

Product datasheet for SC334805

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JKAMP (NM_001284203) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: JKAMP (NM_001284203) Human Untagged Clone

Tag: Tag Free Symbol: JKAMP

Synonyms: C14orf100; C24orf100; CDA06; HSPC213; HSPC327; JAMP

Mammalian Cell

Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001284203, the custom clone sequence may differ by one or

more nucleotides

Restriction Sites: Sgfl-Mlul

ACCN: NM 001284203

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 001284203.1</u>, <u>NP 001271132.1</u>

RefSeq Size: 2461 bp
RefSeq ORF: 747 bp
Locus ID: 51528
UniProt ID: Q9P055
Cytogenetics: 14q23.1

Protein Families: Transmembrane

Gene Summary: May be a regulator of the duration of MAPK8 activity in response to various stress stimuli.

Facilitates degradation of misfolded endoplasmic reticulum (ER) luminal proteins through the

recruitment of components of the proteasome and endoplasmic reticulum-associated

degradation (ERAD) system (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (5) uses an alternate splice site in the 5' region and initiates translation at a downstream in-frame start codon, compared to variant 1. The encoded isoform (5) 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.