

## **Product datasheet for SC330792**

## PAAF1 (NM 001267806) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

**Product Name:** PAAF1 (NM\_001267806) Human Untagged Clone

Tag: Tag Free
Symbol: PAAF1

Synonyms: PAAF; Rpn14; WDR71

**Vector:** pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC330792 representing NM\_001267806.

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

**CTCTGA** 

**Restriction Sites:** Sgfl-Mlul

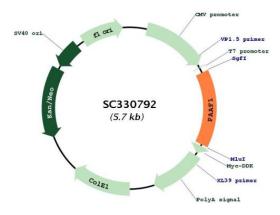
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## Plasmid Map:



**ACCN:** NM\_001267806

**Insert Size:** 834 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: NM 001267806.1

RefSeq Size: 1569 bp RefSeq ORF: 834 bp 80227 Locus ID: **UniProt ID:** Q9BRP4 Cytogenetics: 11q13.4 MW: 30 kDa

This gene encodes a WD repeat-containing protein involved in regulation of association of **Gene Summary:** 

> proteasome components. During HIV infection, the encoded protein is thought to promote provirus transcription through recruitment of the 19S regulatory complex. Alternatively spliced transcript variants encoding multiple isoforms have been observed for this gene.

[provided by RefSeq, Jun 2012]

Transcript Variant: This variant (5) differs in the 5' UTR, lacks an internal exon, and initiates translation at a downstream in-frame start codon, compared to variant 1. The encoded isoform (3) has a shorter N-terminus compared to isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the

transcript record were based on transcript alignments.