

Milon Mondal

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3168119/publications.pdf>

Version: 2024-02-01

19
papers

1,002
citations

567281

15
h-index

752698

20
g-index

23
all docs

23
docs citations

23
times ranked

1462
citing authors

#	ARTICLE	IF	CITATIONS
1	UCHL1 as a novel target in breast cancer: emerging insights from cell and chemical biology. <i>British Journal of Cancer</i> , 2022, 126, 24-33.	6.4	29
2	Photochemical Probe Identification of a Small-Molecule Inhibitor Binding Site in Hedgehog Acyltransferase (HHAT)**. <i>Angewandte Chemie</i> , 2021, 133, 13654-13659.	2.0	0
3	Photochemical Probe Identification of a Small-Molecule Inhibitor Binding Site in Hedgehog Acyltransferase (HHAT)**. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 13542-13547.	13.8	18
4	Light-mediated discovery of surfaceome nanoscale organization and intercellular receptor interaction networks. <i>Nature Communications</i> , 2021, 12, 7036.	12.8	33
5	Discovery of a Potent and Selective Covalent Inhibitor and Activity-Based Probe for the Deubiquitylating Enzyme UCHL1, with Antifibrotic Activity. <i>Journal of the American Chemical Society</i> , 2020, 142, 12020-12026.	13.7	51
6	Recent Developments in Cell Permeable Deubiquitinating Enzyme Activity-Based Probes. <i>Frontiers in Chemistry</i> , 2019, 7, 876.	3.6	25
7	Chimeric peptidomimetic antibiotics against Gram-negative bacteria. <i>Nature</i> , 2019, 576, 452-458.	27.8	231
8	A Peptidomimetic Antibiotic Interacts with the Periplasmic Domain of LptD from <i>Pseudomonas aeruginosa</i> . <i>ACS Chemical Biology</i> , 2018, 13, 666-675.	3.4	68
9	Thanatin targets the intermembrane protein complex required for lipopolysaccharide transport in <i>Escherichia coli</i> . <i>Science Advances</i> , 2018, 4, eaau2634.	10.3	109
10	Design and Synthesis of Bioisosteres of Acylhydrazones as Stable Inhibitors of the Aspartic Protease Endothiapepsin. <i>ChemMedChem</i> , 2018, 13, 2266-2270.	3.2	7
11	Fragment-Based Drug Design Facilitated by Protein-Templated Click Chemistry: Fragment Linking and Optimization of Inhibitors of the Aspartic Protease Endothiapepsin. <i>Chemistry - A European Journal</i> , 2016, 22, 14826-14830.	3.3	16
12	Furoates and thenoates inhibit pyruvate dehydrogenase kinase 2 allosterically by binding to its pyruvate regulatory site. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2016, 31, 170-175.	5.2	4
13	Fragment Linking and Optimization of Inhibitors of the Aspartic Protease Endothiapepsin: Fragment-Based Drug Design Facilitated by Dynamic Combinatorial Chemistry. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 9422-9426.	13.8	55
14	Fragmentverknüpfung und Optimierung von Hemmstoffen der Aspartylprotease Endothiapepsin: Fragmentbasiertes Wirkstoffdesign beschleunigt durch dynamische kombinatorische Chemie. <i>Angewandte Chemie</i> , 2016, 128, 9569-9574.	2.0	21
15	Structure-Based Optimization of Inhibitors of the Aspartic Protease Endothiapepsin. <i>International Journal of Molecular Sciences</i> , 2015, 16, 19184-19194.	4.1	13
16	Fragment growing exploiting dynamic combinatorial chemistry of inhibitors of the aspartic protease endothiapepsin. <i>MedChemComm</i> , 2015, 6, 1267-1271.	3.4	19
17	Fighting Malaria: Structure-Guided Discovery of Nonpeptidomimetic Plasmepepsin Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2015, 58, 5151-5163.	6.4	24
18	Dynamic combinatorial chemistry: a tool to facilitate the identification of inhibitors for protein targets. <i>Chemical Society Reviews</i> , 2015, 44, 2455-2488.	38.1	176

#	ARTICLE	IF	CITATIONS
19	Structure-Based Design of Inhibitors of the Aspartic Protease Endothiapepsin by Exploiting Dynamic Combinatorial Chemistry. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 3259-3263.	13.8	71