

Francois Foulquier

List of Publications by Year in descending order

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73
papers

3,626
citations

126901

33
h-index

138468

58
g-index

76
all docs

76
docs citations

76
times ranked

3800
citing authors

#	ARTICLE	IF	CITATIONS
1	SLC10A7, an orphan member of the SLC10 family involved in congenital disorders of glycosylation. <i>Human Genetics</i> , 2022, 141, 1287-1298.	3.8	3
2	CAMLG-CDG: a novel congenital disorder of glycosylation linked to defective membrane trafficking. <i>Human Molecular Genetics</i> , 2022, , .	2.9	7
3	Towards understanding the extensive diversity of protein N-glycan structures in eukaryotes. <i>Biological Reviews</i> , 2022, 97, 732-748.	10.4	14
4	TMEM165 a new player in proteoglycan synthesis: loss of TMEM165 impairs elongation of chondroitin- and heparan-sulfate glycosaminoglycan chains of proteoglycans and triggers early chondrocyte differentiation and hypertrophy. <i>Cell Death and Disease</i> , 2022, 13, 11.	6.3	13
5	Differential Effects of D-Galactose Supplementation on Golgi Glycosylation Defects in TMEM165 Deficiency. <i>Frontiers in Cell and Developmental Biology</i> , 2022, 10, .	3.7	3
6	Variation of the serum N-glycosylation during the pregnancy of a MPI-CDG patient. <i>JIMD Reports</i> , 2021, 62, 22-29.	1.5	1
7	Fetal bovine serum impacts the observed N-glycosylation defects in TMEM165 KO HEK cells. <i>Journal of Inherited Metabolic Disease</i> , 2020, 43, 357-366.	3.6	11
8	Serum bikunin isoforms in congenital disorders of glycosylation and linkeropathies. <i>Journal of Inherited Metabolic Disease</i> , 2020, 43, 1349-1359.	3.6	12
9	SPCA1 governs the stability of TMEM165 in Hailey-Hailey disease. <i>Biochimie</i> , 2020, 174, 159-170.	2.6	6
10	Biometals and glycosylation in humans: Congenital disorders of glycosylation shed lights into the crucial role of Golgi manganese homeostasis. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2020, 1864, 129674.	2.4	25
11	Novel role for the Golgi membrane protein TMEM165 in control of migration and invasion for breast carcinoma. <i>Oncotarget</i> , 2020, 11, 2747-2762.	1.8	11
12	Dissection of TMEM165 function in Golgi glycosylation and its Mn ²⁺ sensitivity. <i>Biochimie</i> , 2019, 165, 123-130.	2.6	22
13	Mutations in MAGT1 lead to a glycosylation disorder with a variable phenotype. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 9865-9870.	7.1	66
14	Involvement of thapsigargin- and cyclopiazonic acid-sensitive pumps in the rescue of TMEM165-associated glycosylation defects by Mn ²⁺ . <i>FASEB Journal</i> , 2019, 33, 2669-2679.	0.5	21
15	Investigating the functional link between TMEM165 and SPCA1. <i>Biochemical Journal</i> , 2019, 476, 3281-3293.	3.7	12
16	Chemical glycomics enrichment: imaging the recycling of sialic acid in living cells. <i>Journal of Inherited Metabolic Disease</i> , 2018, 41, 515-523.	3.6	12
17	Congenital disorders of glycosylation (CDG): Quo vadis?. <i>European Journal of Medical Genetics</i> , 2018, 61, 643-663.	1.3	191
18	Hypothesis: lobe A (COG1 ⁴)-CDG causes a more severe phenotype than lobe B (COG5 ⁸)-CDG. <i>Journal of Medical Genetics</i> , 2018, 55, 137-142.	3.2	14

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19	Expanding the phenotype of metabolic cutis laxa with an additional disorder of N-linked protein glycosylation. <i>European Journal of Human Genetics</i> , 2018, 26, 618-621.	2.8	12
20	Investigating the function of Gdt1p in yeast Golgi glycosylation. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2018, 1862, 394-402.	2.4	29
21	Use of Endoglycosidase H as a diagnostic tool for MAN1B1â€CDG patients. <i>Electrophoresis</i> , 2018, 39, 3133-3141.	2.4	9
22	Protein N-glycosylation alteration and glycolysis inhibition both contribute to the antiproliferative action of 2-deoxyglucose in breast cancer cells. <i>Breast Cancer Research and Treatment</i> , 2018, 171, 581-591.	2.5	30
23	The extended cytoplasmic tail of the human B4GALNT2 is critical for its Golgi targeting and postâ€Golgi sorting. <i>FEBS Journal</i> , 2018, 285, 3442-3463.	4.7	10
24	Integrating glycomics and genomics uncovers SLC10A7 as essential factor for bone mineralization by regulating post-Golgi protein transport and glycosylation. <i>Human Molecular Genetics</i> , 2018, 27, 3029-3045.	2.9	37
25	TMEM165 deficiencies in Congenital Disorders of Glycosylation type II (CDG-II): Clues and evidences for roles of the protein in Golgi functions and ion homeostasis. <i>Tissue and Cell</i> , 2017, 49, 150-156.	2.2	34
26	Evidence for splice transcript variants of TMEM165, a gene involved in CDG. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2017, 1861, 737-748.	2.4	8
27	Manganese-induced turnover of TMEM165. <i>Biochemical Journal</i> , 2017, 474, 1481-1493.	3.7	44
28	Galactose Supplementation in Patients With TMEM165-CDG Rescues the Glycosylation Defects. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017, 102, 1375-1386.	3.6	61
29	Mutations in <i>TRAPPC11</i> are associated with a congenital disorder of glycosylation. <i>Human Mutation</i> , 2017, 38, 148-151.	2.5	34
30	Mutations in the X-linked <i>ATP6AP2</i> cause a glycosylation disorder with autophagic defects. <i>Journal of Experimental Medicine</i> , 2017, 214, 3707-3729.	8.5	62
31	Yeast Gdt1 is a Golgi-localized calcium transporter required for stress-induced calcium signaling and protein glycosylation. <i>Scientific Reports</i> , 2016, 6, 24282.	3.3	48
32	Glycosylation abnormalities in Gdt1p/TMEM165 deficient cells result from a defect in Golgi manganese homeostasis. <i>Human Molecular Genetics</i> , 2016, 25, 1489-1500.	2.9	92
33	CCDC115 Deficiency Causes a Disorder of Golgi Homeostasis with Abnormal Protein Glycosylation. <i>American Journal of Human Genetics</i> , 2016, 98, 310-321.	6.2	88
34	TMEM199 Deficiency Is a Disorder of Golgi Homeostasis Characterized by Elevated Aminotransferases, Alkaline Phosphatase, and Cholesterol and Abnormal Glycosylation. <i>American Journal of Human Genetics</i> , 2016, 98, 322-330.	6.2	73
35	ALG11-CDG: Three novel mutations and further characterization of the phenotype. <i>Molecular Genetics and Metabolism Reports</i> , 2015, 2, 16-19.	1.1	14
36	Abnormal cartilage development and altered N-glycosylation in Tmem165-deficient zebrafish mirrors the phenotypes associated with TMEM165-CDG. <i>Glycobiology</i> , 2015, 25, 669-682.	2.5	29

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37	Golgi post-translational modifications and associated diseases. <i>Journal of Inherited Metabolic Disease</i> , 2015, 38, 741-751.	3.6	28
38	Cohen syndrome is associated with major glycosylation defects. <i>Human Molecular Genetics</i> , 2014, 23, 2391-2399.	2.9	79
39	Assessing ER and Golgi N-Glycosylation Process Using Metabolic Labeling in Mammalian Cultured Cells. <i>Methods in Cell Biology</i> , 2013, 118, 157-176.	1.1	7
40	Alkynyl monosaccharide analogues as a tool for evaluating Golgi glycosylation efficiency: application to Congenital Disorders of Glycosylation (CDG). <i>Chemical Communications</i> , 2013, 49, 11293.	4.1	24
41	Glycosylation disorders of membrane trafficking. <i>Glycoconjugate Journal</i> , 2013, 30, 23-31.	2.7	53
42	Newly characterized Golgi-localized family of proteins is involved in calcium and pH homeostasis in yeast and human cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6859-6864.	7.1	129
43	MAN1B1 Deficiency: An Unexpected CDG-II. <i>PLoS Genetics</i> , 2013, 9, e1003989.	3.5	63
44	Impact of disease-causing mutations on TMEM165 subcellular localization, a recently identified protein involved in CDG-II. <i>Human Molecular Genetics</i> , 2013, 22, 2914-2928.	2.9	39
45	Insulin signaling controls the expression of O-GlcNAc transferase and its interaction with lipid microdomains. <i>FASEB Journal</i> , 2013, 27, 3478-3486.	0.5	43
46	PUGNAc treatment leads to an unusual accumulation of free oligosaccharides in CHO cells. <i>Journal of Biochemistry</i> , 2012, 151, 439-446.	1.7	20
47	DPM2-CDG: A muscular dystrophy-dystroglycanopathy syndrome with severe epilepsy. <i>Annals of Neurology</i> , 2012, 72, 550-558.	5.3	121
48	TMEM165 Deficiency Causes a Congenital Disorder of Glycosylation. <i>American Journal of Human Genetics</i> , 2012, 91, 15-26.	6.2	162
49	Characterization of O-GlcNAc cycling and proteomic identification of differentially O-GlcNAcylated proteins during G1/S transition. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2012, 1820, 1839-1848.	2.4	56
50	COG5-CDG: expanding the clinical spectrum. <i>Orphanet Journal of Rare Diseases</i> , 2012, 7, 94.	2.7	36
51	Identification of phosphorylated oligosaccharides in cells of patients with a congenital disorders of glycosylation (CDG-I). <i>Biochimie</i> , 2011, 93, 823-833.	2.6	23
52	How Golgi glycosylation meets and needs trafficking: the case of the COG complex. <i>Glycobiology</i> , 2011, 21, 853-863.	2.5	75
53	Overexpression of Man2C1 leads to protein underglycosylation and upregulation of endoplasmic reticulum-associated degradation pathway. <i>Glycobiology</i> , 2011, 21, 363-375.	2.5	22
54	Differential effects of lobe A and lobe B of the Conserved Oligomeric Golgi complex on the stability of β 1,4-galactosyltransferase 1 and β 2,6-sialyltransferase 1. <i>Glycobiology</i> , 2011, 21, 864-876.	2.5	33

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55	Quality control of glycoproteins bearing truncated glycans in an ALC9-defective (CDG-IL) patient. <i>Glycobiology</i> , 2009, 19, 910-917.	2.5	20
56	Golgi function and dysfunction in the first COG4-deficient CDG type II patient. <i>Human Molecular Genetics</i> , 2009, 18, 3244-3256.	2.9	129
57	Deficiency in COG5 causes a moderate form of congenital disorders of glycosylation. <i>Human Molecular Genetics</i> , 2009, 18, 4350-4356.	2.9	104
58	RFT1 deficiency in three novel CDG patients. <i>Human Mutation</i> , 2009, 30, 1428-1434.	2.5	25
59	A new mutation in COG7 extends the spectrum of COG subunit deficiencies. <i>European Journal of Medical Genetics</i> , 2009, 52, 303-305.	1.3	34
60	COG defects, birth and rise!. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2009, 1792, 896-902.	3.8	70
61	Screening for OST deficiencies in unsolved CDG-I patients. <i>Biochemical and Biophysical Research Communications</i> , 2009, 390, 769-774.	2.1	15
62	Impaired glycosylation and cutis laxa caused by mutations in the vesicular H ⁺ -ATPase subunit ATP6VOA2. <i>Nature Genetics</i> , 2008, 40, 32-34.	21.4	330
63	Oligosaccharyltransferase-Subunit Mutations in Nonsyndromic Mental Retardation. <i>American Journal of Human Genetics</i> , 2008, 82, 1150-1157.	6.2	130
64	Deficiencies in subunits of the Conserved Oligomeric Golgi (COG) complex define a novel group of Congenital Disorders of Glycosylation. <i>Molecular Genetics and Metabolism</i> , 2008, 93, 15-21.	1.1	97
65	Cerebrocostomandibular-like syndrome and a mutation in the conserved oligomeric Golgi complex, subunit 1. <i>Human Molecular Genetics</i> , 2008, 18, 517-524.	2.9	40
66	Characterization of two unusual truncating PMM2 mutations in two CDG-Ia patients. <i>Molecular Genetics and Metabolism</i> , 2007, 90, 408-413.	1.1	42
67	A new inborn error of glycosylation due to a Cog8 deficiency reveals a critical role for the Cog1-Cog8 interaction in COG complex formation. <i>Human Molecular Genetics</i> , 2007, 16, 717-730.	2.9	114
68	A rapid mass spectrometric strategy for the characterization of N- and O-glycan chains in the diagnosis of defects in glycan biosynthesis. <i>Proteomics</i> , 2007, 7, 1800-1813.	2.2	47
69	Conserved oligomeric Golgi complex subunit 1 deficiency reveals a previously uncharacterized congenital disorder of glycosylation type II. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 3764-3769.	7.1	175
70	Discrimination between luminal and cytosolic sites of deglycosylation in endoplasmic reticulum-associated degradation of glycoproteins by using benzyl mannose in CHO cell lines. <i>Glycobiology</i> , 2004, 14, 841-849.	2.5	5
71	Endoplasmic reticulum-associated degradation of glycoproteins bearing Man5GlcNAc2 and Man9GlcNAc2 species in the MI8-5 CHO cell line. <i>FEBS Journal</i> , 2004, 271, 398-404.	0.2	25
72	The unfolded protein response in a dolichyl phosphate mannose-deficient Chinese hamster ovary cell line points out the key role of a demannosylation step in the quality-control mechanism of N-glycoproteins. <i>Biochemical Journal</i> , 2002, 362, 491.	3.7	20

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73	The unfolded protein response in a dolichyl phosphate mannose-deficient Chinese hamster ovary cell line points out the key role of a demannosylation step in the quality-control mechanism of N-glycoproteins. <i>Biochemical Journal</i> , 2002, 362, 491-498.	3.7	29