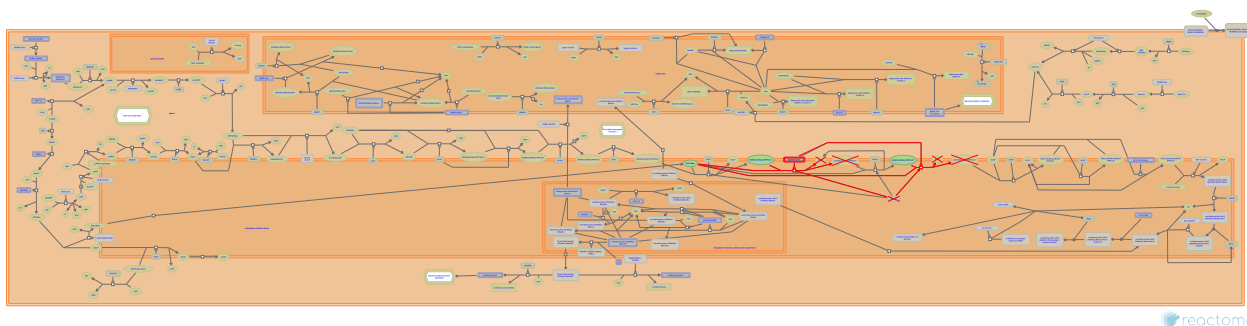


# Defective ALG9 causes CDG-11



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This is just an excerpt of a full-length report for this pathway. To access the complete report, please download it at the [Reactome Textbook](https://reactome.org/page/about-us).

14/05/2024

## Introduction

Reactome is open-source, open access, manually curated and peer-reviewed pathway database. Pathway annotations are authored by expert biologists, in collaboration with Reactome editorial staff and cross-referenced to many bioinformatics databases. A system of evidence tracking ensures that all assertions are backed up by the primary literature. Reactome is used by clinicians, geneticists, genomics researchers, and molecular biologists to interpret the results of high-throughput experimental studies, by bioinformaticians seeking to develop novel algorithms for mining knowledge from genomic studies, and by systems biologists building predictive models of normal and disease variant pathways.

The development of Reactome is supported by grants from the US National Institutes of Health (P41 HG003751), University of Toronto (CFREF Medicine by Design), European Union (EU STRP, EMI-CD), and the European Molecular Biology Laboratory (EBI Industry program).

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- Sidiropoulos, K., Viteri, G., Sevilla, C., Jupe, S., Webber, M., Orlic-Milacic, M. et al. (2017). Reactome enhanced pathway visualization. *Bioinformatics*, 33, 3461-3467. [↗](#)
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- Fabregat, A., Korninger, F., Viteri, G., Sidiropoulos, K., Marin-Garcia, P., Ping, P. et al. (2018). Reactome graph database: Efficient access to complex pathway data. *PLoS computational biology*, 14, e1005968. [↗](#)

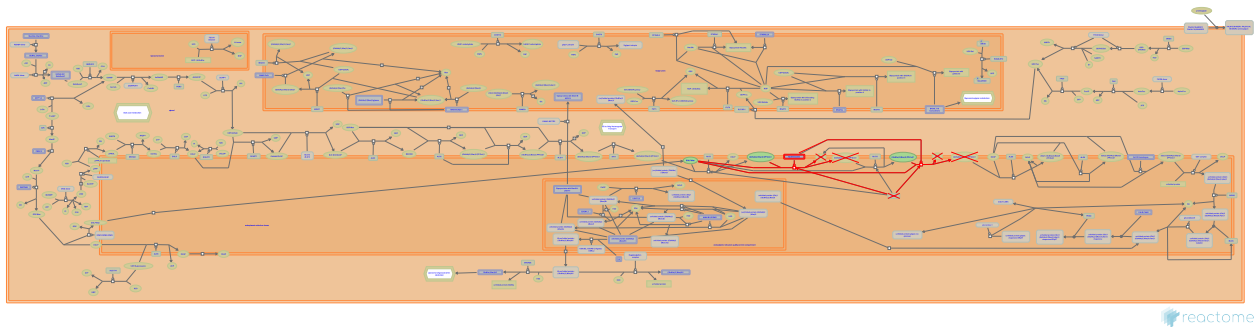
Reactome database release: 88

This document contains 1 pathway and 2 reactions ([see Table of Contents](#))

Defective ALG9 causes CDG-11 ↗

Stable identifier: R-HSA-4720454

Diseases: congenital disorder of glycosylation type I



Alpha-1,2-mannosyltransferase ALG9 (ALG9) normally catalyses the transfer of mannose to the lipid-linked oligosaccharide (LLO) precursor. It adds the 7th and 9th mannose moieties to LLO. Defects in ALG9 are associated with congenital disorder of glycosylation 11 (ALG9-CDG, CDG11; MIM:608776), a multisystem disorder caused by a defect in glycoprotein biosynthesis and characterised by under-glycosylated serum glycoproteins. CDG type 1 diseases result in a wide variety of clinical features, such as defects in the nervous system development, psychomotor retardation, dysmorphic features, hypotonia, coagulation disorders, and immunodeficiency (Frank et al. 2004, Weinstein et al. 2005). The LLO profile showed accumulation of (GlcNAc)2 (Man)6 (PP-Dol)1 and (GlcNAc)2 (Man)8 (PP-Dol)1 fragments, suggesting a defect in ALG9 and correlating with the normal function of ALG9 in adding the 7th and 9th mannose moieties (Frank et al. 2004).

Literature references

Hennet, T., Eyaid, W., Frank, CG., Grubenmann, CE., Berger, EG., Aebi, M. (2004). Identification and functional analysis of a defect in the human ALG9 gene: definition of congenital disorder of glycosylation type 1L. *Am J Hum Genet*, 75, 146-50. ↗

Hennet, T., Clarke, JT., Frank, CG., Seargeant, L., Matthijs, G., Neupert, C. et al. (2005). CDG-1L: an infant with a novel mutation in the ALG9 gene and additional phenotypic features. *Am J Med Genet A*, 136, 194-7. ↗

Editions

2013-10-21	Authored, Edited	Jassal, B.
2014-10-31	Reviewed	Belaya, K.

**Defective ALG9 does not add the seventh mannose to the N-glycan precursor** ↗

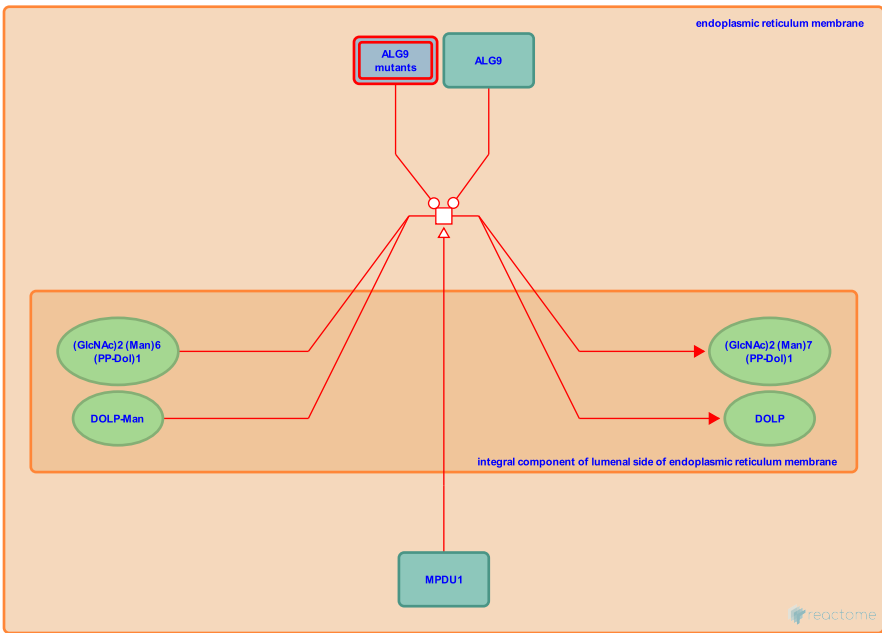
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**Stable identifier:** R-HSA-4720478

**Type:** transition

**Compartments:** endoplasmic reticulum membrane, integral component of luminal side of endoplasmic reticulum membrane

**Diseases:** congenital disorder of glycosylation type I



Alpha-1,2-mannosyltransferase ALG9 (ALG9) normally catalyses the transfer of mannose to the lipid-linked oligosaccharide (LLO) precursor. It adds the 7th and 9th mannose moieties to LLO. Defects in ALG9 are associated with congenital disorder of glycosylation 1I (ALG9-CDG, CDG1I; MIM:608776), a multisystem disorder caused by a defect in glycoprotein biosynthesis and characterised by under-glycosylated serum glycoproteins. CDG type 1 diseases result in a wide variety of clinical features, such as defects in the nervous system development, psychomotor retardation, dysmorphic features, hypotonia, coagulation disorders, and immunodeficiency (Frank et al. 2004, Weinstein et al. 2005). The LLO profile showed accumulation of (GlcNAc)2 (Man)6 (PP-Dol)1 and (GlcNAc)2 (Man)8 (PP-Dol)1 fragments, suggesting a defect in ALG9 and correlating with the normal function of ALG9 in adding the 7th and 9th mannose moieties (Frank et al. 2004). Point mutations that can cause ALG9-CDG are E523K and Y286C (Frank et al. 2004, Weinstein et al. 2005).

**Literature references**

Hennet, T., Eyaid, W., Frank, CG., Grubenmann, CE., Berger, EG., Aebi, M. (2004). Identification and functional analysis of a defect in the human ALG9 gene: definition of congenital disorder of glycosylation type 1I. *Am J Hum Genet*, 75, 146-50. ↗

Hennet, T., Clarke, JT., Frank, CG., Seargeant, L., Matthijs, G., Neupert, C. et al. (2005). CDG-1I: an infant with a novel mutation in the ALG9 gene and additional phenotypic features. *Am J Med Genet A*, 136, 194-7. ↗

**Editions**

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2014-10-31	Reviewed	Belaya, K.

# Defective ALG9 does not add the last mannose to the N-glycan precursor ↗

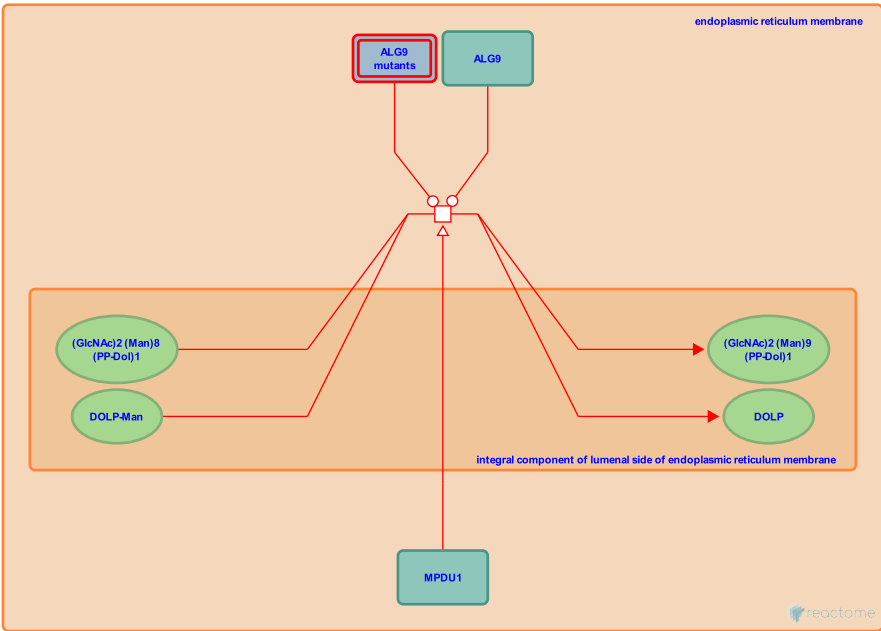
**Location:** Defective ALG9 causes CDG-1I

**Stable identifier:** R-HSA-9035514

**Type:** transition

**Compartments:** endoplasmic reticulum membrane, integral component of luminal side of endoplasmic reticulum membrane

**Diseases:** congenital disorder of glycosylation type I



Alpha-1,2-mannosyltransferase ALG9 (ALG9) normally catalyses the transfer of mannose to the lipid-linked oligosaccharide (LLO) precursor. It adds the 7th and 9th mannose moieties to LLO. Defects in ALG9 are associated with congenital disorder of glycosylation 1I (ALG9-CDG, CDG1I; MIM:608776), a multisystem disorder caused by a defect in glycoprotein biosynthesis and characterised by under-glycosylated serum glycoproteins. CDG type 1 diseases result in a wide variety of clinical features, such as defects in the nervous system development, psychomotor retardation, dysmorphic features, hypotonia, coagulation disorders, and immunodeficiency (Frank et al. 2004, Weinstein et al. 2005). The LLO profile showed accumulation of (GlcNAc)2 (Man)6 (PP-Dol)1 and (GlcNAc)2 (Man)8 (PP-Dol)1 fragments, suggesting a defect in ALG9 and correlating with the normal function of ALG9 in adding the 7th and 9th mannose moieties (Frank et al. 2004). Point mutations that can cause ALG9-CDG are E523K and Y286C (Frank et al. 2004, Weinstein et al. 2005).

## Literature references

Hennet, T., Eyaid, W., Frank, CG., Grubenmann, CE., Berger, EG., Aebi, M. (2004). Identification and functional analysis of a defect in the human ALG9 gene: definition of congenital disorder of glycosylation type 1I. *Am J Hum Genet*, 75, 146-50. ↗

Hennet, T., Clarke, JT., Frank, CG., Seargeant, L., Matthijs, G., Neupert, C. et al. (2005). CDG-1I: an infant with a novel mutation in the ALG9 gene and additional phenotypic features. *Am J Med Genet A*, 136, 194-7. ↗

## Editions

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