

# **Product datasheet for SC329805**

## STEAP4 (NM\_001205316) Human Untagged Clone

#### **Product data:**

**Product Type:** Expression Plasmids

**Product Name:** STEAP4 (NM\_001205316) Human Untagged Clone

Tag: Tag Free Symbol: STEAP4

Synonyms: SchLAH; STAMP2; TIARP; TNFAIP9

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

Fully Sequenced ORF: >SC329805 representing NM\_001205316.

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

TGGGAAAGGAACTCAAAACACTAG

**Restriction Sites:** Sgfl-Mlul

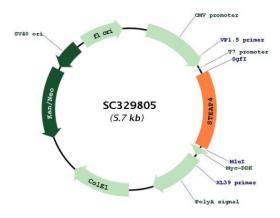
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#### Plasmid Map:



**ACCN:** NM\_001205316

**Insert Size:** 852 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.



### STEAP4 (NM\_001205316) Human Untagged Clone - SC329805

**RefSeq:** <u>NM 001205316.1</u>

 RefSeq Size:
 3960 bp

 RefSeq ORF:
 852 bp

 Locus ID:
 79689

 UniProt ID:
 Q687X5

 Cytogenetics:
 7q21.12

**Protein Families:** Druggable Genome, Transmembrane

MW: 31.3 kDa

**Gene Summary:** The protein encoded by this gene belongs to the STEAP (six transmembrane epithelial antigen

of prostate) family, and resides in the golgi apparatus. It functions as a metalloreductase that has the ability to reduce both Fe(3+) to Fe(2+) and Cu(2+) to Cu(1+), using NAD(+) as acceptor. Studies in mice and human suggest that this gene maybe involved in adipocyte development and metabolism, and may contribute to the normal biology of the prostate cell, as well as prostate cancer progression. Alternatively spliced transcript variants encoding different

isoforms have been found for this gene. [provided by RefSeq, Apr 2011]

Transcript Variant: This variant (3) lacks an in-frame coding exon compared to variant 1, resulting in a shorter isoform (2) missing an internal protein segment compared to isoform 1.