

# Scott A Wildman

## List of Publications by Year in descending order

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32  
papers

2,189  
citations

430442

18  
h-index

414034

32  
g-index

36  
all docs

36  
docs citations

36  
times ranked

3443  
citing authors

#	ARTICLE	IF	CITATIONS
1	Discovery and Mechanism of Small Molecule Inhibitors Selective for the Chromatin-Binding Domains of Oncogenic UHRF1. <i>Biochemistry</i> , 2022, 61, 354-366.	1.2	8
2	TRPV4 receptor as a functional sensory molecule in bladder urothelium: Stretch-independent, tissue-specific actions and pathological implications. <i>FASEB Journal</i> , 2020, 34, 263-286.	0.2	26
3	Predicting kinase inhibitors using bioactivity matrix derived informer sets. <i>PLoS Computational Biology</i> , 2019, 15, e1006813.	1.5	9
4	Identification of small molecule enzyme inhibitors as broad-spectrum anthelmintics. <i>Scientific Reports</i> , 2019, 9, 9085.	1.6	25
5	Piperidine carbamate peptidomimetic inhibitors of the serine proteases HGFA, matriptase and hepsin. <i>MedChemComm</i> , 2019, 10, 1646-1655.	3.5	8
6	A general strategy for diversifying complex natural products to polycyclic scaffolds with medium-sized rings. <i>Nature Communications</i> , 2019, 10, 4015.	5.8	68
7	The antimicrobial potential of <i>Streptomyces</i> from insect microbiomes. <i>Nature Communications</i> , 2019, 10, 516.	5.8	222
8	Identification of a novel class of RIP1/RIP3 dual inhibitors that impede cell death and inflammation in mouse abdominal aortic aneurysm models. <i>Cell Death and Disease</i> , 2019, 10, 226.	2.7	69
9	Identification of 4-Amino-Thieno[2,3- <i>d</i> ]Pyrimidines as QcrB Inhibitors in <i>Mycobacterium tuberculosis</i> . <i>MSphere</i> , 2019, 4, .	1.3	19
10	Practical Model Selection for Prospective Virtual Screening. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 282-293.	2.5	46
11	Small Molecule Inhibitors of Metabolic Enzymes Repurposed as a New Class of Anthelmintics. <i>ACS Infectious Diseases</i> , 2018, 4, 1130-1145.	1.8	18
12	A High-Throughput Screening Strategy to Identify Inhibitors of SSB Protein-Protein Interactions in an Academic Screening Facility. <i>SLAS Discovery</i> , 2018, 23, 94-101.	1.4	23
13	Machine Learning Consensus Scoring Improves Performance Across Targets in Structure-Based Virtual Screening. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 1579-1590.	2.5	84
14	2- <i>tert</i> -Butyl-4-oxo-1,2,3,4-tetrahydro-1H-benzothiazole Serine Protease Inhibitors of Aberrant HGF/c-MET and MSP/RON Kinase Pathway Signaling in Cancer. <i>ChemMedChem</i> , 2016, 11, 585-599.	1.6	32
15	Tumor-Specific Binding of Radiolabeled PEGylated GIRLRG Peptide: A Novel Agent for Targeting Cancers. <i>Journal of Nuclear Medicine</i> , 2016, 57, 1991-1997.	2.8	21
16	Hopeahainol A binds reversibly at the acetylcholinesterase (AChE) peripheral site and inhibits enzyme activity with a novel higher order concentration dependence. <i>Chemico-Biological Interactions</i> , 2016, 259, 78-84.	1.7	5
17	Characterization of parasite-specific indels and their proposed relevance for selective anthelmintic drug targeting. <i>Infection, Genetics and Evolution</i> , 2016, 39, 201-211.	1.0	7
18	Identification of Small Molecule Inhibitors That Block the <i>Toxoplasma gondii</i> Rhoptyry Kinase ROP18. <i>ACS Infectious Diseases</i> , 2016, 2, 194-206.	1.8	20

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19	Structure-based discovery of small molecule hepsin and HGFA protease inhibitors: Evaluation of potency and selectivity derived from distinct binding pockets. <i>Bioorganic and Medicinal Chemistry</i> , 2015, 23, 2328-2343.	1.4	31
20	Inhibitors of HGFA, Matrilptase, and Hepsin Serine Proteases: A Nonkinase Strategy to Block Cell Signaling in Cancer. <i>ACS Medicinal Chemistry Letters</i> , 2014, 5, 1219-1224.	1.3	41
21	Isoform-selective Inhibition of Facilitative Glucose Transporters. <i>Journal of Biological Chemistry</i> , 2014, 289, 16100-16113.	1.6	16
22	The Natural Product Dihydrotanshinone I Provides a Prototype for Uncharged Inhibitors That Bind Specifically to the Acetylcholinesterase Peripheral Site with Nanomolar Affinity. <i>Biochemistry</i> , 2013, 52, 7486-7499.	1.2	10
23	Optimizing Small Molecule Inhibitors of Calcium-Dependent Protein Kinase 1 to Prevent Infection by <i>Toxoplasma gondii</i> . <i>Journal of Medicinal Chemistry</i> , 2013, 56, 3068-3077.	2.9	64
24	High-throughput identification of putative receptors for cancer-binding peptides using biopanning and microarray analysis. <i>Integrative Biology (United Kingdom)</i> , 2013, 5, 342-350.	0.6	6
25	Approaches to Virtual Screening and Screening Library Selection. <i>Current Pharmaceutical Design</i> , 2013, 19, 4787-4796.	0.9	10
26	Drug-like Leads for Steric Discrimination between Substrate and Inhibitors of Human Acetylcholinesterase. <i>Chemical Biology and Drug Design</i> , 2011, 78, 495-504.	1.5	20
27	Structure-Based Drug Design and Optimization of Mannoside Bacterial FimH Antagonists. <i>Journal of Medicinal Chemistry</i> , 2010, 53, 4779-4792.	2.9	220
28	Predicting Kinase Selectivity Profiles Using Free-Wilson QSAR Analysis. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 1851-1867.	2.5	39
29	Validation of DAPPER for 3D QSAR: Conformational Search and Chirality Metric. <i>Journal of Chemical Information and Computer Sciences</i> , 2003, 43, 629-636.	2.8	14
30	Three-dimensional molecular descriptors and a novel QSAR method. <i>Journal of Molecular Graphics and Modelling</i> , 2002, 21, 161-170.	1.3	18
31	Evaluation of Ligand Overlap by Atomic Parameters. <i>Journal of Chemical Information and Computer Sciences</i> , 2001, 41, 446-450.	2.8	11
32	Prediction of Physicochemical Parameters by Atomic Contributions. <i>Journal of Chemical Information and Computer Sciences</i> , 1999, 39, 868-873.	2.8	977