

# Geoffray Monteuiis

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1049981/publications.pdf>

Version: 2024-02-01

11  
papers

335  
citations

1307594

7  
h-index

1372567

10  
g-index

12  
all docs

12  
docs citations

12  
times ranked

648  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mitochondrial fatty acid synthesis, fatty acids and mitochondrial physiology. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2017, 1862, 39-48.	2.4	105
2	The changing paradigm of intron retention: regulation, ramifications and recipes. <i>Nucleic Acids Research</i> , 2019, 47, 11497-11513.	14.5	90
3	Non-canonical translation initiation in yeast generates a cryptic pool of mitochondrial proteins. <i>Nucleic Acids Research</i> , 2019, 47, 5777-5791.	14.5	56
4	A conserved mammalian mitochondrial isoform of acetyl-CoA carboxylase ACC1 provides the malonyl-CoA essential for mitochondrial biogenesis in tandem with ACSF3. <i>Biochemical Journal</i> , 2017, 474, 3783-3797.	3.7	23
5	Holding on to Junk Bonds: Intron Retention in Cancer and Therapy. <i>Cancer Research</i> , 2021, 81, 779-789.	0.9	19
6	Expression and Evolution of the Non-Canonically Translated Yeast Mitochondrial Acetyl-CoA Carboxylase Hfa1p. <i>PLoS ONE</i> , 2014, 9, e114738.	2.5	17
7	Widespread Aberrant Alternative Splicing despite Molecular Remission in Chronic Myeloid Leukaemia Patients. <i>Cancers</i> , 2020, 12, 3738.	3.7	10
8	In-frame deletion in canine PITRM1 is associated with a severe early-onset epilepsy, mitochondrial dysfunction and neurodegeneration. <i>Human Genetics</i> , 2021, 140, 1593-1609.	3.8	9
9	The Fusion of CLEC12A and MIR223HG Arises from a trans-Splicing Event in Normal and Transformed Human Cells. <i>International Journal of Molecular Sciences</i> , 2021, 22, 12178.	4.1	4
10	A hunt for OM45 synthetic petite interactions in <i>Saccharomyces cerevisiae</i> reveals a role for Miro GTPase Gem1p in cristae structure maintenance. <i>MicrobiologyOpen</i> , 2021, 10, e1238.	3.0	1
11	Computational Methods for Intron Retention Identification and Quantification. , 2021, , 63-74.		0