

Product datasheet for SC330354

THTPA (NM_001256062) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: THTPA (NM_001256062) Human Untagged Clone

Tag: Tag Free
Symbol: THTPA

Synonyms: THTP; THTPASE

Vector: pCMV6-Entry (PS100001)

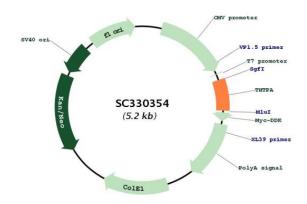
Fully Sequenced ORF: >SC330354 representing NM_001256062.

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

TGTAAGGTGTGCCTGCACAGGAGACAGCACCAGCCAAGCTGA

Restriction Sites: Sgfl-Mlul

Plasmid Map:



ACCN: NM 001256062

Insert Size: 318 bp



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THTPA (NM_001256062) Human Untagged Clone - SC330354

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 001256062.2

RefSeq Size: 1963 bp RefSeq ORF: 318 bp Locus ID: 79178 **UniProt ID:** Q9BU02

14q11.2 **Protein Pathways:** Metabolic pathways, Thiamine metabolism

MW: 12 kDa

Cytogenetics:

Gene Summary: This gene encodes an enzyme which catalyzes the biosynthesis of thiamine disphophate

(vitamin B1) by hydrolysis of thiamine triphosphate. Alternative splicing results in multiple

transcript variants. [provided by RefSeq, Dec 2011]

Transcript Variant: This variant (4) uses an alternate splice site that results in a frameshift in the 3' coding region, compared to variant 1. The encoded isoform (2) has a distinct and shorter C-terminus, compared to isoform 1. Both variants 4 and 5 encode isoform 2.

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.