



PSTPIP1 (NM_003978.5) - CDNA - 2024-05-07

AAACCTTCCT GCGCAGGCCT CGGGCTGCCT GCCTGCCTGC CTGCCTGCC -399
CGGCCCGAGC TCCAGCCTGC CTCTTCCACT GGCCACTGCC TCCCACCCAG -349
GGCTGGCATC CCTGCTCCCT GCCCTGGGTG CCAGACTGTG TCCTCCATCA -299
CCGCAGGGTC GGTGAGGGGC TGGGCTGGAC ACCAGGGCCC GCCCTCCCAT -249 -282G>T
CACTGAGCTC CACTCCTTCC TCATTTGCT GCTGATTCTA GCCCCAAACA -199
AAACAGGTTG AGCTTTTCC TCCCCTCAGA AGCTCCTCTC TGGCTCGTGG -149 c.262G>A
CTGCCTCTG AGTGTTCAG ACGGCGCCGG CCGGGAAAGGG GGGCCTGGGC -99
CAGCCCTGCC AGGACTGGGA CGCTGCTGCT GGCGCCTGGC CCTCCATCAG -49
GCCAGCCTGT GGCAGGAGAG TGAGCTTTGC CGCGGCAGAC GCCTGAGGAT 2
GATGCCCGAG CTGCAGTTCA AAGATGCCTT TTGGTGCAGG GACTTCACAG 52 M2T
CCCACACGGG CTACGAGGTG CTGCTGCAGC GGCTTCTGGA TGGCAGGAAG 102 T20M K34T
ATGTGCAAAG ACATGGAGGA GCTACTGAGG CAGAGGGCCC AGGCGGAGGA 152 A49V
CGGTACGGG AAGGAGTGG TGCAGATCGC ACGGAAGGCA GGTGGCCAGA 202 E51D R52Q L57R R62W A64T
CGGAGATCAA CTCCCTGAGG GCCTCCTTG ACTCCTTGAA GCAGCAAATG 252 T68M
GAGAATGTGG GCAGCTCACA CATCCAGCTG GCCCTGACCC TGCAGTGAGGA 302
GCTGCGGAGT CTCGAGGAGT TTCGTGAG GCAGAAGGAG CAGAGGAAGA 352 E110D
AGTATGAGGC CGTCATGGA CGGGTCCAGA AGAGCAAGCT GTCGCTCTAC 402 Y119C V122I M123 L416delinsT M123 K128delinsTMLLSQ D124 K128delinsGMLLS L133 K136delinsPPDP
AAGAAGGCCA TGGAGTCAA GAAGACATAC GAGCAGAAGT GCCGGGACGC 452
GGACGACGGG GAGCAGGCCT TCGAGCGCAT TAGGCCAAC GGCCACCAGA 502
AGCAGGTGGA GAAGAGTCAG AACAAAGCCA GGCAGTGCAA GGACTCGGCC 552 G543A
ACCGAGGCAG AGCGGGTATA CAGGCAGAGC ATTGCGCAGC TGGAGAAGGT 602

CGGGCTGAG TGGGAGCAGG AGCACCGGAC CACCTGTGAG GCCTTCAGC 652 R202W R210W
TGCAAGAGTT TGACCGGCTG ACCATTCTCC GCAACGCCCT GTGGGTGCAC 702 c.C850T: p.Q284X Q219H E220 I226delinsDROFHPG D222 L416delinsSTRKCG1 R228C A230T
AGCAACCAGC TTCCATGCA GTGTGTCAAG GATGATGAGC TCTACGAGGA 752 N236K Q237 M240delinsRHQO) S239P Q241R D246N E250Q E250K
AGTGCGGCTG ACGCTGGAG GCTGCAGCAT AGACGCCGAC ATCGACAGTT 802 T255M E257K E256G G258R G258S G258A D266N
TCATCCAGGC CAAGAGCAG GGCACAGGC CCCCCGCTCC GGTGCCCTAC 852 T274M E277D
CAGAACTATT ACGATCGGGA GGTCACCCCG CTGACCAGCA GCCTGGCAT 902 D289H P299S
ACAGCCGTCC TGCGGCATGA TAAAGAGGTT CTCTGGACTG CTGCACGGAA 952 C915T
GTCCCAGAC CACTTCGTTG GCAGCTTCTG CTGCGTCCAC AGAGACCTG 1002 S323L A329V
ACCCCCACCC CCGAGGGGAA TGAGGGTGTC TACACAGCCA TCGCAGTGCA 1052 T337Pfs*52 P338 A347delinsLSERRVHTS E339 D341delinsGGM Val344Ile Y345C
GGAAATACAG GGAAACCCGG CCTCACCAGC CCAGGAGTAC CGGGCGCTCT 1102 E352K E352A R365W
ACGATTATAC ACGCGCAGAAC CCAGATGAGC TGGACCTGTC CGCGGGAGAC 1152 D369G A372V D384G
ATCCTGGAGG TGATCCTGGA AGGGGAGAT GGCTGGTGGA CTGTGGAGAG 1202 c.1154T>C G1179A G395S
GAACGGGCAG CGTGGCTTCG TCCCTGGTTC CTACCTGGAG AAGCTTTGAG *1 G403R G403E R405C F407L V408I
GAAGGGCCAG GAGCCCCTTC GGACCTGCC TGCCAGTGGA GCCAGCAGTG *51
CCCCCAGCAC TGTCCCCACC TTGCTAGGGC CCAGAACCAA GCGTCCCCCA *101
GCCCGAGAG GGAGCCTGTC GTCTCCCAGG GAATAAAGGA GTGCGTTCTG *151
TTCTCTTGG TGTGCTGGGG TCCCGTTCTC TTTTTCTCCT GCTCCAGTGT *201 c.*156_158del
CCGAGTGCTC AGTTCAGAGG AGGCAAAGGA ACAAGGGAAG GAGCCTGGAT *251
GTGGAGCTCC CCAACTCAGC CGAGGCTTCA GCTATAGTTG GAGAAGAG

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