

# **Product datasheet for SC202175**

# OriGene Technologies, Inc.

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## CARS (NM\_001014438) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: CARS (NM\_001014438) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: CARS

Synonyms: CARS1; CYSRS; cysteine-tRNA ligase; cysteine translase; cysteine tRNA ligase 1, cytoplasmic;

cysteinyl-tRNA synthetase; MGC:11246; OTTHUMP00000012605

**ACCN:** NM\_001014438

**Insert Size:** 227 bp

Insert Sequence: >SC202175 3'UTR clone of NM\_001014438

The sequence shown below is from the reference sequence of NM\_001014438. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AACAGTGAAAAAAAAAAAA

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 001014438.1</u>





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**Summary:** 

This gene encodes a class 1 aminoacyl-tRNA synthetase, cysteinyl-tRNA synthetase. Each of the twenty aminoacyl-tRNA synthetases catalyzes the aminoacylation of a specific tRNA or tRNA isoaccepting family with the cognate amino acid. This gene is one of several located near the imprinted gene domain on chromosome 11p15.5, an important tumor-suppressor gene region. Alterations in this region have been associated with Beckwith-Wiedemann syndrome, Wilms tumor, rhabdomyosarcoma, adrenocortical carcinoma, and lung, ovarian and breast cancers. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Aug 2010]

**Locus ID:** 833 **MW:** 8.5