

Product datasheet for **RC200382**

THOC5 (NM_001002878) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	THOC5 (NM_001002878) Human Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	THOC5
Synonyms:	C22orf19; Fmip; fSAP79; PK1.3
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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**ORF Nucleotide
Sequence:**

>RC200382 representing NM_001002878
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGCATCGC**

ATGTCATCAGAATCGAGCAAAAAACGGAAGCCCAAAGTGATCCGAAGCGATGGAGCCCCAGCTGAAGGAA
 AGCGGAATCGATCTGACACCGAGCAGGAAGGTAAATACTACAGTGAGGAGCCGAGGTGGATCTGCGGGA
 CCCTGGCAGAGACTATGAGTTATACAAGTACACCTGCCAGGAGCTACAGAGGCTGATGGTGAGATCCAA
 GACCTGAAGAGCAGGGGTGGCAAGGATGTGGCAATAGAAATAGAAGAACGGAGGATCCAGAGCTGTGTGC
 ATTTTCATGACTCTAAAGAAGCTTAACCGATTAGCCACATCAGGTTGAAGAAAGGAAGAGATCAGACCCA
 CGAGGCTAAGCAGAAAGTAGATGCCTATCATCTGCAGCTCCAGAACCTGTTGTATGAGGTGATGCACCTA
 CAGAAGGAGATCACCAAATGTTTGGAGTTAAGTCAAAGCATGAAGAAATTGATCTGGTCAGTTTAGAGG
 AGTTTTATAAGGAGGCTCCACCAGATATCAGCAAGGCCGAAGTCACCATGGGAGACCCCTACCAGCAAAC
 ACTGGCACGTCTGGACTGGGAGCTGGAGCAGCGGAAAAGGCTGGCAGAGAAGTACCGAGAGTGCCTATCT
 AACAGGAGAAGATTCTCAAGGAGATTGAGGTGAAGAAGGAGTACCTGAGCAGCCTCCAGCCCCGCCTCA
 ACAGCATCATGCAGGCTTCCCTTCCGGTGCAGGAGTACCTGTTTATGCCATTTCAGCAGGCTCACAAAGCA
 GTATGAGACAGCCAGACACCTGCCGCCTCCCCTCTATGTCCTCTTTGTTTCAGGCCACTGCGTATGGGCAG
 GCCTGTGATAAGACGTTATCTGTGGCAATCGAAGGCAGTGTGGATGAAGCCAAGGCTCTGTTCAAACCTC
 CAGAGGACTCCCAAGATGACGAGAGTGAATCAGATGCCGAGGAGGAGCAGACTACGAAGCGCCGGAGACC
 CACACTGGGGGTTCAAGTGGACGACAAACGCAAGGAGATGCTGAAGAGGCACCCACTGTCTGTGATGCTC
 GACCTGAAGTGCAAAGATGACAGTGTGCTTACCTGACTTTCTACTACCTCATGAACCTCAACATCATGA
 CAGTAAAAGCCAAAGTGACAATGCCATGGAGCTGATCACCCCATCAGTGCAGGTGACTTGTCTGTCTCC
 TGACTCAGTCCTGAGTTGCTTGTATCCTGGGGATCATGGAAAGAAAACCTCCGAATCCAGCCAATCAGTAT
 CAGTTTGATAAAGTTGGCATCCTGACTTTGAGCGACTATGTACTTGAGCTAGGTACCCCTATTTGTGGG
 TGCAGAAGCTGGGTGGCCTCCACTTCCCAAAGAGCAGCCCCAGCAAACAGTGATTGCTGACCACTCGCT
 GAGCGCCAGCCACATGGAGACCACCATGAACTTCTGAAGACCAGGGTGCAGTCCCGCCTGGCCCTCCAC
 AAACAGTTTGCATCCCTAGAACATGGCATTGTGCCAGTTACCAGTGATTGCCAGTACCTCTTCCCTGCCA
 AGGTTGTCTCTCGCCTGGTGAAATGGGTGACAGTTGCCCATGAGGATTACATGGAGCTGCACTTCACCAA
 AGACATTGTGGATGCGGGACTGGCTGGGACACCAATCTCTACTACATGGCGCTCATCGAAAGGGGCACA
 GCCAAACTGCAGGCCGCTGTGGTGTGAACCCTGGCTACTCCTCCATCCCACCTGTTTTCCAGCTCTGTT
 TGAAGTGAAAGGGGAGAAAACCAACAGCAACGATGACAACATTCGGGCCATGGAGGGCGAAGTCAATGT
 GTGCTACAAGGAGCTGTGTGGCCCTTGGCCAGCCACCAGCTGTTGACCAACCAGCTGCAGCGGCTGTGT
 GTGCTGTGGATGTTTACCTGGAGACCGAGAGCCATGACGACAGTGTGGAGGGGCCCAAGGAATTTCCC
 AGGAGAAGATGTGTCTCGGCTCTTCAAGGGTCTAGCAGGATGAAGCCATTTAAATACAACCATCCTCA
 GGGATTCTTCAGCCATCGC

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >RC200382 representing NM_001002878
 Red=Cloning site Green=Tags(s)

MSESSKKRKPKVIRSDGAPAEGRNRSDEQEGKYYSEAEVDLRDPGRDYELKYTCQELQRLMAEIQ
 DLKSRGGKDVAIIEERRIQSCVHMTLKKLNRLAHIRLKKGRDQTHEAKQKVDAYHLQLQNLLYEVMLH
 QKEITKCLEFKSKHEEIDLVSLEEFYKEAPPDISKAEVMTGDPHQQTARLDWELEQRKRLAEKYRECLS
 NKEKILKEIEVKKEYLSSLQPRLSIMQASLPVQEYLFMPFDQAHKQYETARHLPPPLYVLFVQATAYGQ
 ACDKTLSSVAIEGSVDEAKALFKPPEDSQDDESDSDAEETTKRRRPTLGVLDDKRKEMLRHPLSVML
 DLKCKDDSVLHLTFYYLMNLNIMTVKAKVTTAMELITPISAGDLLSPDSVL SCLYPGDHGKKTNPANQY
 QFDKVGILTLSDYVLELGHYPYLWVQKLGLHFPKEQPQQTVIADHSL SASHMETTMKLLKTRVQSRLALH
 KQFASLEHGIVPVTSDCQYLFPAKVVSRLVKWVTVAHEDYMEHFHFKDIVDAGLAGDTNLYMALIERGT
 AKLQAAVVLNPGYSSIPPVFLCLNWKGEKTSNDDNIRAMEGEVNVCKELCGPWP SHQLLTNQLQRLC
 VLLDVYLETESHDDSVGPKFEPQEKMCLRLFRGP SRMKPFKYNHPQGFFSHR

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Chromatograms: https://cdn.origene.com/chromatograms/mk8104_a05.zip

Restriction Sites: SgfI-MluI

Cloning Scheme:



ACCN: NM_001002878

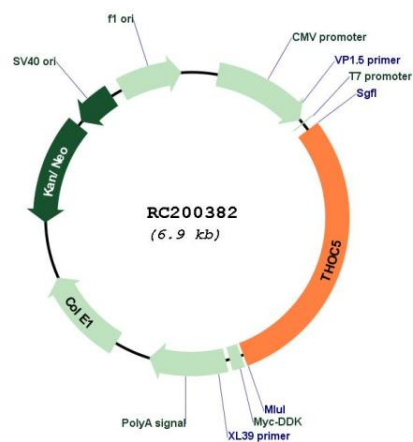
ORF Size: 2049 bp

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.

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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	<u>NM_001002878.1, NP_001002878.1</u>
RefSeq Size:	2694 bp
RefSeq ORF:	2052 bp
Locus ID:	8563
UniProt ID:	<u>Q13769</u>
Cytogenetics:	22q12.2
MW:	78.5 kDa
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. The TREX complex is essential for the export of Kaposi's sarcoma-associated herpesvirus (KSHV) intronless mRNAs and infectious virus production. 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]

Product images:



Circular map for RC200382