

# Magali Taulan

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

775  
citations

759233

12  
h-index

610901

24  
g-index

26  
all docs

26  
docs citations

26  
times ranked

1109  
citing authors

#	ARTICLE	IF	CITATIONS
1	Splicing mutations in the CFTR gene as therapeutic targets. <i>Gene Therapy</i> , 2022, 29, 399-406.	4.5	6
2	Exon identity influences splicing induced by exonic variants and in silico prediction efficacy. <i>Journal of Cystic Fibrosis</i> , 2021, 20, 464-472.	0.7	5
3	miRNA repertoires of cystic fibrosis ex vivo models highlight miR-181a and miR-101 that regulate WISP1 expression. <i>Journal of Pathology</i> , 2021, 253, 186-197.	4.5	7
4	The CYSMA web server: An example of integrative tool for in silico analysis of missense variants identified in Mendelian disorders. <i>Human Mutation</i> , 2020, 41, 375-386.	2.5	6
5	Current and future molecular approaches in the diagnosis of cystic fibrosis. <i>Expert Review of Respiratory Medicine</i> , 2018, 12, 415-426.	2.5	10
6	The HDAC inhibitor SAHA does not rescue CFTR membrane expression in Cystic Fibrosis. <i>International Journal of Biochemistry and Cell Biology</i> , 2017, 88, 124-132.	2.8	13
7	Targeted RNA-Seq profiling of splicing pattern in the DMD gene: exons are mostly constitutively spliced in human skeletal muscle. <i>Scientific Reports</i> , 2017, 7, 39094.	3.3	23
8	CCSP G38A polymorphism environment interactions regulate CCSP levels differentially in COPD. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2016, 311, L696-L703.	2.9	11
9	New Molecular Diagnosis Approaches From the Identification of Mutations to their Characterization. , 2015, , .		1
10	Role of Non-coding RNAs in Cystic Fibrosis. , 2015, , .		2
11	Small-scale high-throughput sequencing-based identification of new therapeutic tools in cystic fibrosis. <i>Genetics in Medicine</i> , 2015, 17, 796-806.	2.4	31
12	Should diffuse bronchiectasis still be considered a CFTR-related disorder?. <i>Journal of Cystic Fibrosis</i> , 2015, 14, 646-653.	0.7	20
13	Transcription factors and miRNAs that regulate fetal to adult CFTR expression change are new targets for cystic fibrosis. <i>European Respiratory Journal</i> , 2015, 45, 116-128.	6.7	65
14	A Classification Model Relative to Splicing for Variants of Unknown Clinical Significance: Application to the CFTR Gene. <i>Human Mutation</i> , 2013, 34, 774-784.	2.5	23
15	Phosphorylated C/EBP $\beta$ Influences a Complex Network Involving YY1 and USF2 in Lung Epithelial Cells. <i>PLoS ONE</i> , 2013, 8, e60211.	2.5	9
16	Functional analysis of a promoter variant identified in the CFTR gene in cis of a frameshift mutation. <i>European Journal of Human Genetics</i> , 2012, 20, 180-184.	2.8	6
17	Lethal factor VII deficiency due to novel mutations in the F7 promoter: Functional analysis reveals disruption of HNF4 binding site. <i>Thrombosis and Haemostasis</i> , 2012, 108, 277-283.	3.4	5
18	Identification of a novel duplication CFTRdup2 and functional impact of large rearrangements identified in the CFTR gene. <i>Gene</i> , 2012, 500, 194-198.	2.2	9

#	ARTICLE	IF	CITATIONS
19	Variants in CFTR untranslated regions are associated with congenital bilateral absence of the vas deferens. <i>Journal of Medical Genetics</i> , 2011, 48, 152-159.	3.2	22
20	NF-E2-related factor 2, a key inducer of antioxidant defenses, negatively regulates the CFTR transcription. <i>Cellular and Molecular Life Sciences</i> , 2010, 67, 2297-2309.	5.4	27
21	A novel double deletion underscores the importance of characterizing end points of the CFTR large rearrangements. <i>European Journal of Human Genetics</i> , 2009, 17, 1683-1687.	2.8	10
22	Large genomic rearrangements in the CFTR gene contribute to CBAVD. <i>BMC Medical Genetics</i> , 2007, 8, 22.	2.1	42
23	Comprehensive analysis of the renal transcriptional response to acute uranyl nitrate exposure. <i>BMC Genomics</i> , 2006, 7, 2.	2.8	49
24	Binding of serum response factor to cystic fibrosis transmembrane conductance regulator CArG-like elements, as a new potential CFTR transcriptional regulation pathway. <i>Nucleic Acids Research</i> , 2005, 33, 5271-5290.	14.5	23
25	Renal Toxicogenomic Response to Chronic Uranyl Nitrate Insult in Mice. <i>Environmental Health Perspectives</i> , 2004, 112, 1628-1635.	6.0	54
26	Oxytocin and Vasopressin V1a and V2 Receptors Form Constitutive Homo- and Heterodimers during Biosynthesis. <i>Molecular Endocrinology</i> , 2003, 17, 677-691.	3.7	296