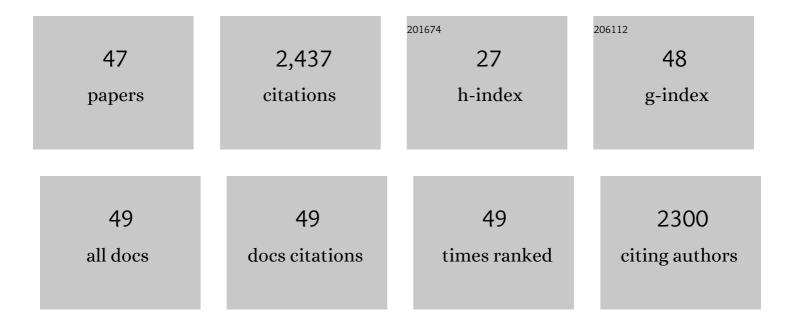
Eric ChabriÃ"re

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Enzyme Nanoreactor for <i>In Vivo</i> Detoxification of Organophosphates. ACS Applied Materials & Interfaces, 2022, , .	8.0	9
2	Applying molecular and phenotypic screening assays to identify efficient quorum quenching lactonases. Enzyme and Microbial Technology, 2022, 160, 110092.	3.2	5
3	Organophosphorus poisoning in animals and enzymatic antidotes. Environmental Science and Pollution Research, 2021, 28, 25081-25106.	5.3	17
4	Disrupting quorum sensing alters social interactions in Chromobacterium violaceum. Npj Biofilms and Microbiomes, 2021, 7, 40.	6.4	30
5	Enzymatic Decontamination of G-Type, V-Type and Novichok Nerve Agents. International Journal of Molecular Sciences, 2021, 22, 8152.	4.1	20
6	Engineering acyl-homoserine lactone-interfering enzymes toward bacterial control. Journal of Biological Chemistry, 2020, 295, 12993-13007.	3.4	42
7	Lactonase Specificity Is Key to Quorum Quenching in Pseudomonas aeruginosa. Frontiers in Microbiology, 2020, 11, 762.	3.5	35
8	Steady-State Kinetics of Enzyme-Catalyzed Hydrolysis of Echothiophate, a P–S Bonded Organophosphorus as Monitored by Spectrofluorimetry. Molecules, 2020, 25, 1371.	3.8	7
9	Enzymatic decontamination of paraoxon-ethyl limits long-term effects in planarians. Scientific Reports, 2020, 10, 3843.	3.3	3
10	Lactonase SsoPox modulates CRISPR-Cas expression in gram-negative proteobacteria using AHL-based quorum sensing systems. Research in Microbiology, 2019, 170, 296-299.	2.1	16
11	Quorum Quenching Lactonase Strengthens Bacteriophage and Antibiotic Arsenal Against Pseudomonas aeruginosa Clinical Isolates. Frontiers in Microbiology, 2019, 10, 2049.	3.5	41
12	Evaluation of a robust engineered enzyme towards organophosphorus insecticide bioremediation using planarians as biosensors. Chemico-Biological Interactions, 2019, 306, 96-103.	4.0	9
13	Structural and Biochemical Characterization of AaL, a Quorum Quenching Lactonase with Unusual Kinetic Properties. Scientific Reports, 2018, 8, 11262.	3.3	38
14	Interference in Bacterial Quorum Sensing: A Biopharmaceutical Perspective. Frontiers in Pharmacology, 2018, 9, 203.	3.5	230
15	Biotechnological applications of quorum quenching enzymes. Chemico-Biological Interactions, 2017, 267, 104-115.	4.0	138
16	New Approaches to Prevent Healthcare-Associated Infection. Clinical Infectious Diseases, 2017, 65, S50-S54.	5.8	25
17	Rational engineering of a native hyperthermostable lactonase into a broad spectrum phosphotriesterase. Scientific Reports, 2017, 7, 16745.	3.3	39
18	Enzymatic degradation of organophosphorus insecticides decreases toxicity in planarians and enhances survival. Scientific Reports, 2017, 7, 15194.	3.3	34

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19	Effect of Quorum Quenching Lactonase in Clinical Isolates of Pseudomonas aeruginosa and Comparison with Quorum Sensing Inhibitors. Frontiers in Microbiology, 2017, 08, 227.	3.5	120
20	Malaria: Massive open online courses MOOC. Travel Medicine and Infectious Disease, 2016, 14, 636.	3.0	4
21	Harnessing hyperthermostable lactonase from Sulfolobus solfataricus for biotechnological applications. Scientific Reports, 2016, 6, 37780.	3.3	38
22	Current and emerging strategies for organophosphate decontamination: special focus on hyperstable enzymes. Environmental Science and Pollution Research, 2016, 23, 8200-8218.	5.3	72
23	Crystal structure of VmoLac, a tentative quorum quenching lactonase from the extremophilic crenarchaeon Vulcanisaeta moutnovskia. Scientific Reports, 2015, 5, 8372.	3.3	44
24	Surface plasmon resonance imaging of pathogens: the Yersinia pestis paradigm. BMC Research Notes, 2015, 8, 259.	1.4	6
25	Inhaled Lactonase Reduces Pseudomonas aeruginosa Quorum Sensing and Mortality in Rat Pneumonia. PLoS ONE, 2014, 9, e107125.	2.5	97
26	Sac Pox from the thermoacidophilic crenarchaeon Sulfolobus acidocaldarius is a proficient lactonase. BMC Research Notes, 2014, 7, 333.	1.4	25
27	Ancestral mutations as a tool for solubilizing proteins: The case of a hydrophobic phosphateâ€binding protein. FEBS Open Bio, 2014, 4, 121-127.	2.3	13
28	Crystallization and preliminary X-ray diffraction analysis of the lactonaseVmoLac fromVulcanisaeta moutnovskia. Acta Crystallographica Section F: Structural Biology Communications, 2013, 69, 1235-1238.	0.7	15
29	Crystallization and preliminary X-ray diffraction analysis of the organophosphorus hydrolase OPHC2 from <i>Pseudomonas pseudoalcaligenes</i> . Acta Crystallographica Section F: Structural Biology Communications, 2013, 69, 73-76.	0.7	13
30	Structural and Enzymatic Characterization of the Phosphotriesterase OPHC2 from Pseudomonas pseudoalcaligenes. PLoS ONE, 2013, 8, e77995.	2.5	50
31	DING Proteins from Phylogenetically Different Species Share High Degrees of Sequence and Structure Homology and Block Transcription of HIV-1 LTR Promoter. PLoS ONE, 2013, 8, e69623.	2.5	10
32	Differential Active Site Loop Conformations Mediate Promiscuous Activities in the Lactonase SsoPox. PLoS ONE, 2013, 8, e75272.	2.5	89
33	Characterisation of the organophosphate hydrolase catalytic activity of SsoPox. Scientific Reports, 2012, 2, 779.	3.3	82
34	The molecular basis of phosphate discrimination in arsenate-rich environments. Nature, 2012, 491, 134-137.	27.8	209
35	Structural and Enzymatic characterization of the lactonase SisLac from Sulfolobus islandicus. PLoS ONE, 2012, 7, e47028.	2.5	70
36	Human-Phosphate-Binding-Protein inhibits HIV-1 gene transcription and replication. Virology Journal, 2011, 8, 352.	3.4	18

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37	The DING family of proteins: ubiquitous in eukaryotes, but where are the genes?. BioEssays, 2009, 31, 570-580.	2.5	31
38	Structural determinants of the high thermal stability of SsoPox from the hyperthermophilic archaeon Sulfolobus solfataricus. Extremophiles, 2009, 13, 461-470.	2.3	60
39	Elucidation of the Phosphate Binding Mode of DING Proteins Revealed by Subangstrom X-ray Crystallography. Journal of the American Chemical Society, 2009, 131, 7879-7886.	13.7	50
40	Tandem use of Xâ€ray crystallography and mass spectrometry to obtain ab initio the complete and exact amino acids sequence of HPBP, a human 38â€kDa apolipoprotein. Proteins: Structure, Function and Bioinformatics, 2008, 71, 1708-1720.	2.6	36
41	Structural Basis for Natural Lactonase and Promiscuous Phosphotriesterase Activities. Journal of Molecular Biology, 2008, 379, 1017-1028.	4.2	159
42	Structure-function relationships in a bacterial DING protein. FEBS Letters, 2007, 581, 3455-3460.	2.8	29
43	Human paraoxonase: A promising approach for pre-treatment and therapy of organophosphorus poisoning. Toxicology, 2007, 233, 47-59.	4.2	137
44	Crystallization and preliminary X-ray diffraction analysis of the hyperthermophilic <i>Sulfolobus solfataricus</i> phosphotriesterase. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 553-555.	0.7	19
45	Crystallization, diffraction data collection and preliminary crystallographic analysis of DING protein from <i>Pseudomonas fluorescens</i> . Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 590-592.	0.7	17
46	Tandem purification of two HDL-associated partner proteins in human plasma, paraoxonase (PON1) and phosphate binding protein (HPBP) using hydroxyapatite chromatography. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2006, 836, 15-21.	2.3	93
47	Serendipitous Discovery and X-Ray Structure of a Human Phosphate Binding Apolipoprotein. Structure, 2006, 14, 601-609.	3.3	79