

## **Product datasheet for SC210303**

## FTSJ2 (MRM2) (NM\_013393) Human 3' UTR Clone

**Product data:** 

Product Type: 3' UTR Clones

Symbol: FTSJ2

**Synonyms:** FJH1; FTSJ2; HEL97; MTDPS17; RRMJ2

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

**ACCN:** NM\_013393

Insert Size: 957 bp

Insert Sequence: >SC210303 3'UTR clone of NM\_013393

The sequence shown below is from the reference sequence of NM\_013393. The complete sequence of

this clone may contain minor differences, such as  $\ensuremath{\mathsf{SNPs}}\xspace.$ 

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TAGGAACTTGACGCAGCTTTATATTTAATAAAATAATAGAGATTGCATTATTTTAAGGTA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul



**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



## FTSJ2 (MRM2) (NM\_013393) Human 3' UTR Clone | SC210303

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.

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

filter is required.

**RefSeq:** <u>NM\_013393.3</u>

Summary: The protein encoded by this gene is a member of the S-adenosylmethionine-binding protein

family. It is a nucleolar protein and it may be involved in the processing and modification of rRNA. This gene has been suggested to be involved in cell cycle control and DNA repair.

[provided by RefSeq, Jul 2008]

**Locus ID:** 29960

**MW:** 36.6