

## Product datasheet for SC203175

## OriGene Technologies, Inc.

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## SLC35A2 (NM\_005660) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: SLC35A2 (NM\_005660) Human 3' UTR Clone

Symbol: SLC35A2

Synonyms: CDG2M; CDGX; UDP-Gal-Tr; UGALT; UGAT; UGT1; UGT2; UGTL

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_005660

**Insert Size:** 285 bp

The sequence shown below is from the reference sequence of NM\_005660. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAGTTGCTCACCAAGGTGAAGGGTTCCTAGCCGCTGGGATTGAAGACATTGGCCTGGCCTCGTTCTCTC
TTCTTGCCCTTGGCCCAGCTGGGACCAAACTCTGATCAGTATTAGGGGTAGAGTGAGGTAGACACTGGA
CTCCCTGTCCCCACCACCCCTGCCCACCCAGGGCCAACATGACTAAGCTCTCTCATGACCCACCTCAG
CTCAGCCCCCCAGCCCCTGCCAGCTCCACACTATCTCTTAGCTGAGTTTTTGCAAAATAAAATGTGTTGT

GTATCTTGC

**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 005660.3</u>





## SLC35A2 (NM\_005660) Human 3' UTR Clone - SC203175

Summary: This gene encodes a member of the nucleotide-sugar transporter family. The encoded

protein is a multi-pass membrane protein. It transports UDP-galactose from the cytosol into Golgi vesicles, where it serves as a glycosyl donor for the generation of glycans. Mutations in this gene cause congenital disorder of glycosylation type IIm (CDG2M). Multiple alternatively spliced transcript variants encoding distinct isoforms have been found for this gene.

[provided by RefSeq, Oct 2014]

**Locus ID:** 7355 **MW:** 10