## Douglas S Daniels

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7294977/publications.pdf

Version: 2024-02-01

20 papers 1,512 citations

567281 15 h-index 19 g-index

20 all docs

20 does citations

times ranked

20

2048 citing authors

#	Article	IF	CITATIONS
1	The Autophagy-Related Beclin-1 Protein Requires the Coiled-Coil and BARA Domains To Form a Homodimer with Submicromolar Affinity. Biochemistry, 2017, 56, 6639-6651.	2.5	14
2	Point mutations at the catalytic site of PCSK9 inhibit folding, autoprocessing, and interaction with the LDL receptor. Protein Science, 2016, 25, 2018-2027.	7.6	11
3	A Maltose-Binding Protein Fusion Construct Yields a Robust Crystallography Platform for MCL1. PLoS ONE, 2015, 10, e0125010.	2.5	26
4	A Small Molecule That Binds and Inhibits the ETV1 Transcription Factor Oncoprotein. Molecular Cancer Therapeutics, 2014, 13, 1492-1502.	4.1	52
5	New hypotheses about the structure–function of proprotein convertase subtilisin/kexin type 9: Analysis of the epidermal growth factorâ€like repeat A docking site using WaterMap. Proteins: Structure, Function and Bioinformatics, 2010, 78, 2571-2586.	2.6	65
6	$\hat{l}^2$ -Peptides with improved affinity for hDM2 and hDMX. Bioorganic and Medicinal Chemistry, 2009, 17, 2038-2046.	3.0	66
7	Minimally Cationic Cell-Permeable Miniature Proteins via $\hat{l}_{\pm}$ -Helical Arginine Display. Journal of the American Chemical Society, 2008, 130, 2948-2949.	13.7	102
8	High-Resolution Structure of a $\hat{l}^2$ -Peptide Bundle. Journal of the American Chemical Society, 2007, 129, 1532-1533.	13.7	195
9	Biophysical and Structural Characterization of a Robust Octameric $\hat{l}^2$ -Peptide Bundle. Journal of the American Chemical Society, 2007, 129, 14746-14751.	13.7	63
10	Biophysical Characterization of a $\hat{l}^2$ -Peptide Bundle: $\hat{A}$ Comparison to Natural Proteins. Journal of the American Chemical Society, 2007, 129, 5344-5345.	13.7	54
11	Intrinsically Cell-Permeable Miniature Proteins Based on a Minimal Cationic PPII Motif. Journal of the American Chemical Society, 2007, 129, 14578-14579.	13.7	108
12	DNA binding and nucleotide flipping by the human DNA repair protein AGT. Nature Structural and Molecular Biology, 2004, 11, 714-720.	8.2	275
13	Characterization of the electrophile binding site and substrate binding mode of the 26-kDa glutathione S-transferase fromSchistosoma japonicum. Proteins: Structure, Function and Bioinformatics, 2003, 51, 137-146.	2.6	43
14	CysG structure reveals tetrapyrrole-binding features and novel regulation of siroheme biosynthesis. Nature Structural and Molecular Biology, 2003, 10, 1064-1073.	8.2	78
15	Full-length archaeal Rad51 structure and mutants: mechanisms for RAD51 assembly and control by BRCA2. EMBO Journal, 2003, 22, 4566-4576.	7.8	239
16	DNA damage recognition and repair pathway coordination revealed by the structural biochemistry of DNA repair enzymes. Progress in Molecular Biology and Translational Science, 2001, 68, 315-347.	1.9	30
17	Conserved structural motifs governing the stoichiometric repair of alkylated DNA by O6-alkylguanine-DNA alkyltransferase. Mutation Research DNA Repair, 2000, 460, 151-163.	3.7	76
18	Mea Culpa: Formal Education and the Dis-Integrated World. , 1999, , 107-128.		0

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#	Article	lF	CITATIONS
19	Structuring the Liberal (Arts) Education in Chemistry. The Chemical Educator, 1996, 1, 1-32.	0.0	8
20	I Scream, You Scream: A New Twist on the Liquid Nitrogen Demonstrations. Journal of Chemical Education, 1994, 71, 1080.	2.3	7