

# Paul N Mortenson

## List of Publications by Year in descending order

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Version: 2024-02-01

30  
papers

2,106  
citations

331259

21  
h-index

433756

31  
g-index

32  
all docs

32  
docs citations

32  
times ranked

2879  
citing authors

#	ARTICLE	IF	CITATIONS
1	Diverse, High-Quality Test Set for the Validation of Protein-Ligand Docking Performance. <i>Journal of Medicinal Chemistry</i> , 2007, 50, 726-741.	2.9	546
2	Energy Landscapes: From Clusters to Biomolecules. <i>Advances in Chemical Physics</i> , 2007, , 1-111.	0.3	153
3	Protein-Ligand Docking against Non-Native Protein Conformers. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 2214-2225.	2.5	129
4	Efficient exploration of chemical space by fragment-based screening. <i>Progress in Biophysics and Molecular Biology</i> , 2014, 116, 82-91.	1.4	127
5	Docking Performance of Fragments and Druglike Compounds. <i>Journal of Medicinal Chemistry</i> , 2011, 54, 5422-5431.	2.9	109
6	Assessing the lipophilicity of fragments and early hits. <i>Journal of Computer-Aided Molecular Design</i> , 2011, 25, 663-667.	1.3	95
7	Energy landscapes of model polyanilines. <i>Journal of Chemical Physics</i> , 2002, 117, 1363-1376.	1.2	78
8	Energy landscapes, global optimization and dynamics of the polyaniline Ac(ala)8NHMe. <i>Journal of Chemical Physics</i> , 2001, 114, 6443-6454.	1.2	77
9	Crystals of binary Lennard-Jones solids. <i>Physical Review B</i> , 2001, 64, .	1.1	77
10	Discovery of 2-(6-[[6-Fluoroquinolin-2-yl)methyl]amino}bicyclo[3.1.0]hex-3-yl)- <i>N</i> -hydroxypyrimidine-5-carboxamide (CHR-3996), a Class I Selective Orally Active Histone Deacetylase Inhibitor. <i>Journal of Medicinal Chemistry</i> , 2010, 53, 8663-8678.	2.9	74
11	DNA gyrase (GyrB)/topoisomerase IV (ParE) inhibitors: Synthesis and antibacterial activity. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2009, 19, 894-899.	1.0	70
12	Discovery of novel inhibitors of <i>Trypanosoma cruzi</i> trans-sialidase from in silico screening. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2009, 19, 589-596.	1.0	68
13	Fragment-to-Lead Medicinal Chemistry Publications in 2018. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 4430-4444.	2.9	61
14	Structure-Activity and Structure-Conformation Relationships of Aryl Propionic Acid Inhibitors of the Kelch-like ECH-Associated Protein 1/Nuclear Factor Erythroid 2-Related Factor 2 (KEAP1/NRF2) Protein-Protein Interaction. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 4683-4702.	2.9	59
15	Fragment-Based Discovery of Potent and Selective DDR1/2 Inhibitors. <i>ACS Medicinal Chemistry Letters</i> , 2015, 6, 798-803.	1.3	49
16	Fragment-to-Lead Medicinal Chemistry Publications in 2017. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 3857-3872.	2.9	47
17	Fragment-to-Lead Medicinal Chemistry Publications in 2016. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 1774-1784.	2.9	41
18	Fragment-to-Lead Medicinal Chemistry Publications in 2019. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 15494-15507.	2.9	41

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19	Fragment-Based Discovery of 7-Azabenzimidazoles as Potent, Highly Selective, and Orally Active CDK4/6 Inhibitors. <i>ACS Medicinal Chemistry Letters</i> , 2012, 3, 445-449.	1.3	34
20	Predicting "Hot" and "Warm" Spots for Fragment Binding. <i>Journal of Medicinal Chemistry</i> , 2017, 60, 4036-4046.	2.9	32
21	C-H functionalisation tolerant to polar groups could transform fragment-based drug discovery (FBDD). <i>Chemical Science</i> , 2021, 12, 11976-11985.	3.7	30
22	ParaMol: A Package for Automatic Parameterization of Molecular Mechanics Force Fields. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 2026-2047.	2.5	22
23	Fragment-Based Discovery of 6-Azaindazoles As Inhibitors of Bacterial DNA Ligase. <i>ACS Medicinal Chemistry Letters</i> , 2013, 4, 1208-1212.	1.3	21
24	Allosteric Inhibition of the Neuropeptidase Neurolysin. <i>Journal of Biological Chemistry</i> , 2014, 289, 35605-35619.	1.6	18
25	The discovery of quinoline-3-carboxamides as hematopoietic prostaglandin D synthase (H-PGDS) inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2019, 27, 1456-1478.	1.4	17
26	Fragment-Based Approaches to the Discovery of Kinase Inhibitors. <i>Methods in Enzymology</i> , 2014, 548, 69-92.	0.4	13
27	Identification of orally bioavailable small-molecule inhibitors of hematopoietic prostaglandin D2 synthase using X-ray fragment based drug discovery. <i>MedChemComm</i> , 2014, 5, 134-141.	3.5	8
28	The exploration of aza-quinolines as hematopoietic prostaglandin D synthase (H-PGDS) inhibitors with low brain exposure. <i>Bioorganic and Medicinal Chemistry</i> , 2020, 28, 115791.	1.4	4
29	A knowledge-based, structural-aided discovery of a novel class of 2-phenylimidazo[1,2-a]pyridine-6-carboxamide H-PGDS inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2021, 47, 128113.	1.0	3
30	Generation of Quantum Configurational Ensembles Using Approximate Potentials. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 7021-7042.	2.3	2