

# Nicolas Lenfant

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2643092/publications.pdf>

Version: 2024-02-01

19  
papers

1,566  
citations

687220

13  
h-index

794469

19  
g-index

19  
all docs

19  
docs citations

19  
times ranked

2375  
citing authors

#	ARTICLE	IF	CITATIONS
1	FGF10 promotes cardiac repair through a dual cellular mechanism increasing cardiomyocyte renewal and inhibiting fibrosis. <i>Cardiovascular Research</i> , 2022, 118, 2625-2637.	1.8	16
2	An evolutionary perspective on the first disulfide bond in members of the cholinesterase-carboxylesterase (COesterase) family: Possible outcomes for cholinesterase expression in prokaryotes. <i>Chemico-Biological Interactions</i> , 2019, 308, 179-184.	1.7	3
3	Broad-specificity GH131 $\beta$ -glucanases are a hallmark of fungi and oomycetes that colonize plants. <i>Environmental Microbiology</i> , 2019, 21, 2724-2739.	1.8	18
4	Lytic xylan oxidases from wood-decay fungi unlock biomass degradation. <i>Nature Chemical Biology</i> , 2018, 14, 306-310.	3.9	269
5	A bioinformatics analysis of 3400 lytic polysaccharide oxidases from family AA9. <i>Carbohydrate Research</i> , 2017, 448, 166-174.	1.1	55
6	Natural genomic amplification of cholinesterase genes in animals. <i>Journal of Neurochemistry</i> , 2017, 142, 73-81.	2.1	8
7	CAZyme content of <i>Pochonia chlamydosporia</i> reflects that chitin and chitosan modification are involved in nematode parasitism. <i>Environmental Microbiology</i> , 2016, 18, 4200-4215.	1.8	41
8	Relationships of human $\beta$ -hydrolase fold proteins and other organophosphate-interacting proteins. <i>Chemico-Biological Interactions</i> , 2016, 259, 343-351.	1.7	9
9	Dividing the Large Glycoside Hydrolase Family 43 into Subfamilies: a Motivation for Detailed Enzyme Characterization. <i>Applied and Environmental Microbiology</i> , 2016, 82, 1686-1692.	1.4	173
10	The molecular basis of polysaccharide cleavage by lytic polysaccharide monooxygenases. <i>Nature Chemical Biology</i> , 2016, 12, 298-303.	3.9	264
11	Structure-Function Analysis of a Mixed-linkage $\beta$ -Glucanase/Xyloglucanase from the Key Ruminant Bacteroidetes <i>Prevotella bryantii</i> B14. <i>Journal of Biological Chemistry</i> , 2016, 291, 1175-1197.	1.6	38
12	Structure and boosting activity of a starch-degrading lytic polysaccharide monooxygenase. <i>Nature Communications</i> , 2015, 6, 5961.	5.8	254
13	Molecular characterization of an acetylcholinesterase from the hemichordate <i>Saccoglossus kowalevskii</i> . <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2015, 181, 50-58.	0.7	3
14	Tracking the Origin and Divergence of Cholinesterases and Neuroligins: The Evolution of Synaptic Proteins. <i>Journal of Molecular Neuroscience</i> , 2014, 53, 362-369.	1.1	11
15	Proteins with an alpha/beta hydrolase fold: Relationships between subfamilies in an ever-growing superfamily. <i>Chemico-Biological Interactions</i> , 2013, 203, 266-268.	1.7	39
16	The Human PDZome: A Gateway to PSD95-Disc Large-Zonula Occludens (PDZ)-mediated Functions. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 2587-2603.	2.5	59
17	Prevalence, Specificity and Determinants of Lipid-Interacting PDZ Domains from an In-Cell Screen and In Vitro Binding Experiments. <i>PLoS ONE</i> , 2013, 8, e54581.	1.1	23
18	ESTHER, the database of the $\beta$ -hydrolase fold superfamily of proteins: tools to explore diversity of functions. <i>Nucleic Acids Research</i> , 2012, 41, D423-D429.	6.5	244

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19	A genome-wide study of PDZ-domain interactions in <i>C. elegans</i> reveals a high frequency of non-canonical binding. <i>BMC Genomics</i> , 2010, 11, 671.	1.2	39