

Product datasheet for SC200991

C11orf10 (TMEM258) (NM_014206) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Symbol: Cllorf10

Synonyms: Cllorf10; Kud; Kuduk

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

ACCN: NM_014206

Insert Size: 327 bp

Insert Sequence: >SC200991 3'UTR clone of NM_014206

The sequence shown below is from the reference sequence of NM_014206. The complete sequence of

this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

ATTTGTGTAGCACCGGTGTTTTGTCATTGGAATTTAAGGCTTACCTGTGGA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Safl-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).

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.



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com

EU: info-de@origene.com CN: techsupport@origene.cn



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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_014206.4</u>

Summary: Subunit of the oligosaccharyl transferase (OST) complex that catalyzes the initial transfer of a

defined glycan (Glc(3)Man(9)GlcNAc(2) in eukaryotes) from the lipid carrier dolichol-

pyrophosphate to an asparagine residue within an Asn-X-Ser/Thr consensus motif in nascent

polypeptide chains, the first step in protein N-glycosylation. N-glycosylation occurs

cotranslationally and the complex associates with the Sec61 complex at the channel-forming translocon complex that mediates protein translocation across the endoplasmic reticulum

(ER). All subunits are required for a maximal enzyme activity (PubMed:26472760,

PubMed:27974209). Involved in ER homeostasis in the colonic epithelium (By similarity).

[UniProtKB/Swiss-Prot Function]

Locus ID: 746

MW: 12.1