

Product datasheet for SC206192

RTEL1 (NM_016434) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: RTEL1

Synonyms: C20orf41; DKCA4; DKCB5; NHL; PFBMFT3; RTEL

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

ACCN: NM_016434

Insert Size: 490 bp

Insert Sequence: >SC206192 3'UTR clone of NM_016434

The sequence shown below is from the reference sequence of NM_016434. The complete sequence of

this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GTGCCCA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

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).



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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.

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

filter is required.

RefSeq: <u>NM_016434.4</u>

Summary: This gene encodes a DNA helicase which functions in the stability, protection and elongation of

telomeres and interacts with proteins in the shelterin complex known to protect telomeres during DNA replication. Mutations in this gene have been associated with dyskeratosis congenita and Hoyerall-Hreidarsson syndrome. Read-through transcription of this gene into the neighboring downstream gene, which encodes tumor necrosis factor receptor superfamily,

member 6b, generates a non-coding transcript. Alternative splicing results in multiple

transcript variants encoding different isoforms. [provided by RefSeq, Sep 2013]

Locus ID: 51750

MW: 16.7