

## **Product datasheet for SC201751**

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## UTP14A (NM\_001166221) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: UTP14A (NM 001166221) Human 3' UTR Clone

Symbol: UTP14A

Synonyms: dJ537K23.3; NYCO16; SDCCAG16; Utp14

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_001166221

**Insert Size:** 174 bp

Insert Sequence: >SC201751 3'UTR clone of NM\_001166221

The sequence shown below is from the reference sequence of NM\_001166221. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAACAGCTGAAGAAATGCTCTGTAGATTGAGTTGCTGGAGGAGTGACAGCCAGGAGCCCTGACTTCACTTCCTTTGGTCCAGTTTTACTCTGATACAGGGTGGATTCCAAAACTGGCTCAGTACATTGCATGATTA

AGCCACATTTTAAAAATAAAGGCATTTTTTAATCTA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul

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 001166221.2</u>





## UTP14A (NM\_001166221) Human 3' UTR Clone - SC201751

Summary: This gene encodes a member of the uridine triphosphate 14 family. As an essential

component of a large ribonucleoprotein complex bound to the U3 small nucleolar RNA, the encoded protein is involved in ribosome biogenesis and 18S rRNA synthesis. An autosomal retrotransposed copy of this X-linked gene exists on chromosome 13. Alternative splicing

results in multiple transcript variants.[provided by RefSeq, Oct 2009]

**Locus ID:** 10813

MW: 6.4