

Product datasheet for SC332124

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

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

SET (NM_001248001) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: SET (NM_001248001) Human Untagged Clone

Tag: Tag Free

Symbol: SET

Synonyms: 2PP2A; I2PP2A; IGAAD; IPP2A2; MRD58; PHAPII; TAF-I; TAF-IBETA

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC332124 representing NM_001248001.

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

Restriction Sites: Sgfl-Mlul

ACCN: NM 001248001

Insert Size: 801 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 001248001.1</u>

RefSeq Size: 2562 bp
RefSeq ORF: 801 bp
Locus ID: 6418
UniProt ID: Q01105
Cytogenetics: 9q34.11

Protein Families: Druggable Genome, Phosphatase, Stem cell - Pluripotency

MW: 31.1 kDa

Gene Summary: The protein encoded by this gene inhibits acetylation of nucleosomes, especially histone H4,

by histone acetylases (HAT). This inhibition is most likely accomplished by masking histone

lysines from being acetylated, and the consequence is to silence HAT-dependent transcription. The encoded protein is part of a complex localized to the endoplasmic reticulum but is found in the nucleus and inhibits apoptosis following attack by cytotoxic T lymphocytes. This protein can also enhance DNA replication of the adenovirus genome. Several transcript variants encoding different isoforms have been found for this gene.

[provided by RefSeq, Oct 2011]

Transcript Variant: This variant (4) differs in the 5' UTR and coding sequence compared to variant 1. The resulting isoform (4) has a shorter and distinct 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.