

Product datasheet for MR223275L3V

OriGene Technologies, Inc.

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Thoc5 (NM_172438) Mouse Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Thoc5 (NM_172438) Mouse Tagged ORF Clone Lentiviral Particle

Symbol: Thoc5

Synonyms: 1700060C24Rik; A430085L24Rik; Fmip; PK1.3; Sgk2

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-Myc-DDK-P2A-Puro (PS100092)

 Tag:
 Myc-DDK

 ACCN:
 NM_172438

 ORF Size:
 2049 bp

ORF Nucleotide

Sequence:

The ORF insert of this clone is exactly the same as(MR223275).

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through

naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

RefSeq: <u>NM 172438.3, NP 766026.1</u>

RefSeq Size: 2235 bp
RefSeq ORF: 2052 bp
Locus ID: 107829
UniProt ID: Q8BKT7
Cytogenetics: 11 A1





Gene Summary:

Acts as component of the THO subcomplex of the TREX complex which is thought to couple mRNA transcription, processing and nuclear export, and which specifically associates with spliced mRNA and not with unspliced pre-mRNA. TREX is recruited to spliced mRNAs by a transcription-independent mechanism, binds to mRNA upstream of the exon-junction complex (EJC) and is recruited in a splicing- and cap-dependent manner to a region near the 5' end of the mRNA where it functions in mRNA export to the cytoplasm via the TAP/NFX1 pathway. THOC5 in conjunction with ALYREF/THOC4 functions in NXF1-NXT1 mediated nuclear export of HSP70 mRNA; both proteins enhance the RNA binding activity of NXF1 and are required for NXF1 localization to the nuclear rim. Involved in transcription elongation and genome stability. Involved in alternative polyadenylation site choice by recruiting CPSF6 to 5' region of target genes; probably mediates association of the TREX and CFIm complexes. [UniProtKB/Swiss-Prot Function]