

## **Product datasheet for SC205425**

## SIL1 (NM 001037633) Human 3' UTR Clone

## **Product data:**

**Product Type:** 3' UTR Clones

**Product Name:** SIL1 (NM 001037633) Human 3' UTR Clone

Symbol: SIL1

Synonyms: BAP; MSS; ULG5

**Mammalian Cell** 

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_001037633

**Insert Size:** 423 bp

Insert Sequence: >SC205425 3'UTR clone of NM\_001037633

The sequence shown below is from the reference sequence of NM\_001037633. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AATCACGTA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul

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.



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

CN: techsupport@origene.cn

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



## SIL1 (NM\_001037633) Human 3' UTR Clone - SC205425

**RefSeq:** <u>NM 001037633.2</u>

Summary: This gene encodes a resident endoplasmic reticulum (ER), N-linked glycoprotein with an N-

terminal ER targeting sequence, 2 putative N-glycosylation sites, and a C-terminal ER

retention signal. This protein functions as a nucleotide exchange factor for another unfolded protein response protein. Mutations in this gene have been associated with Marinesco-Sjogren syndrome. Alternate transcriptional splice variants have been characterized.

[provided by RefSeq, Jul 2008]

Locus ID: 64374 MW: 14.7