGCAAATG 252 T68M
 hrGluIleAs nSerLeuArg AlaSerPheA spSerLeuLy sGlnGlnMet 84

GAGAATGTGG GCAGCTCACA CATCCAGCTG GCCCTGACCC TGCGTGAGGA 302
 GluAsnValG lySerSerHi sIleGlnLeu AlaLeuThrL euArgGluGl 101

GCTGCGGAGT CTCGAGGAGT TTCGTGAGGAG GCAGAAGGAG CAGAGGAAGA 352 E110D
 uLeuArgSer LeuGluGluP heArgGluAr gGlnLysGlu GlnArgLysL 118

AGTATGAGGC CGTCATGGAC CGGGTCCAGA AGAGCAAGCT GTCGCTCTAC 402 Y119C V122I M123 L416delinsT M123 K128delinsTMLLSO D124 K128delinsGMLLS L133 K136delinsPPDP)
 ysTyrGluAl aValMetAsp ArgValGlnL ysSerLysLe uSerLeuTyr 134

AAGAAGGCCA TGGAGTCCAA GAAGACATAC GAGCAGAAGT GCCGGGACGC 452
 LysLysAlaM etGluSerLy sLysThrTyr GluGlnLysC ysArgAspAl 151

GGACGACGCG GAGCAGGCCT TCGAGCGCAT TAGCGCCAAC GGCCACCAGA 502
aAspAspAla GluGlnAlaP heGluArgIl eSerAlaAsn GlyHisGlnL 168

AGCAGGTGGA GAAGAGTCAG AACAAAGCCA GGCAGTGCAA GGACTCGGCC 552 G543A
ysGlnValGl uLysSerGln AsnLysAlaA rgGlnCysLy sAspSerAla 184

ACCGAGGCAG AGCGGGTATA CAGGCAGAGC ATTGCGCAGC TGGAGAAGGT 602
ThrGluAlaG luArgValTy rArgGlnSer ileAlaGlnL euGluLysVa 201

CCGGGCTGAG TGGGAGCAGG AGCACCGGAC CACCTGTGAG GCCTTTCAGC 652 R202W R210W
lArgAlaGlu TrpGluGlnG luHisArgTh rThrCysGlu AlaPheGlnL 218

TGCAAGAGTT TGACCGGCTG ACCATTCTCC GCAACGCCCT GTGGGTGCAC 702 c.C850T: p.Q284X Q219H E220 I226delinsDRQFHPG D222 L416delinsSTRKCG) R228C A230T
euGlnGluPh eAspArgLeu ThrIleLeuA rgAsnAlaLe uTrpValHis 234

AGCACCAGC TCTCCATGCA GTGTGTCAAG GATGATGAGC TCTACGAGGA 752 N236K Q237 M240delinsRHRQ) S239P D246N E250Q E250K
SerAsnGlnL euSerMetGl nCysValLys AspAspGluL euTyrGluGl 251

AGTGCGGCTG ACGCTGGAAG GCTGCAGCAT AGACCCGAC ATCGACAGTT 802 T255M E257K E256G G258R G258A D266N
uValArgLeu ThrLeuGluG lyCysSerIl eAspAlaAsp ileAspSerP 268

TCATCCAGGC CAAGAGCACG GGCACAGAGC CCCCCGCTCC GGTGCCCTAC 852 T274M E277D
heIleGlnAl aLysSerThr GlyThrGluP roProAlaPr oValProTyr 284

CAGAACTATT ACGATCGGGA GGTCACCCCG CTGACCAGCA GCCCTGGCAT 902 D289H
GlnAsnTyrT yrAspArgGl uValThrPro LeuThrSerS erProGlyIl 301

ACAGCCGTCC TGCGGCATGA TAAAGAGGTT CTCTGGACTG CTGCACGGAA 952 C915T
eGlnProSer CysGlyMetI leLysArgPh eSerGlyLeu LeuHisGlyS 318

GTCCCAAGAC CACTTCGTTG GCAGCTTCTG CTGCGTCCAC AGAGACCCTG 1002 S323L A329V
erProLysTh rThrSerLeu AlaAlaSerA laAlaSerTh rGluThrLeu 334

ACCCCCACCC CCGAGCGGAA TGAGGGTGTC TACACAGCCA TCGCAGTGCA 1052 T337Pfs*52 P338 A347delinsLSERRVHTS E339 D341delinsGGM Val344Ile Y345C
ThrProThrP roGluArgAs nGluGlyVal TyrThrAlaI leAlaValGl 351

GGAGATACAG GGAAACCCGG CCTCACCAGC CCAGGAGTAC CGGGCGCTCT 1102 E352K R365W
nGluIleGln GlyAsnProA laSerProAl aGlnGluTyr ArgAlaLeuT 368

ACGATTATAC AGCGCAGAAC CCAGATGAGC TGGACCTGTC CGCGGGAGAC 1152 [D369G](#) [A372V](#) [D384G](#)
yrAspTyrTh rAlaGlnAsn ProAspGluL euAspLeuSe rAlaGlyAsp 384

ATCCTGGAGG TGATCCTGGA AGGGGAGGAT GGGCTGGTGA CTGTGGAGAG 1202 [c.1154T>C](#) [G1179A](#) [G395S](#)
IleLeuGluV alIleLeuGl uGlyGluAsp GlyTrpTrpT hrValGluAr 401

GAACGGGCAG CGTGGCTTCG TCCCTGGTTC CTACCTGGAG AAGCTTTGAG *1 [G403R](#) [G403E](#) [R405C](#) [F407L](#) [V408I](#)
gAsnGlyGln ArgGlyPheV alProGlySe rTyrLeuGlu LysLeuStop 418

GAAGGGCCAG GAGCCCCTTC GGACCTGCCC TGCCAGTGGA GCCAGCAGTG *51
CCCCCAGCAC TGTCCCACC TTGCTAGGGC CCAGAACCAA GCGTCCCCCA *101
GCCCCGAGAG GGAGCCTGTC GTCTCCCAGG GAATAAAGGA GTGCGTTCTG *151
TTCTCCTTGG TGTGCTGGGG TCCCGTTCTC TTTTCTCCT GCTCCAGTGT *201 [c.*156_158del](#)
CCGAGTGCTC AGTTCAGAGG AGGCAAAGGA ACAAGGGAAG GAGCCTGGAT *251
GTGGAGCTCC CCAACTCAGC CGAGGCTTCA GCTATAGTTG GAGAAGAG

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