



POMP (NM_015932.6) - cDNA + Protein - 2024-05-19

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 TCCATTCTT TCAAGCTCAA 295

uPheLysAla ValGlnGlnV alGlnArgLe uProPheLeu SerSerSerA 99

ATCTTTCACT GGATGTTTTG AGGGGTAATG ATGAGACIAT TGGATTIGAG 345 p.t111fs 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 TTTGTAAGT AATTTGATGT *69

ACACAACATT AAAAGTACTG ACACCTGAGA ATTTCTGCTC AAGTAGTATC *119

AGTGATCATT TAAAATTTGG AGGGGTCTTT GGTTCACAGC CATGTGACAA *169

TTAAAAGCAC TAAAGGGAGA TCATGTTAAA GCTCTTAATT TATATTAAAA *219

CAGTAGCCTT TGTCTTTAAA AAAGTTGTTG CTCATGAATA TTATAAAATG *269

ATCTACAGGT TTCAATTCAA CCTGTTTCTA GGTTTTTTTG 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

ATTCTTTATG AAGTTGATAT ATAAAAATTT ACATTTTATG AACATTAGTG *769

AATGGATCAT CTTTTACAAT TAAAAGTATA TTTTGATTAT CAGTTTCTTA *819

GGAACATCTT TGATTAGATA AATTACTGAT TTGAAACA

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