

Product datasheet for SC202366

PPIH (NM_006347) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: PPIH (NM_006347) Human 3' UTR Clone

Symbol: PPIH

Synonyms: CYP-20; CYPH; SnuCyp-20; USA-CYP

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM 006347

Insert Size: 226 bp

Insert Sequence: >SC202366 3'UTR clone of NM_006347

The sequence shown below is from the reference sequence of NM_006347. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAGTTTTTTTGTATGCGTA

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.

RefSeq: <u>NM 006347.4</u>



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



ORIGENE

Summary: The protein encoded by this gene is a member of the peptidyl-prolyl cis-trans isomerase

(PPlase) family. PPlases catalyze the cis-trans isomerization of proline imidic peptide bonds in oligopeptides and accelerate the folding of proteins. This protein is a specific component of the complex that includes pre-mRNA processing factors PRPF3, PRPF4, and PRPF18, as well as U4/U5/U6 tri-snRNP. This protein has been shown to possess PPlase activity and may act as a protein chaperone that mediates the interactions between different proteins inside the

spliceosome. [provided by RefSeq, Jul 2008]

Locus ID: 10465

MW: 8.5