

## Product datasheet for SC201146

## PPP2R5C (NM\_178587) Human 3' UTR Clone

## **Product data:**

## OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	PPP2R5C (NM_178587) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	PPP2R5C
Synonyms:	B56G; B56gamma; PR61G
ACCN:	NM_178587
Insert Size:	158 bp
Insert Sequence:	<pre>&gt;SC201146 3'UTR clone of NM_178587 The sequence shown below is from the reference sequence of NM_178587. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CCCCAGGTACTAAAAAAGAGAATAACATGAAAACGCCCAGGGTTACTTGAATGTTTTTATAAGATAGGA ATATATGTCTTCACCATGGGGGGGGTCTCGATTTCACTAACGTTGTATATGAAAATGTCTGCAATAAAA AGTACTTTTAAACTTTGTAA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACC</pre>
<b>Restriction Sites:</b>	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.
RefSeq:	<u>NM 178587.3</u>



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Summary:	The product of this gene belongs to the phosphatase 2A regulatory subunit B family. Protein phosphatase 2A is one of the four major Ser/Thr phosphatases, and it is implicated in the negative control of cell growth and division. It consists of a common heteromeric core enzyme, which is composed of a catalytic subunit and a constant regulatory subunit, that associates with a variety of regulatory subunits. The B regulatory subunit might modulate substrate selectivity and catalytic activity. This gene encodes a gamma isoform of the regulatory subunit B56 subfamily. Alternatively spliced transcript variants encoding different isoforms have been identified. [provided by RefSeq, Jul 2008]
Locus ID:	5527
MW:	5.9

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