strategies, being tested and used in these disorders. NA – Not available.	Supplementary Table 1 - Overview of	of the ongoing clinical trials ((CTs), available CDG in	<i>vitro</i> and <i>in vivo</i> model	s and biomarkers and d	ietary supplementation
	strategies, being tested and used in t	hese disorders. NA – Not av	vailable.			

	ALG1-CDG Chitobiosyldiphospho- dolichol β- mannosyltransferase	ALG6-CDG α-1,3- glucosyltransferase	ALG13-CDG UDP-GlcNAc transferase	ATP6AP1-CDG Accessory subunit of the vacuolar (V)- ATPase protein pump	CAD-CDG Tri-functional protein (ATase, CPSase, ATCase and DHOase)	CCDC115-CDG Coiled-coil domain- Containing protein 115
CTs	Interventional NCT02955264	Interventional NCT02955264	NA		NA	NA
ls	S. cerevisiae alg1 mutant (K57-6C strain) [24,34,41]	<i>S. cerevisiae alg6</i> mutant [87,88]	<i>S. cerevisiae</i> "Tet-Off " strain	E. cerevisiae t-Off " strain L047W::kanR- et07-TATA S. cerevisiae (voa1::H A3::CMV-tTA vma21QQ strain) a his3-1 leu2-0 [97] met15-0)	<i>CAD</i> -deficient CHO-G9C [100]	S. cerevisiae strains: HY13 (vma22A::LEU2) KHY34 (vma22A::LEU2 pep4-3)
In vitro mode	S. cerevisiae alg1 mutant (PRY56 strain)	voisiae alg1 mutant PRY56 strain) MI8-5 Alg6-/- teol MATa his3-1 1 met15-0)	tet07-TATA URA3::CMV-tTA MATa his3-1 leu2-0 met15-0)			KHY38 (vma22A::URA3) KHY39 (vma22A::URA3 pep4-3) [105]
	[31]		[281]			HeLa (CCDC115 knockdown using CRISPR/Cas9) [106]
odels				Zebrafish	C. elegans pyr-1(cu8) mutant [85]	
In vivo me	NA	NA	NA	(Atp6ap1b ^{a82/a82} ,Atp6ap1b knockdown) [98]	D. melanogaster Rudimentary mutant [86]	NA

Suppleme	entary	Table	2 – co	ontinued.

	ALG1-CDG Chitobiosyldiphospho- dolichol β- mannosyltransferase	ALG6-CDG α-1,3- glucosyltransferase	ALG13-CDG UDP-GlcNAc transferase	ATP6AP1-CDG Accessory subunit of the vacuolar (V)- ATPase protein pump	CAD-CDG Tri-functional protein (ATase, CPSase, ATCase and DHOase)	CCDC115-CDG Coiled-coil domain- Containing protein 115
In vivo models	NA	NA	NA	Chimeric Mouse (w/ <i>Atp6ap1</i> reduced expression) [99]	Zebrafish <i>Perplexed</i> (<i>plx</i> ⁴⁵²) mutant [87] Zebrafish Transgenic <i>Tg</i> (<i>p2xr3.2:gfp</i>) ^{sl23} mutant [88]	NA
ers	GlcNAc2-PP-dolichol [31]	Dolichol-linked ManºGlcNAc2 [177]	GlcNAc-PP-dolichol			
3iomarko	N-tetrasaccharide (Neu₅Ac_2,6Gal_1,4-	AGA [166]	ICAM-1	NA	NA	NA
	GlcNAc_1,4GlcNAc) [34,178]	β-trace protein [180,189]	[190]			
Dietery therapy	Man suppl [178]	NA	Gal suppl [190]	NA	Uridine suppl [69,100]	Iron suppl [106]
Transplantation	NA	NA	NA	Liver transplantation (approved therapy in Europe)	NA	Liver transplantation (approved therapy in Europe) [224]

Suppleme	entary	Table	3 -	continued.

	ALG1-CDG Chitobiosyldiphospho- dolichol β- mannosyltransferase	ALG6-CDG α-1,3- glucosyltransferase	ALG13-CDG UDP-GlcNAc transferase	ATP6AP1-CDG Accessory subunit of the vacuolar (V)- ATPase protein pump	CAD-CDG Tri-functional protein (ATase, CPSase, ATCase and DHOase)	CCDC115-CDG Coiled-coil domain- Containing protein 115
Gene therapy	NA	NA	NA	NA	NA	NA
Antisense therapy	NA	NA	NA	NA	NA	NA
PCs	NA	NA	NA	NA	NA	NA
Others	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256]
	DOLK-CDG Dolichol kinase	GNE-CDG UDP-GlcNAc 2- epimerase/ ManNAc kinase	ISPD-CDG Isoprenoid synthase domain- containing protein	MAGT1-CDG Magnesium transporter 1	MPI-CDG Mannose-6- phosphate isomerase	NANS-CDG CMP-N- acetylneuraminic acid synthetase
CTs	NA	Interventional NCT02346461 NCT01634750 NCT01517880	NA	NA	NA	SA trial approved but discontinued due unfavorable results in GNE-CDG

Suppleme	entary Table 4 - continued.					
	DOLK-CDG Dolichol kinase	GNE-CDG UDP-GlcNAc 2- epimerase/ ManNAc kinase	ISPD-CDG Isoprenoid synthase domain- containing protein	MAGT1-CDG Magnesium transporter 1	MPI-CDG Mannose-6- phosphate isomerase	NANS-CDG CMP-N- acetylneuraminic acid synthetase
CTs	-	Interventional NCT01830972 NCT02377921 NCT02736188 NCT02731690 NCT01359319 NCT00195637 NCT0195637 NCT01236898 NCT02196909 Observational NCT01784679 NCT01902940 NCT01417533	-	-	-	-
In vitro models	<i>S. cerevisiae Sec59</i> mutant [18,29]	CHO Gne-deficient Lec3 mutant [19,26,28,107,108] HEK293 (D176V-Gne, V572L-Gne and Gne knockdown) [39,109] BJA-B K20 (D176V-Gne and M712L-Gne) [110,111,206,207] HL60-I [110,111] Sf9 M712T-Gne [40] Gne ^{-/-} mice ESC [74 112 113 115]	HEK293 <i>Ispd</i> knockout [68] HAP1 <i>Ispd</i> knockout [150]	<i>S. cerevisiae Alr1∆</i> strain [90] HEK293 <i>Magt1</i> knock down [90]	HT-29 <i>Mpi</i> knockdown [59]	NA

Suppleme	entary Table 5 - continued.					
	DOLK-CDG Dolichol kinase	GNE-CDG UDP-GlcNAc 2- epimerase/ ManNAc kinase	ISPD-CDG Isoprenoid synthase domain- containing protein	MAGT1-CDG Magnesium transporter 1	MPI-CDG Mannose-6- phosphate isomerase	NANS-CDG CMP-N- acetylneuraminic acid synthetase
In vivo models		Mouse (<i>Gne^{+/-,} Gne^{+/-}</i>) [74,116] Mouse (<i>Gne</i> ^{M712T/M712T}) [76,81,117]	Mouse (<i>Ispd</i> ^{L79*/L79*}) [151]	Zebrafish (<i>Magt1</i> knock-out) [90]	Mouse (<i>Mpi</i> ^{Y255C/Y255C}) [72]	
	NA	Mouse (Gne ^(-/-) hGNED176V-Tg) [118,119] Mouse (Gne ^{V572L/V572L}) [76,81,117]	Mouse (Knock-out) [152]		Mouse (Mpi≁) [91]	Zebrafish model (<i>nansa</i> and <i>nansb</i> knockdown) [122]
		Mouse (Transgenic FVBN- GNR-R263L) [120]	Zebrafish (Knock-out) [153]	NA	Zebrafish (w/ 13% of <i>mpi</i> activity) [92]	
		Zebrafish (<i>gne</i> knock-out) [121]				
Biomarkers	NA	GM3 and GD3 gangliosides [181,182]	NA		N-tetrasaccharide (Neu5Ac_2,6Gal_1,4- GlcNAc_1,4GlcNAc [178]	
		NCAM [116 183]		NA	AGA [166] ICAM-I [174,175]	NA
		Thomsen-Friedenreich (T)-antigen [184]				

Suppleme	entary Table 6 - continued.					
	DOLK-CDG Dolichol kinase	GNE-CDG UDP-GlcNAc 2- epimerase/ ManNAc kinase	ISPD-CDG Isoprenoid synthase domain- containing protein	MAGT1-CDG Magnesium transporter 1	MPI-CDG Mannose-6- phosphate isomerase	NANS-CDG CMP-N- acetylneuraminic acid synthetase
Diatery therapy	NA	ManNAc suppl [66,80–82,117] ManN suppl [78,83] SA suppl [78,82,83] 6'-sialyllactose suppl [82,84]	Ribitol suppl [68,150]	Mg²⁺ suppl [63,215]	Man suppl [49,160–169]	SA suppl [122]
		Ac4ManNAc suppl [83,207]				
Transplantation	Heart transplantation [252,253]	NA	NA	Hematopoietic cell transplantation [215]	Liver transplantation (approved therapy in Europe) [171]	NA
Gene therapy	NA	AVV8 [61,243] AVV-TS [244] GNE-lipoplex [28,246–248]	NA	NA	NA	NA

Supplem	entary	Table 7	7 – 0	continue	ed.

	DOLK-CDG Dolichol kinase	GNE-CDG UDP-GlcNAc 2- epimerase/ ManNAc kinase	ISPD-CDG Isoprenoid synthase domain-containing protein	MAGT1-CDG Magnesium transporter 1	MPI-CDG Mannose-6- phosphate isomerase	NANS-CDG CMP-N- acetylneuraminic acid synthetase
Antisense therapy	NA	NA	NA	NA	NA	NA
PCs	NA	NA	NA	NA	NA	NA
Others	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256] GNEM-FAS [261] REMUDY [262,263]	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256]
	PGM1-CDG	PGM3-CDG	PIGA-CDG Phosphatidylinositol	PIGM-CDG GPI α-1,4-	PIGO-CDG GPI ethanolamine	PMM2-CDG
	Phosphogluco- mutase 1	Phosphogluco- mutase 3	N-acetylglucosaminyl- transferase (subunit A)	mannosyltransferase I	phosphate transferase 3	Phosphomannomutase 2
CTs	Phosphogluco- mutase 1 Interventional NCT02955264	Phosphogluco- mutase 3 NA	N-acetylglucosaminyl- transferase (subunit A) NA	mannosyltransferase I NA	phosphate transferase 3 NA	Phosphomannomutase 2 Interventional NCT03250728 2017-000810-44 Observational NCT03173300
nodels CTs	Phosphogluco- mutase 1 Interventional NCT02955264	Phosphogluco- mutase 3 NA	N-acetylglucosaminyl- transferase (subunit A) NA hiPSC (hypomorphic	mannosyltransferase I NA Ramos517 cells <i>Pigm</i> - deficient [148]	phosphate transferase 3 NA CHO <i>Pigo</i> -deficient [35,37]	Phosphomannomutase 2 Interventional NCT03250728 2017-000810-44 Observational NCT03173300 iPSC (hypomorphic PMM2 ⁴²² C=A/357C=A and DM 2422C=A/357C=A ard

Supplem	upplementary Table 8 - continued.								
	PGM1-CDG Phosphogluco- mutase 1	PGM3-CDG Phosphogluco- mutase 3	PIGA-CDG Phosphatidylinositol N-acetylglucosaminyl- transferase (subunit A)	PIGM-CDG GPI α-1,4- mannosyltransferase I	PIGO-CDG GPI ethanolamine phosphate transferase 3	PMM2-CDG Phosphomanno- mutase 2			
In vivo models	NA	Mouse (<i>Pgm3^{mld1}</i>) [124] Mouse (<i>Pgm3st</i>) [124]	Chimeric Mouse (<i>Piga</i> -deficient) [146] Mouse (Partial exon2 excision) [147]	NA	NA	Mouse (Knock-out) [75] Mouse (Pmm2 ^{R137H/R137H}) [77] Mouse (Pmm2 ^{F118L/F118L}) [77] Mouse (Pmm2 ^{R137H/F118L}) [77] Mouse (Pmm2 ^{R137H/F115L}) [94] Zebrafish (Knock-out) [95] Drosophila melanogaster (pmm2-null) [96] D.melanogaster (pmm2 knockdown) [96] Xenopus laevis (Pmm2-null) [282]			
Biomarkers	NA	NA	NA	NA	NA	N-tetrasaccharide (Neu5Ac_2,6Gal_1,4- GlcNAc_1,4GlcNAc) [178]			

Supplementary Table 9 - continue

	PGM1-CDG Phosphogluco- mutase 1	PGM3-CDG Phosphogluco- mutase 3	PIGA-CDG Phosphatidylinositol N-acetylglucosaminyl- transferase (subunit A)	PIGM-CDG GPI α-1,4- mannosyltransferase I	PIGO-CDG GPI ethanolamine phosphate transferase 3	PMM2-CDG Phosphomanno- mutase 2
Biomarkers	NA	NA	NA	NA	NA	Band 3 and glycophorin A [185] Glycosphingolipids (Gb3, GM2, GD3 and GD1a) [186] AGA [166] ICAM-I [174,175] α_1 -acid glycoprotein [155,187] Ceruloplasmin [155,187] α_1 -antichymotrypsin [155,187] α_1 B-glycoprotein [155,187]
tery therapy	Gal suppl [65,208,209] Uridine suppl [209,210] Glucose IV	GlcNAc Suppl [67]	Ketogenic diet [216]	Sodium Phenylbutyrate suppl [57]	Vitamine B6 suppl [220]	Man (Man-1-P) suppl http://glycomine.com/ [46,47,52,77,194,202]
Die	Glucose IV administration [211]					Glc starvation [194]

	PGM1-CDG Phosphogluco- mutase 1	PGM3-CDG Phosphogluco- mutase 3	PIGA-CDG Phosphatidylinositol N-acetylglucosaminyl- transferase (subunit A)	PIGM-CDG GPI α-1,4- mannosyltransferase I	PIGO-CDG GPI ethanolamine phosphate transferase 3	PMM2-CDG Phosphomanno- mutase 2
Transplantation	Heart transplantation [65]	Hematopoietic cell transplantation [42]	NA NA NA		NA	
Gene therapy	NA	NA	NA	NA	NA	NA
Antisense therapy	NA	NA	NA	NA	NA	AMO (c.640-15479C>T) [239]
PCs	NA	NA	NA	NA	NA	1-(3-chlorophenyl)-3-3- bis(pyridine-2-yl)urea [231]
Others	NPCRS [255,256] TPCRS [260]	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256]	NPCRS [255,256]	Metformin [200] MPI inhibitors [59,237] NPCRS [255–258] ICARS [257,258]

	SLC35A1-CDG CMP-sialic acid transporter	SLC35A2-CDG UDP-galactose transporter	SLC35C1-CDG GDP-fucose transporter	SLC39A8-CDG Solute carrier family 39 (zinc transporter), member 8	SRD5A3-CDG Steroid 5α- reductase type 3	TMEM165-CDG Transmembrane protein 165
CTs	NA	NA	NA	Interventional NCT02955264 Mn²+ trial (registered)	Interventional NCT02955264	Interventional NCT02955264
els	CHO Lec 2 mutants [17,25,26,38,125]	CHO Lec8 mutant [33]	CHO <i>Slc35c1</i> knockout [130]			S. cerevisiae Gdt1∆ [30,141,142]
i <i>tro</i> mod	CHO MAR-11 mutant [126]	CHO Loce mutant [22]	CHO <i>Slc35c1</i> knockout derived from MAR-11 mutants [128,129]	NA	S.cerevisiae Dfg10-100 [140]	HEK293 <i>TMEM165</i> knockout and knockdown [36,142]
In vi	HAP1 <i>Slc35a1</i> knockout [32]	CHO Lets mutant [33]	ESC <i>Slc35c1</i> knockout [131]		[2.00]	HeLa <i>TMEM165</i> knockdown [142]
In vivo models	NA	C. elegans Srf-1 JA mutants [127]	Mouse (<i>Slc35c1-⁺</i>) [132,133]	Mouse (Slc39a8 ^(neo/neo)) [138 139]		
			Zebrafish (<i>slytherin</i>) [135,136]	[100,107]	Mouse (<i>Srd5a3^{Gt/Gt}</i>) [140]	Zebrafish (<i>tmem165-^{,,}</i>) [143]
			overexpression) [137]	Mouse (Slc39a8+) [85]		
Biomarkers	NA	NA	NA	NA	NA	NA

Supplementary	Table 12 - continued.
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	SLC35A1-CDG CMP-sialic acid transporter	SLC35A2-CDG UDP-galactose transporter	SLC35C1-CDG GDP-fucose transporter	SLC39A8-CDG Solute carrier family 39 (zinc transporter), member 8	SRD5A3-CDG Steroid 5α- reductase type 3	TMEM165-CDG Transmembrane protein 165
herapy	SA, ManNAc or the fetuin suppl	Col suppl	Fue suppl	Mn ²⁺ suppl [226,228] Uridine suppl		Mn ²⁺ suppl [70,142]
Diatery ti	(all w/o beneficial effects) [71]	[33,221]	[23,27,50,51,53,132,136,222]	[226,227] Gal suppl [226,227]	NA	Gal suppl [70]
Transplantation	NA	NA	NA	NA	NA	NA
Gene therapy	NA	NA	NA	NA	NA	NA
Antisense therapy	NA	NA	NA	NA	NA	AMO (c.792+182G>A) [238]
PCs	NA	NA	NA	NA	NA	NA

Supplementary Table 13 – continued	Supp	lementary	Table	13 –	continued
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	SLC35A1-CDG CMP-sialic acid transporter	DG SLC35A2-CDG SLC35C1-CDG acid UDP-galactose transporter GDP-fucose transporter		SLC39A8-CDG Solute carrier family 39 (zinc transporter), member 8	SRD5A3-CDG Steroid 5α- reductase type 3	TMEM165-CDG Transmembrane protein 165
Others	NPCRS	NPCRS	NPCRS	NPCRS	NPCRS	NPCRS
	[255,256]	[255,256]	[255,256]	[255,256]	[255,256]	[255,256]

Supplementary Table 14 – continued.

		TMEM199-CDG Transmembrane protein 199								
CTs	NA	In vivo models	NA	Antisense therapy	NA	PCs	NA			
ro models	DJY62/DJY102 (pep4–3 vma12 Δ::LEU2) DJY63 (vma12 Δ::LEU2) [105,144]	Trans-plantation	Biomarkers		NA	Others	NPCRS [255,256]			
In vit	HeLa cell line (TMEM199 knowdown by CRISPR/Cas9) [106]	Gene therapy	NA	Diatery therapy	Iron supp [106]					



Figure S1 – Drug development process overview.

								-
Animal models	Liver transplant	Heart transplantation	N-acetyl-D- mannosamine (MapNAC)		Congenital disorders of glycosylation	ALG1	ALG1-CDG	ALG6
Yeast	Heart transplant	Transplantation	Mannosamine (MapN)		ALG6-CDG	ALG12	ALG12-CDG	ALG13
Drosophila melanogaster	Bone marrow	Stem cell	Ribitol		ALG13-CDG	ATP6VAP1	ATP6VAP1-CDG	CAD
Zebrafish	Transplant	Oral supplementation	Magnesium (Mg2+)	王	Carbohydrate deficient glycoprotein	CAD-CDG	CCDC115	CCDC115-CDG
Mouse	Pharmacological chaperones	Mannose	N-Acetyi glucosamine (GlcNAc)	.IM N	COG5	COG5-CDG	DOLK	DOLK-CDG
Rat	Chaperones	Galactose	Phenylbutyrate	VIIO	GNE	GNE-CDG	ISPD	ISPD-CDG
Clinical trials	Antisense therapy	Fucose	Guanosine diphosphate	3IN/	MAGT1	MAGT1-CDG	MPI	MPI-CDG
Therapies	Man-1-P	Sialic Acid	Uridine-5'- Diphosphate	WO	NANS	NANS-CDG	PGM1	PGM1-CDG
C.elegans	Man-1-P therapy	Manganese	L-Aspartic acid	N N	PGM3	PGM3-CDG	PIGM	PIGM-CDG
Therapy	Splicing	Uridine	L-Glutamine	1	PMM2	PMM2-CDG	SL39A8	SL39A8-CDG
Therapeutic	Induced pluripotent stem	Metformin	Famotidine		SLC35A2	SLC35A2-CDG	SLC35C1	SLC35C1-CDG
strategies	cells			-	SRD5A3	SRD5A3-CDG	TMEM165	TMEM165-CDG
options	Biomarkers	Acetazolamide	Phosphoric acid		PIGA	PIGA-CDG	PIGO	PIGO-CDG

Figure S2 – List of all keywords used for this systematic literature review.



Figure S3 – Diagram of the inclusion/elimination process used for manuscript selection.