

## **Product datasheet for SC201017**

## NFU1 (NM\_001002757) Human 3' UTR Clone

**Product data:** 

Product Type: 3' UTR Clones

Symbol: NFU1

Synonyms: CGI-33; HIRIP5; MGC142252; MGC142254; Nfu; NifU; NIFUC

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

**ACCN:** NM\_001002757

Insert Size: 157 bp

Insert Sequence: >SC201017 3'UTR clone of NM\_001002757

The sequence shown below is from the reference sequence of NM\_001002757. The complete sequence

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCAGATGAAAAAGAAGCAAACTCACCTTAAAATAATCTGGATTTTCTTTGGGCATAACAGTCAGACTTG
TTGATAATATATCAAGTTTTTATTAATATGCTGAGGAACTTGAAGATTAATAAAAATATGCTCTT

CAGAGAATGATATAAAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Safl-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.



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Note: Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um

filter is required.

**RefSeq:** <u>NM\_001002757.1</u>

Summary: This gene encodes a protein that is localized to mitochondria and plays a critical role in iron-

sulfur cluster biogenesis. The encoded protein assembles and transfers 4Fe-4S clusters to target apoproteins including succinate dehydrogenase and lipoic acid synthase. Mutations in this gene are a cause of multiple mitochondrial dysfunctions syndrome-1, and pseudogenes of

this gene are located on the short arms of chromosomes 1 and 3. Alternatively spliced

transcript variants encoding multiple isoforms have been observed for this gene. [provided by

RefSeq, Dec 2011]

**Locus ID:** 27247

**MW:** 6.2