Frederic Bard

List of Publications by Year in descending order

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159585 197818 4,253 50 30 49 citations h-index g-index papers 61 61 61 7171 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	A genome-wide RNAi screen reveals determinants of human embryonic stem cell identity. Nature, 2010, 468, 316-320.	27.8	407
2	Podosomes Display Actin Turnover and Dynamic Self-Organization in Osteoclasts Expressing Actin-Green Fluorescent Protein. Molecular Biology of the Cell, 2003, 14, 407-416.	2.1	400
3	Functional genomics reveals genes involved in protein secretion and Golgi organization. Nature, 2006, 439, 604-607.	27.8	337
4	Apatite-mediated Actin Dynamics in Resorbing Osteoclasts. Molecular Biology of the Cell, 2004, 15, 5231-5241.	2.1	248
5	Protein kinase D regulates basolateral membrane protein exit from trans-Golgi network. Nature Cell Biology, 2004, 6, 106-112.	10.3	225
6	Location, location, location: new insights into O-GalNAc protein glycosylation. Trends in Cell Biology, 2011, 21, 149-158.	7.9	200
7	Deterministic Restriction on Pluripotent State Dissolution by Cell-Cycle Pathways. Cell, 2015, 162, 564-579.	28.9	185
8	The Formation of TGN-to-Plasma-Membrane Transport Carriers. Annual Review of Cell and Developmental Biology, 2006, 22, 439-455.	9.4	183
9	Regulation of <i>O</i> -glycosylation through Golgi-to-ER relocation of initiation enzymes. Journal of Cell Biology, 2010, 189, 843-858.	5. 2	178
10	Systematic Identification of Factors for Provirus Silencing in Embryonic Stem Cells. Cell, 2015, 163, 230-245.	28.9	162
11	Initiation of GalNAc-type O-glycosylation in the endoplasmic reticulum promotes cancer cell invasiveness. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E3152-61.	7.1	158
12	RNAi screening reveals a large signaling network controlling the Golgi apparatus in human cells. Molecular Systems Biology, 2012, 8, 629.	7.2	121
13	WLS Retrograde Transport to the Endoplasmic Reticulum during Wnt Secretion. Developmental Cell, 2014, 29, 277-291.	7.0	113
14	Transposon mutagenesis identifies genes driving hepatocellular carcinoma in a chronic hepatitis B mouse model. Nature Genetics, 2014, 46, 24-32.	21.4	105
15	Organelle Specific O-Glycosylation Drives MMP14 Activation, Tumor Growth, and Metastasis. Cancer Cell, 2017, 32, 639-653.e6.	16.8	102
16	Short O-GalNAc glycans: regulation and role in tumor development and clinical perspectives. Biochimica Et Biophysica Acta - General Subjects, 2016, 1860, 1623-1639.	2.4	98
17	Src Regulates Golgi Structure and KDEL Receptor-dependent Retrograde Transport to the Endoplasmic Reticulum. Journal of Biological Chemistry, 2003, 278, 46601-46606.	3.4	97
18	Cracking the Glycome Encoder: Signaling, Trafficking, and Glycosylation. Trends in Cell Biology, 2016, 26, 379-388.	7.9	82

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19	The small <scp>GTP</scp> ase Arf1 modulates mitochondrial morphology and function. EMBO Journal, 2014, 33, 2659-2675.	7.8	81
20	Addicted to secrete – novel concepts and targets in cancer therapy. Trends in Molecular Medicine, 2014, 20, 242-250.	6.7	72
21	Genome-Wide RNAi Screens Identify Genes Required for Ricin and PE Intoxications. Developmental Cell, 2011, 21, 231-244.	7.0	61
22	Nuclear envelope-associated endosomes deliver surface proteins to the nucleus. Nature Communications, 2015, 6, 8218.	12.8	61
23	Genome-Wide Screen Reveals Valosin-Containing Protein Requirement for Coronavirus Exit from Endosomes. Journal of Virology, 2015, 89, 11116-11128.	3.4	54
24	ERK8 is a negative regulator of O-GalNAc glycosylation and cell migration. ELife, 2014, 3, e01828.	6.0	52
25	TRPM5-mediated calcium uptake regulates mucin secretion from human colon goblet cells. ELife, 2013, 2, e00658.	6.0	49
26	RNAi Reveals Phase-Specific Global Regulators of Human Somatic Cell Reprogramming. Cell Reports, 2016, 15, 2597-2607.	6.4	47
27	EVI1 oncoprotein interacts with a large and complex network of proteins and integrates signals through protein phosphorylation. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E2885-94.	7.1	44
28	Molecular complexes that contain both c-Cbl and c-Src associate with Golgi membranes. European Journal of Cell Biology, 2002, 81, 26-35.	3.6	41
29	Zip14 expression induced by lipopolysaccharides in macrophages attenuates inflammatory response. Inflammation Research, 2013, 62, 133-143.	4.0	37
30	VAMP3/Syb and YKT6 are required for the fusion of constitutive secretory carriers with the plasma membrane. PLoS Genetics, 2017, 13, e1006698.	3.5	37
31	Analysis of Collagen Synthesis and Assembly in Culture by Immortalized Mouse Chondrocytes in the Presence or Absence of $\hat{l}\pm 1$ (IX) Collagen Chains. Experimental Cell Research, 1995, 219, 257-265.	2.6	32
32	The NAE Pathway: Autobahn to the Nucleus for Cell Surface Receptors. Cells, 2019, 8, 915.	4.1	25
33	ER-resident oxidoreductases are glycosylated and trafficked to the cell surface to promote matrix degradation by tumour cells. Nature Cell Biology, 2020, 22, 1371-1381.	10.3	24
34	Sar1, a Novel Regulator of ER-Mitochondrial Contact Sites. PLoS ONE, 2016, 11, e0154280.	2.5	22
35	ScreenSifter: analysis and visualization of RNAi screening data. BMC Bioinformatics, 2013, 14, 290.	2.6	19
36	The GalNAc-T Activation (GALA) Pathway: Drivers and markers. PLoS ONE, 2019, 14, e0214118.	2.5	15

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37	Loss of C2orf69 defines a fatal autoinflammatory syndrome in humans and zebrafish that evokes a glycogen-storage-associated mitochondriopathy. American Journal of Human Genetics, 2021, 108, 1301-1317.	6.2	11
38	Quiescin sulfhydryl oxidase 1 (QSOX1) glycosite mutation perturbs secretion but not Golgi localization. Glycobiology, 2018, 28, 580-591.	2.5	10
39	Digging deep into Golgi phenotypic diversity with unsupervised machine learning. Molecular Biology of the Cell, 2017, 28, 3686-3698.	2.1	8
40	Targeting c-Myc with a novel Peptide Nuclear Delivery Device. Scientific Reports, 2020, 10, 17762.	3.3	8
41	Exploratory analysis of cell-based screening data for phenotype identification in drug-siRNA study. International Journal of Computational Biology and Drug Design, 2011, 4, 194.	0.3	6
42	New developments and novel applications in high throughput and high content imaging. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2016, 89, 705-707.	1.5	6
43	Red-COLA1: a human fibroblast reporter cell line for type I collagen transcription. Scientific Reports, 2020, 10, 19723.	3.3	6
44	Comment on "The GalNAc-T Activation Pathway (GALA) is not a general mechanism for regulating mucin-type O-glycosylation― PLoS ONE, 2017, 12, e0180005.	2.5	6
45	Src activates retrograde membrane traffic through phosphorylation of GBF1. ELife, 2021, 10, .	6.0	6
46	The Ubiquitin Ligase CBLC Maintains the Network Organization of the Golgi Apparatus. PLoS ONE, 2015, 10, e0138789.	2.5	4
47	RNAi Screens for Genes Involved in Golgi Glycosylation. Methods in Molecular Biology, 2015, 1270, 411-426.	0.9	4
48	Pushing the boundaries of high content imaging. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2017, 91, 113-114.	1.5	3
49	RNAi screening reveals a large signaling network controlling the Golgi apparatus in human cells. Molecular Systems Biology, 2013, 9, 677.	7.2	1
50	HCS-PhenoCluster., 2018,,.		0