

OMIM Entry - # 613612 - CONGENITAL DISORDER OF ...<https://www.omim.org/entry/613612> ▾

Jul 07, 2020 · In an Iraqi girl with a congenital disorder of glycosylation, Paesold-Burda et al. (2009) identified a homozygous intronic substitution (606821.0001) leading to exon skipping and severely...

Congenital disorders of glycosylation (CDG): It's (nearly ...<https://www.researchgate.net/publication/50305323...>

Congenital disorder of glycosylation type 1a (CDG-1a, OMIM #212065) is the most common type in this group, also known as phosphomannomutase 2 (PMM2) deficiency (OMIM ...

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Congenital disorders of glycosylation | Request PDF<https://www.researchgate.net/publication/329968096...>

Request PDF | Congenital disorders of glycosylation | Congenital disorders of glycosylation are a genetically and clinically heterogeneous group of >130 diseases caused by defects in various steps ...

(PDF) Vascular ring anomaly in a patient with ...<https://www.researchgate.net/publication/343941738...>

Herein we report for the first time on a baby with congenital disorder of glycosylation type 1a with atrial septal defect and make a comparison of changes in atrial septal defect by follow-ups to ...

Mutations in PMM2 that cause congenital disorders of ...<https://www.researchgate.net/publication/12267798...>

Congenital disorder of glycosylation type 1a (CDG-1a, recently named PMM2-CDG) is an inherited autosomal recessive rare disease caused by mutations in the phosphomannomutase 2 (PMM2) gene ...

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(PDF) A compound heterozygous mutation in DPAGT1 ...<https://www.researchgate.net/publication/233948658...>

Dec 19, 2012 · A compound heterozygous mutation in DPAGT1 results in a congenital disorder of glycosylation with a relatively mild phenotype.pdf Content uploaded by Shahzad Mohsin Author content

Neonatal presentation of COG6-CDG with prominent skin

Congenital Disorders Of Glycosylation Center ...

<https://allmedx.com/allmedicine/rare+diseases/...>

Jul 17, 2020 · Feb 14th, 2020 - The **ATP6AP1 gene** encodes for ATPase H⁺ transporting protein.

ATP6AP1 gene mutations are associated with congenital disorders of glycosylation (CDG) and can affect multiple organ system. Descriptions of postnatal phenotype include immunodeficiency, hepatopathy and cognitive impairment.

Diagnostic Approach to Acute Liver Failure in Children: A ...

<https://www.sciencedirect.com/science/article/pii/S1590865821000979>

May 01, 2021 · **Congenital disorders of glycosylation (CDG)** are multiorgan diseases, with variable degree of liver involvement in about 20% of types. Some CDGs have a predominant or isolated liver involvement (eg. MPI-CDG, CCDC115-CDG, TMEM199-CDG and ATP6AP1-CDG) and may present with ALF in ...

Cited by: 1

Author: A. Di Giorgio, E. Bartolini, P.L. Calvo, M. Can...

Publish Year: 2021

Glyco25, XXV International Symposium on Glycoconjugates ...

<https://link.springer.com/article/10.1007/s10719-019-09880-4>

Jul 09, 2019 · **Congenital defects of glycosylation (CDG)** are inherited diseases due to hypoglycosylation of glycoproteins and glycolipids, and to defective biosynthesis of glycosylphosphatidylinositol anchors. The first clinical report appeared in 1980. Since then, the number of published CDG has shown an exponential expansion. Hundred thirty-five CDG are known.

IJMS | Free Full-Text | The V-ATPase a3 Subunit: Structure ...

Name of Journal: *World Journal of Clinical Cases*

Manuscript NO: 66540

Manuscript Type: CASE REPORT

Congenital disorder of glycosylation caused by mutation of ATP6AP1 gene (c.1036G>A) in a Chinese infant: A case report

Yang X *et al.* CDG with ATP6AP1 gene mutation

Abstract

BACKGROUND

The ATP6AP1 gene codes for the accessory protein Ac45 of the vacuolar-type adenosine triphosphatases (V-ATPase) is located on chromosome Xq28. Defects in certain subunits or accessory subunits of the V-ATPase can lead to congenital disorders of glycosylation (CDG). CDG is a group of metabolic disorders in which defective protein and lipid glucosylation processes affect multiple tissues and organs. Therefore, the clinical

Match Overview

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Congenital disorder of glycosylation caused by mutation of ATP6A



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ClinVar archives and aggregates information about relationships among variation and human health.

NM_000249.3(MLH1):c.116+5G>C AND Hereditary cancer ...

<https://www.ncbi.nlm.nih.gov/clinvar/RCV000776333> ▾

ClinVar archives and aggregates information about relationships among variation and human health.

Congenital Disorders Of Glycosylation Center ...

<https://allmedx.com/allmedicine/rare+diseases/> ▾

Jul 17, 2020 - Apr 15th, 2020 - Congenital disorders of glycosylation are a growing group of rare genetic disorders caused by deficient protein and lipid glycosylation. Here, we report the clinical, biochemical, and molecular features of seven patients from four families with GALNT2-congenital disorder of glycosylation (GALNT2-CDG), an O-linked glycosylation ...

rs756898983 RefSNP Report - dbSNP - NCBI

<https://www.ncbi.nlm.nih.gov/snp/rs756898983> ▾

Apr 21, 2020 - Welcome to the Reference SNP (rs) Report. All alleles are reported in the Forward orientation. Click on the Variant Details tab for details on Genomic Placement, Gene, and Amino Acid changes. HGVS names are in the HGVS tab.

www.science.gov

www.science.gov/topicpages/a/autosomal+dominant+pkd.html ▾

May 31, 2018 - Comprehensive PKD 1 and PKD 2 Mutation Analysis in Prenatal Autosomal Dominant Polycystic Kidney Disease. PubMed. Audr  zet, Marie-Pierre; Corbiere, Christine; Lebbah, Said; Mor

List of variants in gene IQCB1 studied for abdominal and ...

<https://clinvarminer.genetics.utah.edu/.../gene/IQCB1> ▾

List of variants in gene IQCB1 studied for abdominal and pelvic region disorder Included ClinVar conditions (1028): 3 beta-Hydroxysteroid dehydrogenase deficiency

List of variants in gene TYR reported as pathogenic