

Vladimir Chupakhin

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

608
citations

840585

11
h-index

752573

20
g-index

23
all docs

23
docs citations

23
times ranked

1047
citing authors

#	ARTICLE	IF	CITATIONS
1	Development and implementation of an enterprise-wide predictive model for early absorption, distribution, metabolism and excretion properties. <i>Future Medicinal Chemistry</i> , 2021, 13, 1639-1654.	1.1	7
2	Industry-scale application and evaluation of deep learning for drug target prediction. <i>Journal of Cheminformatics</i> , 2020, 12, 26.	2.8	23
3	Repurposing High-Throughput Image Assays Enables Biological Activity Prediction for Drug Discovery. <i>Cell Chemical Biology</i> , 2018, 25, 611-618.e3.	2.5	176
4	HyperLoom. , 2018, , .		13
5	HyperLoom Possibilities for Executing Scientific Workflows on the Cloud. <i>Advances in Intelligent Systems and Computing</i> , 2018, , 397-406.	0.5	2
6	Discovery of <i>N</i> -(Pyridin-4-yl)-1,5-naphthyridin-2-amines as Potential Tau Pathology PET Tracers for Alzheimer's Disease. <i>Journal of Medicinal Chemistry</i> , 2017, 60, 1272-1291.	2.9	31
7	ExCAPE-DB: an integrated large scale dataset facilitating Big Data analysis in chemogenomics. <i>Journal of Cheminformatics</i> , 2017, 9, 17.	2.8	116
8	Macau: Scalable Bayesian factorization with high-dimensional side information using MCMC. , 2017, , .		23
9	Synthesis of novel bridged dinitrogen heterocycles and their evaluation as potential fragments for the design of biologically active compounds. <i>Tetrahedron</i> , 2014, 70, 7854-7864.	1.0	18
10	Simple Ligand-Receptor Interaction Descriptor (SILIRID) for alignment-free binding site comparison. <i>Computational and Structural Biotechnology Journal</i> , 2014, 10, 33-37.	1.9	25
11	Asymmetric synthesis and molecular docking study of enantiomerically pure pyrrolidine derivatives with potential antithrombin activity. <i>Tetrahedron: Asymmetry</i> , 2013, 24, 838-843.	1.8	15
12	Determination of binding points of methylene blue and cationic phenoxazine dyes on human butyrylcholinesterase. <i>Archives of Biochemistry and Biophysics</i> , 2013, 532, 32-38.	1.4	10
13	Synthesis of PDE IV inhibitors. First asymmetric synthesis of two of GlaxoSmithKline's highly potent Rolipram analogues. <i>Organic and Biomolecular Chemistry</i> , 2013, 11, 8082.	1.5	17
14	Predicting Ligand Binding Modes from Neural Networks Trained on Protein-Ligand Interaction Fingerprints. <i>Journal of Chemical Information and Modeling</i> , 2013, 53, 763-772.	2.5	47
15	Design of small-molecule thrombin inhibitors based on the cis-5-phenylproline scaffold. <i>Russian Chemical Bulletin</i> , 2011, 60, 685-693.	0.4	7
16	Design, synthesis and biotest of a bicyclo[3.3.1]nonane analogue of 2-amino-5,6-dihydro-4H-1,3-thiazine. <i>Mendeleev Communications</i> , 2010, 20, 323-325.	0.6	18
17	Ionotropic GABA receptors: modelling and design of selective ligands. <i>Journal of Cheminformatics</i> , 2010, 2, .	2.8	0
18	Molecular modeling of ligand-receptor interactions in GABAC receptor. <i>Journal of Molecular Graphics and Modelling</i> , 2009, 27, 813-821.	1.3	21

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19	Computer-aided design of selective ligands of the benzodiazepine-binding site of the GABAA receptor. Doklady Chemistry, 2008, 422, 227-230.	0.2	7
20	Modeling and analysis of ligand-receptor interactions in the GABAC receptor. Doklady Biochemistry and Biophysics, 2007, 412, 25-28.	0.3	7
21	Design of potential NO-synthase inhibitors on the basis of 2-amino-5,6-dihydro-4H-1,3-thiazine derivatives. Moscow University Chemistry Bulletin, 2007, 62, 243-245.	0.2	4
22	Modeling the open and closed forms of GABAA receptor: Analysis of ligand-receptor interactions for the GABA-binding site. Doklady Biochemistry and Biophysics, 2006, 408, 169-174.	0.3	10