

Lars Richter

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6901697/publications.pdf>

Version: 2024-02-01

14
papers

615
citations

933447

10
h-index

1058476

14
g-index

14
all docs

14
docs citations

14
times ranked

1171
citing authors

#	ARTICLE	IF	CITATIONS
1	Diazepam-bound GABAA receptor models identify new benzodiazepine binding-site ligands. <i>Nature Chemical Biology</i> , 2012, 8, 455-464.	8.0	175
2	Kinetics for Drug Discovery: an industry-driven effort to target drug residence time. <i>Drug Discovery Today</i> , 2017, 22, 896-911.	6.4	165
3	An Updated Unified Pharmacophore Model of the Benzodiazepine Binding Site on γ -Aminobutyric Acid Receptors: Correlation with Comparative Models. <i>Current Medicinal Chemistry</i> , 2007, 14, 2755-2775.	2.4	68
4	Ligand Desolvation Steers On-Rate and Impacts Drug Residence Time of Heat Shock Protein 90 (Hsp90) Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 4397-4411.	6.4	37
5	The Application of the Open Pharmacological Concepts Triple Store (Open PHACTS) to Support Drug Discovery Research. <i>PLoS ONE</i> , 2014, 9, e115460.	2.5	31
6	Molecular analysis of the site for 2-arachidonylglycerol (2-AG) on the γ 2 subunit of GABA _A receptors. <i>Journal of Neurochemistry</i> , 2013, 126, 29-36.	3.9	26
7	Medicinal chemistry in the era of big data. <i>Drug Discovery Today: Technologies</i> , 2015, 14, 37-41.	4.0	24
8	Moving targets in drug discovery. <i>Scientific Reports</i> , 2020, 10, 20213.	3.3	23
9	Structure based classification for bile salt export pump (BSEP) inhibitors using comparative structural modeling of human BSEP. <i>Journal of Computer-Aided Molecular Design</i> , 2017, 31, 507-521.	2.9	20
10	Exploiting open data: a new era in pharmacoinformatics. <i>Future Medicinal Chemistry</i> , 2014, 6, 503-514.	2.3	18
11	Topliss Batchwise Schemes Reviewed in the Era of Open Data Reveal Significant Differences between Enzymes and Membrane Receptors. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 2575-2583.	5.4	9
12	A structure-kinetic relationship study using matched molecular pair analysis. <i>RSC Medicinal Chemistry</i> , 2020, 11, 1285-1294.	3.9	9
13	A residue close to γ 1 loop F disrupts modulation of GABA _A receptors by benzodiazepines while their binding is maintained. <i>Journal of Neurochemistry</i> , 2010, 115, 1478-1485.	3.9	5
14	SAR-Guided Scoring Function and Mutational Validation Reveal the Binding Mode of CGS-8216 at the γ 1/ γ 2 Benzodiazepine Site. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 1682-1696.	5.4	5