

Li Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

320
citations

933447

10
h-index

888059

17
g-index

17
all docs

17
docs citations

17
times ranked

350
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of 1-phenyl-4-cyano-5-aminopyrazoles as novel ecdysone receptor ligands by virtual screening, structural optimization, and biological evaluations. <i>Chemical Biology and Drug Design</i> , 2021, 97, 184-195.	3.2	4
2	New lead discovery of insect growth regulators based on the scaffold hopping strategy. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2020, 30, 127500.	2.2	12
3	Design, Synthesis, and Biological Activity of Novel Heptacyclic Pyrazolamide Derivatives: A New Candidate of Dual-Target Insect Growth Regulators. <i>Journal of Agricultural and Food Chemistry</i> , 2020, 68, 6347-6354.	5.2	22
4	Virtual screening, synthesis, and bioactivity evaluation for the discovery of \hat{I}^2 -N-acetyl-D-hexosaminidase inhibitors. <i>Pest Management Science</i> , 2020, 76, 3030-3037.	3.4	9
5	Computational Screening of Potential Inhibitors of \hat{I}^2 -N-Acetyl-D-Hesosaminidases Using Combined Core-Fragment Growth and Pharmacophore Restraints. <i>Applied Biochemistry and Biotechnology</i> , 2019, 189, 1262-1273.	2.9	5
6	Pocket-based Lead Optimization Strategy for the Design and Synthesis of Chitinase Inhibitors. <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 3575-3582.	5.2	24
7	Insights into the structure-affinity relationships and solvation effects between OfHex1 and inhibitors using molecular dynamics simulations. <i>Journal of Molecular Graphics and Modelling</i> , 2019, 90, 1-8.	2.4	4
8	Computational Study for the Unbinding Routes of \hat{I}^2 -N-Acetyl-d-Hexosaminidase Inhibitor: Insight from Steered Molecular Dynamics Simulations. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1516.	4.1	4
9	Identification of novel agonists and antagonists of the ecdysone receptor by virtual screening. <i>Journal of Molecular Graphics and Modelling</i> , 2018, 81, 77-85.	2.4	14
10	Structure-Based Virtual Screening, Compound Synthesis, and Bioassay for the Design of Chitinase Inhibitors. <i>Journal of Agricultural and Food Chemistry</i> , 2018, 66, 3351-3357.	5.2	45
11	Steered molecular dynamics for studying ligand unbinding of ecdysone receptor. <i>Journal of Biomolecular Structure and Dynamics</i> , 2018, 36, 3819-3828.	3.5	17
12	Functional recognition of structure-diverse odor molecules in drinking water based on QSOR study. <i>Chemosphere</i> , 2018, 211, 371-378.	8.2	1
13	Exploration of the binding affinities between ecdysone agonists and EcR/USP by docking and MM-PB/GBSA approaches. <i>Journal of Molecular Modeling</i> , 2017, 23, 166.	1.8	8
14	Design, synthesis and biological activity of novel substituted pyrazole amide derivatives targeting EcR/USP receptor. <i>Chinese Chemical Letters</i> , 2016, 27, 566-570.	9.0	19
15	Target-based design, synthesis and biological activity of new pyrazole amide derivatives. <i>Chinese Chemical Letters</i> , 2016, 27, 251-255.	9.0	19
16	Understanding the Mechanism of Drug Resistance Due to a Codon Deletion in Protoporphyrinogen Oxidase through Computational Modeling. <i>Journal of Physical Chemistry B</i> , 2009, 113, 4865-4875.	2.6	47
17	Quantitative Structure-Activity Relationship for Cyclic Imide Derivatives of Protoporphyrinogen Oxidase Inhibitors: A Study of Quantum Chemical Descriptors from Density Functional Theory. <i>Journal of Chemical Information and Computer Sciences</i> , 2004, 44, 2099-2105.	2.8	66