## A Geoffrey Skillman

List of Publications by Year in descending order

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759233 1125743 3,810 13 12 13 citations h-index g-index papers 13 13 13 5175 docs citations times ranked citing authors all docs

#	Article	lF	CITATIONS
1	Mechanistic Insights into Passive Membrane Permeability of Drug-like Molecules from a Weighted Ensemble of Trajectories. Journal of Chemical Information and Modeling, 2022, 62, 1891-1904.	5.4	7
2	SAMPL6 challenge results from \$\$pK_a\$\$ predictions based on a general Gaussian process model. Journal of Computer-Aided Molecular Design, 2018, 32, 1165-1177.	2.9	20
3	Efficient calculation of SAMPL4 hydration free energies using OMEGA, SZYBKI, QUACPAC, and Zap TK. Journal of Computer-Aided Molecular Design, 2014, 28, 289-298.	2.9	13
4	Conformer Generation with OMEGA: Algorithm and Validation Using High Quality Structures from the Protein Databank and Cambridge Structural Database. Journal of Chemical Information and Modeling, 2010, 50, 572-584.	5 <b>.</b> 4	1,302
5	How to do an evaluation: pitfalls and traps. Journal of Computer-Aided Molecular Design, 2008, 22, 179-190.	2.9	126
6	Comparison of Shape-Matching and Docking as Virtual Screening Tools. Journal of Medicinal Chemistry, 2007, 50, 74-82.	6.4	826
7	A novel mechanism for inhibition of HIV-1 reverse transcriptase. Bioorganic Chemistry, 2002, 30, 443-458.	4.1	27
8	Design, docking, and evaluation of multiple libraries against multiple targets. Proteins: Structure, Function and Bioinformatics, 2001, 42, 296-318.	2.6	66
9	DOCK 4.0: search strategies for automated molecular docking of flexible molecule databases. Journal of Computer-Aided Molecular Design, 2001, 15, 411-428.	2.9	1,022
10	Novel Cathepsin D Inhibitors Block the Formation of Hyperphosphorylated Tau Fragments in Hippocampus. Journal of Neurochemistry, 2000, 74, 1469-1477.	3.9	67
11	A rapid method for exploring the protein structure universe. Proteins: Structure, Function and Bioinformatics, 1999, 34, 317-332.	2.6	15
12	Potent, Low-Molecular-Weight Non-Peptide Inhibitors of Malarial Aspartyl Protease Plasmepsin II. Journal of Medicinal Chemistry, 1999, 42, 1428-1440.	6.4	173
13	Structure-based design and combinatorial chemistry yield low nanomolar inhibitors of cathepsin D. Chemistry and Biology, 1997, 4, 297-307.	6.0	146