

Product datasheet for **SC204739**

ALG14 (NM_144988) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	ALG14 (NM_144988) Human 3' UTR Clone
Symbol:	ALG14
Synonyms:	CMS15; IDDEBF; MEPCA
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_144988
Insert Size:	2000 bp



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Insert Sequence: >SC204739 3'UTR clone of NM_144988
 The sequence shown below is from the reference sequence of NM_144988. The complete sequence of this clone may contain minor differences, such as SNPs.
 Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
AAATCGGTGTACCTTGGGCGAATTGTTTGACAAAATGGCAACTGACTTCTTTAGAATTTTGCAGTTAACA
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GAATTATTGATGGTAAGGAATAAAAAATGTACAGATGACTCAGTGAAGAACTGAGGCTTCTTTATGA
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AATATGTATTACTACCTGCAAATTTCTTCTGGCTGTTTTAGTAGTATTTTTTTTACAGAACTAAATAT
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TTCTGCATTGTAAAAGAAGAGAGAAAAAAGTATTTCTGAGTTTTGGTTCAGATACAAAAGGCTTTATT
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GCCTGGGGCAGCCAAATGATAGTTTATTGTTGCTTTGATTTAATCCTCAAATCTCAGTCACTGGTTGCT
TCCCTGGAGTATCAGAGCGTATTAGTATCATAACCAGGGGCTGAGGTGAAATAAGCCTTGACTGTTT
TAGAAAGAAAGTATTTTTTTCAGGGACAACAGCATCTGCATTTTATTGCTGTTTAAAGCCTTCTCTGC
ACGCGT AAGCGGCCGCGGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
  
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Restriction Sites: Sgfl-MluI

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: [NM_144988.4](#)

Summary: This gene is a member of the glycosyltransferase 1 family. The encoded protein and ALG13 are thought to be subunits of UDP-GlcNAc transferase, which catalyzes the first two committed steps in endoplasmic reticulum N-linked glycosylation. Mutations in this gene have been linked to congenital myasthenic syndrome (CMSWTA). Alternatively spliced transcript variants have been identified. [provided by RefSeq, Mar 2015]

Locus ID: 199857

MW: 77.3