

Product datasheet for **SC206637**

PRPF40B (NM_001031698) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: PRPF40B (NM_001031698) Human 3' UTR Clone
Vector: pMirTarget (PS100062)
Symbol: PRPF40B
Synonyms: HYPC
ACCN: NM_001031698
Insert Size: 503 bp
Insert Sequence: >SC206637 3'UTR clone of NM_001031698
The sequence shown below is from the reference sequence of NM_001031698. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CTCCTACAGCAGCTGGATGATCACCAGTGAACCAATGAGCTGTTCTCTGCCTCGGGTCTGTGTGAGGCC
ATGGCTCCTGGGCCACCCTCACCCTGCCTCAGACTTCTCCTTAGTCTGGTCTGTGTCCACTTTTTTC
TAAAGTAACCCACCCAGCACACCATTTGTTGGCACCTCTCAAGTTGCTCTTGGTGTTC AAGGGTCC
CCTACTCCTGGACTAGTGCAGTCTTGCCTCAGCCCCAGACCAGAGATGGGTGGTATATGCCATGTG
GGGTGGGTGATGCCAGTAGATAAAAGTGTGAGAGAAGGGGTCTCCAGGGAAGAGTACAGGCTGTTGGA
CGCAGCCTGGGTGGCAGAGGGCAGGGTCATCACCTCTAGCATCAGTGCCTGCTCCTGCCTGCCCTGGC
CCTGAGGCTCCACTTCTTCTCCACCCAGGACCTAATGTACGTGTTTTGTTTTTTGTTTTTAA
ATAACAATATTTATAACATG
ACGCGTAAGCGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
```

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: [NM_001031698.3](#)



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Summary: This gene encodes a WW-domain containing protein similar to yeast splicing factor PRP40. This protein has been shown to interact with Huntingtin and methyl CpG binding protein 2 (MeCP2). Alternative splicing results in different transcript variants. [provided by RefSeq, Aug 2014]

Locus ID: 25766

MW: 17.6