

## **Product datasheet for SC330188**

## SMUG1 (NM 001243789) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

**Product Name:** SMUG1 (NM\_001243789) Human Untagged Clone

Tag: Tag Free Symbol: SMUG1

**Synonyms:** FDG; HMUDG; UNG3

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

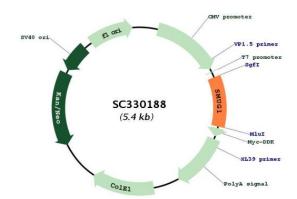
Fully Sequenced ORF: >SC330188 representing NM\_001243789.

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

CAGTGTGACAGAGTCCAAGTCAGAAGACCTGGCTTTTCATCCCAGCTTTGA

**Restriction Sites:** Sgfl-Mlul

Plasmid Map:





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## SMUG1 (NM\_001243789) Human Untagged Clone - SC330188

**ACCN:** NM\_001243789

**Insert Size:** 534 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 001243789.1

 RefSeq Size:
 1046 bp

 RefSeq ORF:
 534 bp

 Locus ID:
 23583

 UniProt ID:
 Q53HV7

 Cytogenetics:
 12q13.13

Protein Families: Druggable Genome
Protein Pathways: Base excision repair

**MW:** 19.6 kDa

**Gene Summary:** This gene encodes a protein that participates in base excision repair by removing uracil from

single- and double-stranded DNA. Many alternatively spliced transcript variants exist for this gene; the full-length nature is known for some but not all of the variants. [provided by RefSeq,

Aug 2011]

Transcript Variant: This variant (4) differs in the 5' UTR, 3' UTR, and coding region compared to variant 1. The resulting isoform (2) has a shorter and distinct C-terminus compared to isoform

1. Variants 4-11 all encode the same isoform (2).