

Product datasheet for **SC332521**

GLT28D1 (ALG13) (NM_001257231) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	GLT28D1 (ALG13) (NM_001257231) Human Untagged Clone
Tag:	Tag Free
Symbol:	ALG13
Synonyms:	CDG1S; CXorf45; DEE36; EIEE36; GLT28D1; MDS031; TDRD13; YGL047W
Vector:	pCMV6-Entry (PS100001)



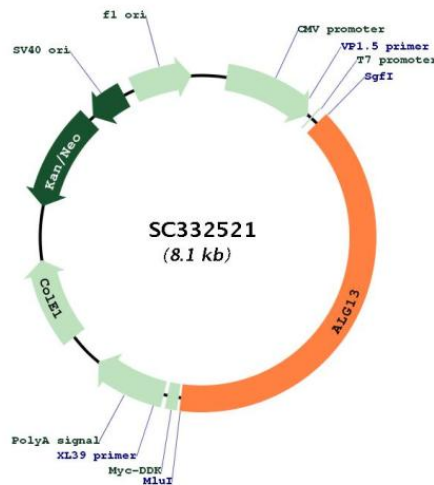
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Fully Sequenced ORF: >SC332521 representing NM_001257231.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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 GATTATTCTGTGTTCCCCCTGGCATCCAGTTGGTACAGCATATGGTGGTTCTTCTCAAATTCATGGT
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 ATGTAA

Restriction Sites: SgfI-MluI

Plasmid Map:



ACCN: NM_001257231

Insert Size: 3180 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_001257231.1](#)

RefSeq Size: 4193 bp

RefSeq ORF: 3180 bp

Locus ID: 79868

UniProt ID: [Q9NP73](#)

Cytogenetics: Xq23

Protein Pathways: Metabolic pathways, N-Glycan biosynthesis

MW: 117.5 kDa

Gene Summary: The protein encoded by this gene is a subunit of a bipartite UDP-N-acetylglucosamine transferase. It heterodimerizes with asparagine-linked glycosylation 14 homolog to form a functional UDP-GlcNAc glycosyltransferase that catalyzes the second sugar addition of the highly conserved oligosaccharide precursor in endoplasmic reticulum N-linked glycosylation. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Dec 2009]

Transcript Variant: This variant (7) differs in the 5' UTR, initiates translation at an alternate start codon and uses an alternate splice site in the 5' coding region, but maintains the reading frame, compared to variant 1. The encoded isoform (6) is shorter and has a distinct N-terminus, compared to isoform 1.