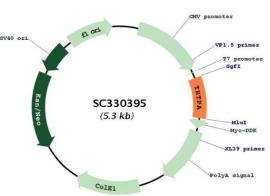


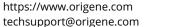
## Product datasheet for SC330395

## THTPA (NM\_001256322) Human Untagged Clone

## Product data:

## **Product Type: Expression Plasmids Product Name:** THTPA (NM 001256322) Human Untagged Clone Tag: Tag Free Symbol: THTPA THTP; THTPASE Synonyms: Vector: pCMV6-Entry (PS100001) **Fully Sequenced ORF:** >SC330395 representing NM\_001256322. Blue=Insert sequence Red=Cloning site Green=Tag(s) GAGTTGGGGGGGCACCCTGGAGTACCGGGTCACCTTCCGAGACACCTACTATGACACCCCTGAGCTGAGC CTCATGCAGGCTGACCACTGGCTGCGACGACGAGGAGGATAGTGGGAGGCTCAAATGTCCTGGAGCA GCAGGTGTCTTAGGACCCCACACGGAGTATAAGGAACTCACAGCGGAACCTACAATTGTGGCCCAACTC TGTGTGCCTGCACAGGAGACAGCACCAGCCAAGCTGATTGTGTATCTACAGCGTTTCCGGCCTCAAGAC TATCAGCGCCTGCTAGAAGTGAACAGCTCCAGAGAGAGGCCACAGGAGACTGAAGATCCTGACCACTGC CTGGGCTAG **Restriction Sites:** Sgfl-Mlul **Plasmid Map:**





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<b>GRIGENE</b> THTPA	(NM_001256322) Human Untagged Clone – SC330395
ACCN:	NM_001256322
Insert Size:	423 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:	<ol> <li>Centrifuge at 5,000xg for 5min.</li> <li>Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li> <li>Close the tube and incubate for 10 minutes at room temperature.</li> <li>Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li> <li>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.</li> </ol>
RefSeq:	<u>NM 001256322.2</u>
RefSeq Size:	1958 bp
RefSeq ORF:	423 bp
Locus ID:	79178
UniProt ID:	<u>Q9BU02</u>
Cytogenetics:	14q11.2
Protein Pathways:	Metabolic pathways, Thiamine metabolism
MW:	16 kDa
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 (6) uses an alternate in-frame splice site in the central coding region, compared to variant 1, resulting in an isoform (3) that is shorter than isoform 1. Both variants 6 and 7 encode isoform 3.

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