

Product datasheet for SC211291

TUT7 (NM 024617) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: TUT7 (NM_024617) Human 3' UTR Clone

Symbol: TUT7

Synonyms: PAPD6; TENT3B; ZCCHC6

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_024617

Insert Size: 974 bp

Insert Sequence: >SC211291 3'UTR clone of NM_024617

The sequence shown below is from the reference sequence of NM_024617. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CCTGTAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul



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TUT7 (NM_024617) Human 3' UTR Clone - SC211291

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 024617.4</u>

Summary: Uridylyltransferase that mediates the terminal uridylation of mRNAs with short (less than 25

nucleotides) poly(A) tails, hence facilitating global mRNA decay (PubMed:19703396,

PubMed:25480299). Essential for both oocyte maturation and fertility. Through 3' terminal

uridylation of mRNA, sculpts, with TUT7, the maternal transcriptome by eliminating

transcripts during oocyte growth (By similarity). Involved in microRNA (miRNA)-induced gene silencing through uridylation of deadenylated miRNA targets (PubMed:25480299). Also

functions as an integral regulator of microRNA biogenesiS using 3 different uridylation

mechanisms (PubMed:25979828). Acts as a suppressor of miRNA biogenesis by mediating the terminal uridylation of some miRNA precursors, including that of let-7 (pre-let-7). Uridylated

pre-let-7 RNA is not processed by Dicer and undergo degradation. Pre-let-7 uridylation is

strongly enhanced in the presence of LIN28A (PubMed:22898984). In the absence of LIN28A,

TUT7 and TUT4 monouridylate group II pre-miRNAs, which includes most of pre-let7

members, that shapes an optimal 3' end overhang for efficient processing

(PubMed:25979828, PubMed:28671666). Add oligo-U tails to truncated pre-miRNAS with a 5'

overhang which may promote rapid degradation of non-functional pre-miRNA species (PubMed:25979828). Does not play a role in replication-dependent histone mRNA

degradation (PubMed:18172165). Due to functional redundancy between TUT4 and TUT7, the

identification of the specific role of each of these proteins is difficult (PubMed:25979828,

PubMed:25480299, PubMed:19703396, PubMed:22898984, PubMed:18172165, PubMed:28671666). TUT4 and TUT7 restrict retrotransposition of long interspersed element-

1 (LINE-1) in cooperation with MOV10 counteracting the RNA chaperonne activity of L1RE1.

TUT7 uridylates LINE-1 mRNAs in the cytoplasm which inhibits initiation of reverse

transcription once in the nucleus, whereas uridylation by TUT4 destabilizes mRNAs in

cytoplasmic ribonucleoprotein granules (PubMed:30122351).[UniProtKB/Swiss-Prot Function]

Locus ID: 79670

MW: 37.3