

Product datasheet for SC208742

OriGene Technologies, Inc.

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MSL3L1 (MSL3) (NM_078630) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: MSL3L1 (MSL3) (NM_078630) Human 3' UTR Clone

Symbol: MSL3L1

Synonyms: DKFZp586J1822; drosophila MSL3-like 1; male-specific lethal 3 homolog (Drosophila); male-

specific lethal 3-like 1; MSL3L1; OTTHUMP00000022911; OTTHUMP00000022912

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_078630

Insert Size: 702 bp

Insert Sequence: >SC208742 3'UTR clone of NM_078630

The sequence shown below is from the reference sequence of NM_078630. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGCGGAAAGATCGCCGTG

 ${\sf TAACAATTGGCAGAGCTCAGAATTCAA}{\sf GCGATCGCC}$

CTTGTTTGAAAG

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul





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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 078630.1</u>

Summary: This gene encodes a nuclear protein that is similar to the product of the Drosophila male-

specific lethal-3 gene. The Drosophila protein plays a critical role in a dosage-compensation pathway, which equalizes X-linked gene expression in males and females. Thus, the human protein is thought to play a similar function in chromatin remodeling and transcriptional regulation, and it has been found as part of a complex that is responsible for histone H4 lysine-16 acetylation. This gene can undergo X inactivation. Alternative splicing results in multiple transcript variants. Related pseudogenes have been identified on chromosomes 2, 7

and 8. [provided by RefSeq, Jul 2010]

Locus ID: 10943 MW: 26.4