

## **Product datasheet for SC204269**

## OriGene Technologies, Inc.

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## Seryl tRNA synthetase (SARS) (NM\_006513) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: Seryl tRNA synthetase (SARS) (NM 006513) Human 3' UTR Clone

**Symbol:** Seryl tRNA synthetase

Synonyms: NEDMAS; SARS; SERRS; SERS

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_006513

**Insert Size:** 324 bp

Insert Sequence: >SC204269 3'UTR clone of NM\_006513

The sequence shown below is from the reference sequence of NM\_006513. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

CTGCAGAACATGGAGGTCACCGATGCTTGAACATTCCTGCCTCCCTATTTGCCAGGCTTTCATTTCTGT CTGCTGAGATCTCAGAGCCTGCCCCAACAGCAGGGAAGCCAAGCACCCATTCATCCCCCTGCCCCCATCT GACTGCGTAGCTGAGAGGGGAACAGTGCCATGTACCACACAGATGTTCCTGTCTCCTCGCATGGGCATA GGGACCCATCATTGATGACTGATGAAACCATGTAATAAAGCATCTCTGGGGAGGGCTTAGGACTCTTCC

TCAGTCTTCTCCCCGGGCTTGAACCCCGCCTCTGAGGTTCTCCCTGA

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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.

**RefSeq:** <u>NM 006513.4</u>





## Seryl tRNA synthetase (SARS) (NM\_006513) Human 3' UTR Clone - SC204269

Summary: This gene belongs to the class II amino-acyl tRNA family. The encoded enzyme catalyzes the

transfer of L-serine to tRNA (Ser) and is related to bacterial and yeast counterparts. Multiple alternatively spliced transcript variants have been described but the biological validity of all

variants is unknown. [provided by RefSeq, Jul 2010]

**Locus ID:** 6301

**MW:** 11.8