



POMP (NM_015932.6) - cDNA + Protein - 2025-04-03

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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 AAAAAATTC CAGCTCAACC AAGATAAAAT GAATTTTTCC 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 p.t111fs Ile112Trpfs*3 Phe114Leufs*18
snLeuSerLe uAspValLeu ArgGlyAsnA spGluThrIl eGlyPheGlu 115
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GATATTCCTTA 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 TATATTAATA *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

TTTACTGTGT CAGTGCTATT TTAGGATTAT AGTTATTGTT TGATTATTTT *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

ATTCCTTATG AAGTTGATAT ATAAAAATTT ACATTTTATG AACATTAGTG *769

AATGGATCAT CTTTTACAAT TAAAAGTATA TTTTGATTAT CAGTTTCTTA *819

GGAACATCTT TGATTAGATA AATTACTGAT TTGAAACA

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