

## Product datasheet for **RR212067**

### Tut1 (NM\_001033901) Rat Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Tut1 (NM_001033901) Rat Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Tut1
Synonyms:	MGC125034
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



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**ORF Nucleotide Sequence:**

>RR212067 representing NM\_001033901  
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**CGGATCGCC**

ATGGCGCGGTGGATTTCGGATGTTGTATCGCTGCCGCGTGGCCGCTTTCGCTGCTGCCTCTGCGATGTTA  
 CTACAGCCAACCGACCCAGCCTAGATGCCCACTTGAAAGGCCGGAAGCACCGGGATCTGGTACAACCTACG  
 AGCTACCAGAAAAGGCCAGGGACTTCGAAGTGTGTTTGTCACTGGCTTCCAGGGATGTAGGCTCTGCT  
 CAGCTCTCTGAATACTCCAGACATTTGGCCCTGTGGCCAATATTGTTATGGACAAGGACAAGGGGGTGT  
 TTGCCATCGTGGAGATGGGAGACATAAGTGTCTGGGAGGCTGTCTTATCTCAGCCCAAGCACAGCCTTGG  
 GGGACACACACTGCGTGTCCGGCAAGGGAGCAGAAGGAGTTCAGAGCCAGCTTCCAAGTCTCCCAAA  
 GGAGTGGACTCAAATAGTACCAGCTGGCCAAGCACTGGCTGAGGCCGAGATGTAGGGGCACAGATGG  
 TGAAGCTTGTGGAAGTGGGGAGTTGTCTGAGGCTGAGCGGCAGCTTCGGACCCTGTTGTGGCCCTGAT  
 GCAGGAGTCTTACAGAGTTCTTCTGGCTGTGTGGTCCATCCTTTGGCTCTTCTGTCAATAGCTTT  
 GATGTTTATGGCTGTGATCTCGACCTCTTCTGGACTTGGGAGATATGGAAGAGCCCAAGCCAGACCCAC  
 AGACTCCAAAGCTTCCAGAGGCTTCGTCCTTGGACTCAACCTTGGCTTCTTCCCTGGATCCTCAAGTGT  
 GGCTGCACCCAGCTTCTAGACTCACTGTCTCCAACATCTCTCCAAGATTCTGAAGCCCTGGACTTT  
 GAAACCCCTTCTTCTCTGGCACCACAGACCCAGACTCTGCTTTGGGCTCTGACTGTACCTCTCCCC  
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 AGCAGAAGCCTCAAAGGACGAGAAGGAGGAGGCAACAGCAGTGTAGAGCTGGTGGGATCTATTCTCCGT  
 GGCTGTGTCCCTGGAGTGTACCGAGTCCAACTGTGCCCTGCCAGGCGTCTGTGGTCAAGTTCTGTC  
 ATCGCCCTCAGGTCTTATGAGACATCTCCCTCAGTAACCGGTTGGCCTGTATAATTCCCGCTTCT  
 GAACCTATGCTCTGAGATGGATAGTCGAGTCCGGCCCTTGTGTATACCTTACGCTGCTGGGCTCAGCAT  
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 AGACCAGAGACCTCCAGTGCTTCTACTGTGGCCAGCTTACGCAGAGATCAGGTGAAGGGGAGCAGGT  
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 CCTCTCAGTTCCCTGCTGGCCAGTTCTTCTCTGCGTGTCTTGTGGGATCTTCTGGCTCCCTGCTGT  
 CCCTGCGGAAGGTCAGGCACTGATGGTGGCAGGGGCTGCCTTCTGATCTCTGGGAAGGGCTGCGCT  
 TGGCCCATGAATCTCCAGGACCCCTTTGATCTGAGTACAATGTTGCAGCCAATGTGACCAGCCGCGT  
 GCTAAGCGTCTGCAGAGCAGTTGTGGAGCAGCAGTAGTTACTGCCGAGTCTCCAGTACCAGCAGCGCT  
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 CAAGCTTATCCCTTACCCTCTGCCCTTCCACAGATCATTACAGCTTTGGTGAAGTGTGTTAAGGGAA  
 GCACTAGGATGCCATATAGAACAGGGAACCAAGAGACGACGGTCGGAAGGTGCCAGATCTAAGACTCTC  
 CCCTGGGAGGGGCAAAACAAAAGACCAAGACTTAGTGGCAAGAAAAGAGCTGTGAGGAAGGGAAAGAGGA  
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 ACAAGACTGCTGAGCAGAATCCCATGAACTGAGGGGCTGGAGAAGGGTCCCGGTGAGACAGAGAA  
 GGAGGCTTACACCCGTATCAGTGAAGTGGCGCTGTGCTTGTGGCACCATAATGGCAGGGGAGGGCGG  
 CGTGCCCGGAGACGATTTACAGCAGAAACCAAGGAAGAAGGCAGAGGAGTCCCAGCACCGGAGCAGAGT  
 GGCTGGCAGTGGAAAGCAGGTAACCCAGGAAGTGAAGGGACCAAGAGTGAAGCAGCAGAGGCTACAGGG  
 GGAGCCCTCCTCACCTTGTGGCATCGGCTCCCAAGCTGAGCAGACACTCACTGTGGCACCCTCCAG  
 GATCCCAAGGCTGTTCCCTGCTTTCATATTTTCTACAGGTTTTTATCCCTCAAGCACTTAAAAATC  
 TCCTCAAG

**ACGCGT**ACGCGGCGGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >RR212067 representing NM\_001033901  
 Red=Cloning site Green=Tags(s)

MAAVSDVVSLSRGRFRCCLCDVTTANRPSLDAHLKGRKHRDLVQLRATRKAQGLRSVFSVGFPRDVGSA  
 QLSEYFQTFGPVANIVMDKDKGVFAIVEMGDISAREAVLSQPKHSLGGHTLRVPRPQKEFQSPASKSPK  
 GVDNSHQLAQALAEAADVGAQMVKLVLELRELSAERQLRTLVLVLMQEVFTEFFPGCVVHPFGSSVNSF  
 DVHGCDLDFLDLGDMEEPQDPQTPKLEASSLDSTLASSLDPQVLACTPASLDSLSPSLQDSEALDF  
 ETPSSLAPQTPDSALGSDTVTSPQSLPPVSPLEEDRGEKGRKELELAESKDEKEEATAVLELVGSILR  
 GCVPGVYRVQTVPSARRPVVKFCHRPSGLHGDISLSNRLALYNSRFLNLCSEMDSRVRPLVYTLRCWAQH  
 NGLSGGGPLLNNYALTLVVIYFLQTRDPPVLPVVAQLTQRSGEQVEVDGWDCSFPKASRLEPSTNVE  
 PLSLLAQFFSCVSCWDLSGSLLSLREGQALMVAGGLPSDLWEGLRGLGPMNLQDPFDLSHNVAANVTSRV  
 AKRLQSSCGAAASYCRSLQYQQRSSRGRDWGLLPLLQPSPPSLLSAKLIPLPSAFPPIITALVSVLRE  
 ALGCHIEQGTKRRRSEGARSKDSPLGGANKRPRLSGQEKSCGEGKEEPQGCAGDHSENEVEEMVIELRET  
 PQDWALLHCGPPGELPLMTAKLCKTAEQNPMEPEGAGEGSPGETEKEASHPSSVSWRCALWHQIQGRR  
 RARRRFQQQTKEEGRGGPSTGAEWLAVEARVTQELKGPKEQQRLQGEPLLLTFVASASQAEQTLTVAPLQ  
 DPQGLFPLHHFLQVFIQALKNLLK

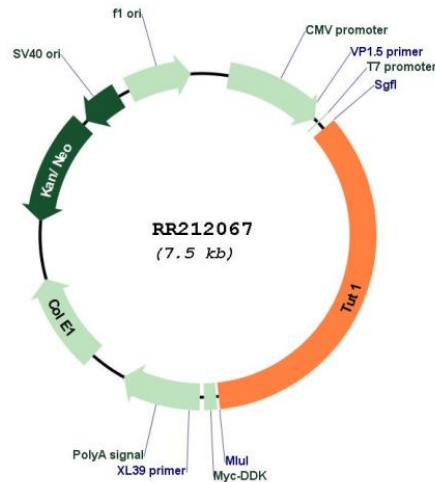
TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Restriction Sites:

Sgfl-MluI

Cloning Scheme:



**Plasmid Map:**


**ACCN:** NM\_001033901

**ORF Size:** 2598 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.

**RefSeq:** [NM\\_001033901.1](#), [NP\\_001029073.1](#)

**RefSeq Size:** 2699 bp

**RefSeq ORF:** 2601 bp

**Locus ID:** 499314

**UniProt ID:** [Q3MHT4](#)

**Cytogenetics:** 1q43

**MW:** 94.4 kDa

**Gene Summary:** Poly(A) polymerase that creates the 3'-poly(A) tail of specific pre-mRNAs. Localizes to nuclear speckles together with PIP5K1A and mediates polyadenylation of a select set of mRNAs, such as HMOX1. In addition to polyadenylation, it is also required for the 3'-end cleavage of pre-mRNAs: binds to the 3' UTR of targeted pre-mRNAs and promotes the recruitment and assembly of the CPSF complex on the 3' UTR of pre-mRNAs. In addition to adenylyltransferase activity, also has uridylyltransferase activity. However, the ATP ratio is higher than UTP in cells, suggesting that it functions primarily as a poly(A) polymerase. Acts as a specific terminal uridylyltransferase for U6 snRNA in vitro: responsible for a controlled elongation reaction that results in the restoration of the four 3'-terminal UMP-residues found in newly transcribed U6 snRNA. Not involved in replication-dependent histone mRNA degradation (By similarity). [UniProtKB/Swiss-Prot Function]