

## **Product datasheet for SC333169**

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## SLC39A1 (NM\_001271961) Human Untagged Clone

#### **Product data:**

**Product Type:** Expression Plasmids

Product Name: SLC39A1 (NM\_001271961) Human Untagged Clone

Tag: Tag Free

Symbol: SLC39A1

Synonyms: ZIP1; ZIRTL

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

Fully Sequenced ORF: >SC333169 representing NM\_001271961.

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

**GCATGA** 

**Restriction Sites:** Sgfl-Mlul

**ACCN:** NM\_001271961

**Insert Size:** 351 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.



### SLC39A1 (NM\_001271961) Human Untagged Clone - SC333169

**RefSeq:** NM 001271961.1

 RefSeq Size:
 2024 bp

 RefSeq ORF:
 351 bp

 Locus ID:
 27173

 UniProt ID:
 Q9NY26

 Cytogenetics:
 1q21.3

**Protein Families:** Transmembrane

**MW:** 11.8 kDa

**Gene Summary:** This gene encodes a member of the zinc-iron permease family. The encoded protein is

localized to the cell membrane and acts as a zinc uptake transporter. This gene has been linked to prostate cancer, breast cancer, and Alzheimer's disease. Alternative splicing results

in multiple transcript variants. [provided by RefSeq, Dec 2012]

Transcript Variant: This variant (6) differs in the 5' UTR and lacks an exon in the 3' coding region which results in a frameshift, compared to variant 1. The encoded isoform (b) is

shorter and has a distinct C-terminus, compared to isoform a.