

Product datasheet for **SC330949**

THADA (NM_001271644) Human Untagged Clone

Product data:

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|---------------|---|
| Product Type: | Expression Plasmids |
| Product Name: | THADA (NM_001271644) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | THADA |
| Synonyms: | ARMC13; GITA |
| Vector: | pCMV6-Entry (PS100001) |

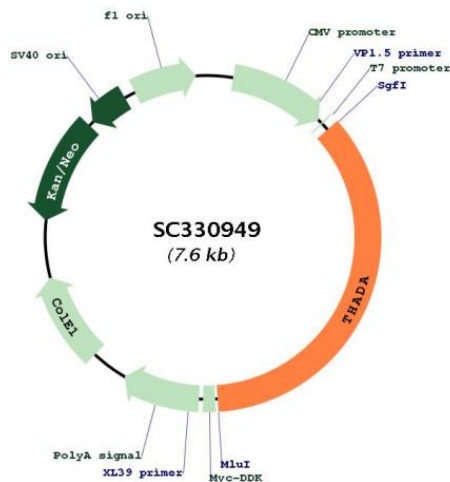


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Fully Sequenced ORF: >SC330949 representing NM_001271644.
Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGGGTGTAAAGAAGAAGAAAGAAATGCAAGTTGCTGCGCTGACCATTTGCCATCAGGACCTTGAAACT
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Restriction Sites: SgfI-MluI

Plasmid Map:


ACCN: NM_001271644

Insert Size: 2685 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: [NM_001271644.1](#)

RefSeq Size: 3066 bp

RefSeq ORF: 2685 bp

Locus ID: 63892

UniProt ID: [Q6YHU6](#)

Cytogenetics: 2p21

MW: 100.7 kDa

Gene Summary:

This gene is the target of 2p21 chromosomal aberrations in benign thyroid adenomas. Single nucleotide polymorphisms (SNPs) in this gene may be associated with type 2 diabetes and polycystic ovary syndrome. The encoded protein is likely involved in the death receptor pathway and apoptosis. [provided by RefSeq, Sep 2016]

Transcript Variant: This variant (5) differs in the 3' coding region and 3' UTR, compared to variant 1. The resulting isoform (c) has a distinct C-terminus and is shorter than isoform a.