

# **Product datasheet for SC333726**

## PHPT1 (NM 001287343) Human Untagged Clone

### **Product data:**

**Product Type:** Expression Plasmids

Product Name: PHPT1 (NM\_001287343) Human Untagged Clone

Tag: Tag Free Symbol: PHPT1

Synonyms: CGI-202; HEL-S-132P; HSPC141; PHP; PHP14

Mammalian Cell Neomycin

Selection:

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

Fully Sequenced ORF: >SC333726 representing NM\_001287343.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

CTATGGTCCTGCCCAGCACGCCATTTCAACTGA

**ACGCGTACGCGGCCGCTC**GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

**Restriction Sites:** Sgfl-Mlul

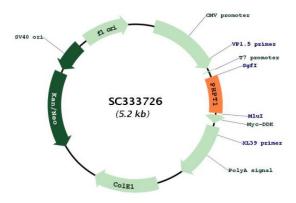
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#### Plasmid Map:



**ACCN:** NM\_001287343

**Insert Size:** 378 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 001287343.1</u>



### PHPT1 (NM\_001287343) Human Untagged Clone - SC333726

RefSeq Size: 1257 bp
RefSeq ORF: 378 bp
Locus ID: 29085
Cytogenetics: 9q34.3

**Protein Families:** Druggable Genome

**Protein Pathways:** Fructose and mannose metabolism, Metabolic pathways, Riboflavin metabolism, Thiamine

metabolism

MW: 13.8 kDa

**Gene Summary:** This gene encodes an enzyme that catalyzes the reversible dephosphorylation of histidine

residues in proteins. It may be involved in the dephosphorylation of G-beta and ATP citrate lyase and in negatively regulating CD4 T lymphocytes by dephosphorylation and inhibition of KCa3.1 channels. Alternative splicing results in multiple transcript variants. [provided by

RefSeq, Dec 2013]

Transcript Variant: This variant (5) uses an alternate in-frame splice site in the 5' coding region and includes an alternate exon in the 3' coding region, compared to variant 4. This results in a frameshift and an early stop codon. It encodes isoform 5, which is shorter and has a distinct

C-terminus, compared to isoform 4.