

Ana C Puhl

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

841
citations

759233

12
h-index

713466

21
g-index

33
all docs

33
docs citations

33
times ranked

1365
citing authors

#	ARTICLE	IF	CITATIONS
1	Exploiting machine learning for end-to-end drug discovery and development. <i>Nature Materials</i> , 2019, 18, 435-441.	27.5	334
2	Mode of Peroxisome Proliferator-Activated Receptor β Activation by Luteolin. <i>Molecular Pharmacology</i> , 2012, 81, 788-799.	2.3	84
3	D α -Tuberculin: Stimulating open drug discovery for SARS-CoV-2. <i>Drug Discovery Today</i> , 2020, 25, 928-941.	6.4	81
4	Mechanisms of Peroxisome Proliferator Activated Receptor β Regulation by Non-steroidal Anti-inflammatory Drugs. <i>Nuclear Receptor Signaling</i> , 2015, 13, nrs.13004.	1.0	63
5	Repurposing the Ebola and Marburg Virus Inhibitors Tilorone, Quinacrine, and Pyronaridine: <i>In Vitro</i> Activity against SARS-CoV-2 and Potential Mechanisms. <i>ACS Omega</i> , 2021, 6, 7454-7468.	3.5	56
6	Machine Learning Models Identify Inhibitors of SARS-CoV-2. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 4224-4235.	5.4	31
7	Recent advances in drug repurposing using machine learning. <i>Current Opinion in Chemical Biology</i> , 2021, 65, 74-84.	6.1	28
8	Machine Learning for Discovery of GSK3 β Inhibitors. <i>ACS Omega</i> , 2020, 5, 26551-26561.	3.5	22
9	New targets for HIV drug discovery. <i>Drug Discovery Today</i> , 2019, 24, 1139-1147.	6.4	18
10	Development of Machine Learning Models and the Discovery of a New Antiviral Compound against Yellow Fever Virus. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 3804-3813.	5.4	16
11	The Antiviral Drug Tilorone Is a Potent and Selective Inhibitor of Acetylcholinesterase. <i>Chemical Research in Toxicology</i> , 2021, 34, 1296-1307.	3.3	15
12	Discovery and Development of Cyclic Peptide Inhibitors of CIB1. <i>ACS Medicinal Chemistry Letters</i> , 2021, 12, 1832-1839.	2.8	14
13	Pyronaridine Protects against SARS-CoV-2 Infection in Mouse. <i>ACS Infectious Diseases</i> , 2022, 8, 1147-1160.	3.8	14
14	Discovery and Characterization of Peptide Inhibitors for Calcium and Integrin Binding Protein 1. <i>ACS Chemical Biology</i> , 2020, 15, 1505-1516.	3.4	11
15	Using Bibliometric Analysis and Machine Learning to Identify Compounds Binding to Sialidase-1. <i>ACS Omega</i> , 2021, 6, 3186-3193.	3.5	11
16	Computational Approaches to Identify Molecules Binding to Mycobacterium tuberculosis KasA. <i>ACS Omega</i> , 2020, 5, 29935-29942.	3.5	8
17	Repurposing the Dihydropyridine Calcium Channel Inhibitor Nicardipine as a Nav1.8 Inhibitor <i>In Vivo</i> for Pitt Hopkins Syndrome. <i>Pharmaceutical Research</i> , 2020, 37, 127.	3.5	7
18	Crystallographic structure and molecular dynamics simulations of the major endoglucanase from <i>Xanthomonas campestris</i> pv. <i>campestris</i> shed light on its oligosaccharide products release pattern. <i>International Journal of Biological Macromolecules</i> , 2019, 136, 493-502.	7.5	5

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19	Crystallization and preliminary diffraction analysis of the catalytic domain of major extracellular endoglucanase from <i>Xanthomonas campestris</i> pv. <i>campestris</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2013, 69, 137-140.	0.7	2
20	Devising effective enzyme replacement therapy for infantile onset neuronal ceroid lipofuscinosis (CLN1 disease). <i>Molecular Genetics and Metabolism</i> , 2021, 132, S28.	1.1	2
21	Structural dataset for the PPAR α V290M mutant. <i>Data in Brief</i> , 2016, 7, 1430-1437.	1.0	1
22	Industrializing enzyme replacement therapy development. <i>Molecular Genetics and Metabolism</i> , 2019, 126, S120.	1.1	0
23	Developing an enzyme replacement therapy and small molecule chaperone combination as a potential treatment for multiple sulfatase deficiency. <i>Molecular Genetics and Metabolism</i> , 2019, 126, S120-S121.	1.1	0