Pascal Mercier

List of Publications by Year in descending order

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567144 642610 2,112 23 15 23 citations h-index g-index papers 23 23 23 3191 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Structural Basis of <i>Tirasemtiv</i> Activation of Fast Skeletal Muscle. Journal of Medicinal Chemistry, 2021, 64, 3026-3034.	2.9	9
2	Characterizing the inhibition of $\hat{l}\pm\hat{a}\in s$ ynuclein oligomerization by a pharmacological chaperone that prevents prion formation by the protein PrP. Protein Science, 2019, 28, 1690-1702.	3.1	9
3	Comparison of computational approaches for identification and quantification of urinary metabolites in ¹ H NMR spectra. Analytical Methods, 2018, 10, 2129-2137.	1.3	4
4	Dataset of urinary metabolites measured by 1 H NMR analysis of normal human urine. Data in Brief, 2017, 10, 227-229.	0.5	1
5	Structure of phosphorylated UBL domain and insights into PINK1-orchestrated parkin activation. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 298-303.	3.3	77
6	Insights into the Mechanism of Action of the Two-Peptide Lantibiotic Lacticin 3147. Journal of the American Chemical Society, 2017, 139, 17803-17810.	6.6	38
7	Assessment of 1 H NMR-based metabolomics analysis for normalization of urinary metals against creatinine. Clinica Chimica Acta, 2017, 464, 37-43.	0.5	11
8	Antimicrobial lipopeptide tridecaptin A ₁ selectively binds to Gram-negative lipid II. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 11561-11566.	3.3	127
9	Disruption of the autoinhibited state primes the E3 ligase parkin for activation and catalysis. EMBO Journal, 2015, 34, 2506-2521.	3.5	160
10	A molecular explanation for the recessive nature of parkin-linked Parkinson's disease. Nature Communications, 2013, 4, 1983.	5.8	123
11	Structure of the HHARI Catalytic Domain Shows Glimpses of a HECT E3 Ligase. PLoS ONE, 2013, 8, e74047.	1.1	12
12	Towards automatic metabolomic profiling of high-resolution one-dimensional proton NMR spectra. Journal of Biomolecular NMR, 2011, 49, 307-323.	1.6	117
13	Is there nascent structure in the intrinsically disordered region of troponin I?. Proteins: Structure, Function and Bioinformatics, 2011, 79, 1240-1250.	1.5	23
14	Metabolite profiling of the intraerythrocytic malaria parasite $\langle i \rangle$ Plasmodium falciparum $\langle i \rangle$ by $\langle \sup 1 \langle \sup H \rangle$ NMR spectroscopy. NMR in Biomedicine, 2009, 22, 292-302.	1.6	101
15	The effect of the cosolvent trifluoroethanol on a tryptophan side chain orientation in the hydrophobic core of troponin C. Protein Science, 2009, 18, 1165-1174.	3.1	7
16	NMR Studies of the Dynamics of a Bifunctional Rhodamine Probe Attached to Troponin C. Journal of the American Chemical Society, 2008, 130, 2602-2609.	6.6	6
17	Targeted Profiling:Â Quantitative Analysis of 1H NMR Metabolomics Data. Analytical Chemistry, 2006, 78, 4430-4442.	3.2	844
18	Effects of Phe-to-Trp mutation and fluorotryptophan incorporation on the solution structure of cardiac troponin C, and analysis of its suitability as a potential probe for in situ NMR studies. Protein Science, 2005, 14, 2447-2460.	3.1	23

#	Article	IF	CITATIONS
19	Structure of Subtilosin A, a Cyclic Antimicrobial Peptide fromBacillus subtiliswith Unusual Sulfur to α-Carbon Cross-Links: Formation and Reduction of α-Thio-α-Amino Acid Derivativesâ€,‡. Biochemistry, 2004, 43, 3385-3395.	1.2	185
20	Structure of Subtilosin A, an Antimicrobial Peptide from Bacillus subtilis with Unusual Posttranslational Modifications Linking Cysteine Sulfurs to \hat{l}_{\pm} -Carbons of Phenylalanine and Threonine. Journal of the American Chemical Society, 2003, 125, 4726-4727.	6.6	111
21	NMR Structure of a Bifunctional Rhodamine Labeled N-Domain of Troponin C Complexed with the Regulatory "Switch―Peptide from Troponin I:  Implications for in Situ Fluorescence Studies in Muscle Fibers,. Biochemistry, 2003, 42, 4333-4348.	1.2	33
22	In Situ Orientations of Protein Domains. Molecular Cell, 2003, 11, 865-874.	4.5	51
23	Structure, Dynamics, and Thermodynamics of the Structural Domain of Troponin C in Complex with the Regulatory Peptide 1â°⁴40 of Troponin Iâ€,‡. Biochemistry, 2001, 40, 10063-10077.	1.2	40