

Product datasheet for SC331072

IFTAP (NM_001276726) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: IFTAP (NM_001276726) Human Untagged Clone

Tag: Tag Free
Symbol: IFTAP

Synonyms: C11orf74; HEPIS; NWC

Vector: pCMV6-Entry (PS100001)

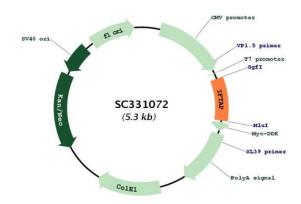
Fully Sequenced ORF: >SC331072 representing NM_001276726.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

AGCCCAGACTTAGAGAAATCCTGTGACTGA

Restriction Sites: Sgfl-Mlul

Plasmid Map:





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IFTAP (NM_001276726) Human Untagged Clone - SC331072

ACCN: NM_001276726

Insert Size: 444 bp

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

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: <u>NM 001276726.1</u>

 RefSeq Size:
 738 bp

 RefSeq ORF:
 444 bp

 Locus ID:
 119710

 UniProt ID:
 Q86VG3

 Cytogenetics:
 11p12

 MW:
 16.8 kDa

Gene Summary: This gene encodes a protein that was identified as a cellular interacting partner of non-

structural protein 10 of the severe acute respiratory syndrome coronavirus (SARS-CoV). The encoded protein may function as a negative regulator of transcription. There is a pseudogene for this gene on chromosome 1. Alternative splicing results in multiple transcript variants.

[provided by RefSeq, Mar 2013]

Transcript Variant: This variant (6) uses an alternate splice site in the 5' UTR and lacks two inframe exons, compared to variant 1. The encoded isoform (b) is shorter than isoform a.

Variants 5, 6, and 7 encode the same isoform (b).