

Infervers - POMP (NM_015932.6) - cDNA + Protein - 2023-02-09

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GGCGGGGTCG ACTGACGGTA ACGGGGCAGA GAGGCTGTTC GCAGAGCTGC -6
GGAAGATGAA TGCCAGAGGA CTTGGATCTG AGCTAAAGGA CAGTATTCCA 45
    MetAs nAlaArgGly LeuGlySerG luLeuLysAs pSerIlePro 15

GTTACTGAAC TTTCAGCAAG TGGACCTTTT GAAAGTCATG ATCTTCTTCG 95
ValThrGluL euSerAlaSe rGlyProPhe GluSerHisA spLeuLeuAr 32

GAAAGGTTTT TCTTGTGTGA AAAATGAACT TTTGCCTAGT CATCCCCTTG 145
gLysGlyPhe SerCysValL ysAsnGluLe uLeuProSer HisProLeuG 49

AATTATCAGA AAAAAATTTT CAGCTCAACC AAGATAAAAT GAATTTTTTC 195
luLeuSerGl uLysAsnPhe GlnLeuAsnG lnAspLysMe tAsnPheSer 65

ACACTGAGAA ACATTCAGGG TCTATTTGCT CCGCTAAAAT TACAGATGGA 245
ThrLeuArgA snIleGlnGl yLeuPheAla ProLeuLysL euGlnMetGl 82

ATTCAAGGCA GTGCAGCAGG TTCAGCGTCT TCCATTTCTT TCAAGCTCAA 295
uPheLysAla ValGlnGlnV alGlnArgLe uPropheLeu SerSerSerA 99

ATCTTTCACT GGATGTTTTG AGGGGTAATG ATGAGACTAT TGGATTGAG 345 Ile112Trpfs*3 Phe114Leufs*18
snLeuSerLe uAspValLeu ArgGlyAsnA spGluThrIl eGlyPheGlu 115

GATATTCTTA ATGATCCATC ACAAAGCGAA GTCATGGGAG AGCCACACTT 395
AspIleLeuA snAspProSe rGlnSerGlu ValMetGlyG luProHisLe 132

GATGGTGGAA TATAAACTTG GTTTACTGTA ATAGTGTGCT GTTCATGGAA *19
uMetValGlu TyrLysLeuG lyLeuLeuSt op

ACCGAGGGCT GCATCTTGTT TATAGTCATC TTTGTACTGT AATTTGATGT *69
ACACAACATT AAAAGTACTG ACACCTGAGA ATTTCTGCTC AAGTAGTATC *119
AGTGATCATT TAAAATTTGG AGGGGTCTTT GGTTTACAGC CATGTGACAA *169
TTAAAAGCAC TAAAGGGAGA TCATGTTAAA GCTCTTAATT TATATTAAAA *219
CAGTAGCCTT TGTCTTTAAA AAAGTTGTTG CTCATGAATA TTATAAAATG *269
ATCTACAGGT TTCAATTCAA CCTGTTTCTA GGTTTTTTTTG TAAATTTAGT *319
TTTGATTAAG CATTATAAGC ATTTGAGTCT ATAAACTTTA TAGTAGCATC *369
TTTCAGAATA AACATTTTTA ATTGATTTCA GTGGCAACTC TCAAATTGAT *419
TACAATATGA GATATATCAG TGTCGTCCAT TAACACTCAT AAGAATAATA *469

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TTTACTGTGT CAGTGCTATT TTAGGATTAT AGTTATTGTT TGATTATTTTC *519
AGGTTGAAAA GTAGAAGTTC CAAGGTTTTG ATTTTGGTCT GGTCTTTAAG *569
TGAAAAATTA AAGCAACCAG TAGATGTAGG TTAAACTTTT ACTTCATAGA *619
CTTAATATGT AATTAATATA TTGCCAAGCA ACACTGTTAA AGAAAAGTAA *669
AACTCATTTT TTCTTGTTCT TAATTTATAT ATTACAAGAT ACTGTAAGGT *719
ATTCTTTATG AAGTTGATAT ATAAAAATTT ACATTTTATG AACATTAGTG *769
AATGGATCAT CTTTTACAAT TAAAAGTATA TTTTGATTAT CAGTTTCTTA *819
GGAACATCTT TGATTAGATA AATTACTGAT TTGAAACA

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