

Product datasheet for SC203606

OriGene Technologies, Inc.

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PIB5PA (INPP5J) (NM 001002837) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: PIB5PA (INPP5J) (NM_001002837) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: INPP5J

Synonyms: INPP5; PIB5PA; PIPP

ACCN: NM_001002837

Insert Size: 324 bp

Insert Sequence: >SC203606 3'UTR clone of NM_001002837

The sequence shown below is from the reference sequence of NM_001002837. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGGCTGGAGGAAGGGGCCCTGGGGCCCTGAGGGTGGGGTAGGCAGATGGGCCAAGGTGACCACCATTCT GCCTCAATCTTTTGCAAGCCCACCTGCCTCTCCTGCTGCTCCTCCAGCTGTATCTGCACCTGCCTCT CTGTCCTGGCCAGGGGTGGACAACTGGGGTCCCCCAAAACTCAGTCCTGGCACCTCAACTGTGACAATC AGCAAAGCCCCACCCAGGCCCCATCTGGGATGATGGGAGAGGCTCTGGCAGATGTCCCAATCCTGGAGG

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeg: NM 001002837.2





PIB5PA (INPP5J) (NM_001002837) Human 3' UTR Clone - SC203606

Summary: Inositol 5-phosphatase, which converts inositol 1,4,5-trisphosphate to inositol 1,4-

bisphosphate. Also converts phosphatidylinositol 4,5-bisphosphate to phosphatidylinositol 4-phosphate and inositol 1,3,4,5-tetrakisphosphate to inositol 1,3,4-trisphosphate in vitro. May be involved in modulation of the function of inositol and phosphatidylinositol polyphosphate-binding proteins that are present at membranes ruffles (By similarity).[UniProtKB/Swiss-Prot

Function]

Locus ID: 27124

MW: 11.5