

Product datasheet for SC210234

MTREX (NM 015360) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: MTREX (NM_015360) Human 3' UTR Clone

Symbol: MTREX

Synonyms: Dob1; fSAP118; KIAA0052; Mtr4; SKIV2L2

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_015360

Insert Size: 813 bp

Insert Sequence: >SC210234 3' UTR clone of NM_015360

The sequence shown below is from the reference sequence of NM_015360. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site

Blue=Stop Codon

CAATTGGCAGAGCTCAGAATTCAAGCGATCGC

TTTGTTATGCCTTCAGCAAATAGCTTCATTTTGCCAATACTGA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCG

Restriction Sites: Sgfl-Mlul



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MTREX (NM_015360) Human 3' UTR Clone - SC210234

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 015360.4</u>

Summary: Component of exosome targeting complexes. Subunit of the trimeric nuclear exosome

targeting (NEXT) complex, a complex that directs a subset of non-coding short-lived RNAs for exosomal degradation. Subunit of the trimeric poly(A) tail exosome targeting (PAXT) complex, a complex that directs a subset of long and polyadenylated poly(A) RNAs for exosomal degradation. The RNA exosome is fundamental for the degradation of RNA in eukaryotic nuclei. Substrate targeting is facilitated by its cofactor MTREX, which links to RNA-binding

protein adapters (PubMed:27871484). Associated with the RNA exosome complex and involved in the 3'-processing of the 7S pre-RNA to the mature 5.8S rRNA (PubMed:17412707, PubMed:29107693). May be involved in pre-mRNA splicing.[UniProtKB/Swiss-Prot Function]

Locus ID: 23517