

Liangren Zhang

List of Publications by Year in descending order

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1,839
citations

236833

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#	ARTICLE	IF	CITATIONS
1	Discovery of novel ataxia telangiectasia mutated (ATM) kinase modulators: Computational simulation, biological evaluation and cancer combinational chemotherapy study. <i>European Journal of Medicinal Chemistry</i> , 2022, 233, 114196.	2.6	4
2	Optimization of 4-arylthiophene-3-carboxylic acid derivatives as inhibitors of ANO1: Lead optimization studies toward their analgesic efficacy for inflammatory pain. <i>European Journal of Medicinal Chemistry</i> , 2022, 237, 114413.	2.6	5
3	CMNPD: a comprehensive marine natural products database towards facilitating drug discovery from the ocean. <i>Nucleic Acids Research</i> , 2021, 49, D509-D515.	6.5	105
4	Discovery of 4-arylthiophene-3-carboxylic acid as inhibitor of ANO1 and its effect as analgesic agent. <i>Acta Pharmaceutica Sinica B</i> , 2021, 11, 1947-1964.	5.7	13
5	Vacuolin-1 inhibits endosomal trafficking and metastasis via CapZ ² . <i>Oncogene</i> , 2021, 40, 1775-1791.	2.6	14
6	The Discovery of Novel ACA Derivatives as Specific TRPM2 Inhibitors that Reduce Ischemic Injury Both In Vitro and In Vivo. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 3976-3996.	2.9	16
7	Target Prediction Model for Natural Products Using Transfer Learning. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4632.	1.8	9
8	Discovery of Novel and Potent <i>N</i> -Methyl- <i>D</i> -aspartate Receptor Positive Allosteric Modulators with Antidepressant-like Activity in Rodent Models. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 5551-5576.	2.9	12
9	Chemical Space, Scaffolds, and Halogenated Compounds of CMNPD: A Comprehensive Chemoinformatic Analysis. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 3323-3336.	2.5	4
10	Synthesis and Biological Evaluation of Novel Triazine Derivatives as Positive Allosteric Modulators of $\alpha 7$ Nicotinic Acetylcholine Receptors. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 12379-12396.	2.9	6
11	Functional Characterization and Crystal Structure of the Bifunctional Thioesterase Catalyzing Epimerization and Cyclization in Skyllamycin Biosynthesis. <i>ACS Catalysis</i> , 2021, 11, 11733-11741.	5.5	8
12	Design, synthesis and biological activities of benzo[d]imidazo[1,2-a]imidazole derivatives as TRPM2-specific inhibitors. <i>European Journal of Medicinal Chemistry</i> , 2021, 225, 113750.	2.6	3
13	A unique ligand-steered strategy for CC chemokine receptor 2 homology modeling to facilitate structure-based virtual screening. <i>Chemical Biology and Drug Design</i> , 2021, 97, 944-961.	1.5	3
14	Challenging Reverse Screening: A Benchmark Study for Comprehensive Evaluation. <i>Molecular Informatics</i> , 2021, , 2100063.	1.4	1
15	DeepScaffold: A Comprehensive Tool for Scaffold-Based De Novo Drug Discovery Using Deep Learning. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 77-91.	2.5	84
16	Design, synthesis and biological activities of piperidine-spirooxadiazole derivatives as $\alpha 7$ nicotinic receptor antagonists. <i>European Journal of Medicinal Chemistry</i> , 2020, 207, 112774.	2.6	8
17	Discovery of 2-(2-aminobenzo[d]thiazol-6-yl) benzo[d]oxazol-5-amine derivatives that regulated HPV relevant cellular pathway and prevented cervical cancer from abnormal proliferation. <i>European Journal of Medicinal Chemistry</i> , 2020, 204, 112556.	2.6	2
18	Medicinal chemistry perspective of TRPM2 channel inhibitors: where we are and where we might be heading?. <i>Drug Discovery Today</i> , 2020, 25, 2326-2334.	3.2	10

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19	TF3P: Three-Dimensional Force Fields Fingerprint Learned by Deep Capsular Network. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 2754-2765.	2.5	9
20	Privileged Scaffold Analysis of Natural Products with Deep Learning-based Indication Prediction Model. <i>Molecular Informatics</i> , 2020, 39, e2000057.	1.4	10
21	Rational modification, synthesis and biological evaluation of 3,4-dihydroquinoxalin-2(1H)-one derivatives as potent and selective c-Jun N-terminal kinase 3 (JNK3) inhibitors. <i>European Journal of Medicinal Chemistry</i> , 2020, 201, 112445.	2.6	17
22	Pose Filter-Based Ensemble Learning Enables Discovery of Orally Active, Nonsteroidal Farnesoid X Receptor Agonists. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 1202-1214.	2.5	7
23	Discovery and antibacterial study of potential PPK1 inhibitors against uropathogenic <i>E. coli</i> . <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2020, 35, 1224-1232.	2.5	12
24	Discovery of fused heterocyclic carboxamide derivatives as novel α -7-nAChR agonists: Synthesis, preliminary SAR and biological evaluation. <i>European Journal of Medicinal Chemistry</i> , 2019, 182, 111618.	2.6	7
25	Multistage Screening Reveals 3-Substituted Indolin-2-one Derivatives as Novel and Isoform-Selective c-Jun N-terminal Kinase 3 (JNK3) Inhibitors: Implications to Drug Discovery for Potential Treatment of Neurodegenerative Diseases. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 6645-6664.	2.9	38
26	Identification and characterization of benzo[d]oxazol-2(3H)-one derivatives as the first potent and selective small-molecule inhibitors of chromodomain protein CDYL. <i>European Journal of Medicinal Chemistry</i> , 2019, 182, 111656.	2.6	11
27	Direct Gating of the TRPM2 Channel by cADPR via Specific Interactions with the ADPR Binding Pocket. <i>Cell Reports</i> , 2019, 27, 3684-3695.e4.	2.9	45
28	Rational Design and Identification of Small-Molecule Allosteric Inhibitors of CD38. <i>ChemBioChem</i> , 2019, 20, 2485-2493.	1.3	6
29	Discovery of novel glycogen synthase kinase-3 β inhibitors: Structure-based virtual screening, preliminary SAR and biological evaluation for treatment of acute myeloid leukemia. <i>European Journal of Medicinal Chemistry</i> , 2019, 171, 221-234.	2.6	19
30	Graph-based generative models for de Novo drug design. <i>Drug Discovery Today: Technologies</i> , 2019, 32-33, 45-53.	4.0	15
31	Design, synthesis and biological activities of 2,3-dihydroquinazolin-4(1H)-one derivatives as TRPM2 inhibitors. <i>European Journal of Medicinal Chemistry</i> , 2018, 152, 235-252.	2.6	29
32	Calcium-Mobilizing Behaviors of Neutral Cyclic ADP-Ribose Mimics that Integrate Modifications to the Nucleobase, Northern Ribose and Pyrophosphate. <i>ChemBioChem</i> , 2018, 19, 1444-1451.	1.3	3
33	Maximal Unbiased Benchmarking Data Sets for Human Chemokine Receptors and Comparative Analysis. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 1104-1120.	2.5	4
34	Selective inhibition of TRPM2 channel by two novel synthesized ADPR analogues. <i>Chemical Biology and Drug Design</i> , 2018, 91, 552-566.	1.5	27
35	Discovery of new GSK-3 β inhibitors through structure-based virtual screening. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2018, 28, 160-166.	1.0	19
36	d-Isothymidine incorporation in the core sequence of aptamer BC15 enhanced its binding affinity to the hnRNP A1 protein. <i>Organic and Biomolecular Chemistry</i> , 2018, 16, 7488-7497.	1.5	5

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37	Synthesis and Biological Evaluation of Fused Tricyclic Heterocycle Piperazine (Piperidine) Derivatives As Potential Multireceptor Atypical Antipsychotics. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 10017-10039.	2.9	39
38	Multi-objective de novo drug design with conditional graph generative model. <i>Journal of Cheminformatics</i> , 2018, 10, 33.	2.8	193
39	Discovery of N-(Naphtho[1,2-b]Furan-5-Yl) Benzenesulfonamides as Novel Selective Inhibitors of Triple-Negative Breast Cancer (TNBC). <i>Molecules</i> , 2018, 23, 678.	1.7	4
40	Scale-Up Synthesis and Identification of GLYX-13, a NMDAR Glycine-Site Partial Agonist for the Treatment of Major Depressive Disorder. <i>Molecules</i> , 2018, 23, 996.	1.7	3
41	Identification of the ADPR binding pocket in the NUDT9 homology domain of TRPM2. <i>Journal of General Physiology</i> , 2017, 149, 219-235.	0.9	49
42	Design, synthesis, and biological evaluation of imidazo[1,2-b]pyridazine derivatives as mTOR inhibitors. <i>European Journal of Medicinal Chemistry</i> , 2017, 129, 135-150.	2.6	24
43	Identifying Glyceraldehyde 3-Phosphate Dehydrogenase as a Cyclic Adenosine Diphosphoribose Binding Protein by Photoaffinity Proteinâ€“Ligand Labeling Approach. <i>Journal of the American Chemical Society</i> , 2017, 139, 156-170.	6.6	30
44	Identification of Novel Vacuolin-1 Analogues as Autophagy Inhibitors by Virtual Drug Screening and Chemical Synthesis. <i>Molecules</i> , 2017, 22, 891.	1.7	17
45	The scoring bias in reverse docking and the score normalization strategy to improve success rate of target fishing. <i>PLoS ONE</i> , 2017, 12, e0171433.	1.1	19
46	Synthesis and biological activities of indolizine derivatives as alpha-7 nAChR agonists. <i>European Journal of Medicinal Chemistry</i> , 2016, 115, 94-108.	2.6	35
47	Cyclic Adenosine 5â€“Diphosphoribose (cADPR) Mimics Used as Molecular Probes in Cell Signaling. <i>Chemical Record</i> , 2015, 15, 511-523.	2.9	4
48	Comparative Analysis of Pharmacophore Features and Quantitative Structureâ€“Activity Relationships for <sc>CD</sc>38 Covalent and Nonâ€“covalent Inhibitors. <i>Chemical Biology and Drug Design</i> , 2015, 86, 1411-1424.	1.5	17
49	Enrichment Assessment of Multiple Virtual Screening Strategies for Tollâ€“like Receptor 8 Agonists Based on a Maximal Unbiased Benchmarking Data Set. <i>Chemical Biology and Drug Design</i> , 2015, 86, 1226-1241.	1.5	11
50	Integrating Pharmacophore into Membrane Molecular Dynamics Simulations to Improve Homology Modeling of <sc>G</sc> Proteinâ€“coupled Receptors with Ligand Selectivity: A_{2A} Adenosine Receptor as an Example. <i>Chemical Biology and Drug Design</i> , 2015, 86, 1438-1450.	1.5	7
51	Benchmarking methods and data sets for ligand enrichment assessment in virtual screening. <i>Methods</i> , 2015, 71, 146-157.	1.9	40
52	An Unbiased Method To Build Benchmarking Sets for Ligand-Based Virtual Screening and its Application To GPCRs. <i>Journal of Chemical Information and Modeling</i> , 2014, 54, 1433-1450.	2.5	46
53	Stability and bioactivity of thrombin binding aptamers modified with <sc>d</sc>-<sc>l</sc>-isothymidine in the loop regions. <i>Organic and Biomolecular Chemistry</i> , 2014, 12, 8866-8876.	1.5	38
54	Unfolding and Conformational Variations of Thrombinâ€“Binding DNA Aptamers: Synthesis, Circular Dichroism and Molecular Dynamics Simulations. <i>ChemMedChem</i> , 2014, 9, 993-1001.	1.6	15

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55	Design, Synthesis and SAR Studies of NAD Analogues as Potent Inhibitors towards CD38 NADase. <i>Molecules</i> , 2014, 19, 15754-15767.	1.7	26
56	Design, synthesis and biological evaluation of indolizine derivatives as HIV-1 Vif-ElonginC interaction inhibitors. <i>Molecular Diversity</i> , 2013, 17, 221-243.	2.1	41
57	Computational Identification of a New Binding Site in Influenza Virus Hemagglutinin for Membrane Fusion Inhibitors. <i>Chemical Biology and Drug Design</i> , 2013, 82, 267-274.	1.5	1
58	Indolizine Derivatives as HIV-1 Vif-ElonginC Interaction Inhibitors. <i>Chemical Biology and Drug Design</i> , 2013, 81, 730-741.	1.5	46
59	Computational investigation of interactions between Cdc37 and celastrol. <i>Molecular Simulation</i> , 2013, 39, 270-278.	0.9	4
60	Toll-Like Receptor 7 Agonists: Chemical Feature Based Pharmacophore Identification and Molecular Docking Studies. <i>PLoS ONE</i> , 2013, 8, e56514.	1.1	14
61	Small-Molecule Inhibition of Human Immunodeficiency Virus Type 1 Replication by Targeting the Interaction between Vif and ElonginC. <i>Journal of Virology</i> , 2012, 86, 5497-5507.	1.5	63
62	A Novel Fluorescent Cell Membrane-permeable Caged Cyclic ADP-ribose Analogue. <i>Journal of Biological Chemistry</i> , 2012, 287, 24774-24783.	1.6	27
63	Synthesis and Calcium Mobilization Activity of cADPR Analogues Which Integrate Nucleobase, Northern and Southern Ribose Modifications. <i>Molecules</i> , 2012, 17, 4343-4356.	1.7	7
64	Catalysis-Based Inhibitors of the Calcium Signaling Function of CD38. <i>Biochemistry</i> , 2012, 51, 555-564.	1.2	31
65	A Cell Permeable NPE Caged ADP-Ribose for Studying TRPM2. <i>PLoS ONE</i> , 2012, 7, e51028.	1.1	23
66	Design, Synthesis and Biological Evaluation of Noncovalent Inhibitors of Human CD38 NADase. <i>ChemMedChem</i> , 2012, 7, 223-228.	1.6	21
67	Design and synthesis of cADPR analogues with simplified ribose and nucleobase. <i>Journal of Chinese Pharmaceutical Sciences</i> , 2012, 21, .	0.4	2
68	Novel nucleobase-simplified cyclic ADP-ribose analogue: A concise synthesis and Ca ²⁺ -mobilizing activity in T-lymphocytes. <i>Organic and Biomolecular Chemistry</i> , 2010, 8, 1843.	1.5	16
69	Concise Syntheses of Trifluoromethylated Cyclic and Acyclic Analogues of cADPR. <i>Molecules</i> , 2010, 15, 8689-8701.	1.7	11
70	Structural basis for the specific interaction of chicken haemoglobin with bromophenol blue: a computational analysis. <i>Molecular Physics</i> , 2010, 108, 215-220.	0.8	1
71	Covalent complexes of proteasome model with peptide aldehyde inhibitors MG132 and MG101: docking and molecular dynamics study. <i>Journal of Molecular Modeling</i> , 2009, 15, 1481-1490.	0.8	30
72	An efficient synthesis of (R)- and (S)-baclofen via desymmetrization. <i>Tetrahedron Letters</i> , 2009, 50, 6166-6168.	0.7	24

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73	Three-dimensional structure of HIV-1 VIF constructed by comparative modeling and the function characterization analyzed by molecular dynamics simulation. <i>Organic and Biomolecular Chemistry</i> , 2007, 5, 617.	1.5	30
74	Synthesis and recognition of novel isonucleoside triphosphates by DNA polymerases. <i>Bioorganic and Medicinal Chemistry</i> , 2007, 15, 3019-3025.	1.4	23
75	Synthesis and Agonist Activity of Cyclic ADP-Ribose Analogues with Substitution of the Northern Ribose by Ether or Alkane Chains. <i>Journal of Medicinal Chemistry</i> , 2006, 49, 5501-5512.	2.9	35
76	Structural insights into the effect of isonucleosides on B-DNA duplexes using molecular-dynamics simulations. <i>Journal of Molecular Modeling</i> , 2006, 12, 781-791.	0.8	6
77	Analysis of the interactions of ribonuclease inhibitor with kanamycin. <i>Journal of Molecular Modeling</i> , 2005, 11, 80-86.	0.8	3
78	A Minimal Structural Analogue of Cyclic ADP-ribose. <i>Journal of Biological Chemistry</i> , 2005, 280, 15952-15959.	1.6	40
79	Amplification and propagation of pacemaker Ca ²⁺ signals by cyclic ADP-ribose and the type 3 ryanodine receptor in T cells. <i>Journal of Cell Science</i> , 2004, 117, 2141-2149.	1.2	41
80	Synthesis and Biological Evaluation of Novel Membrane-Permeant Cyclic ADP-Ribose Mimics: AN1-[(5'-O-Phosphorylethoxy)methyl]-5'-O-phosphorylino-5',5'-Cyclicpyrophosphate (CIDPRE) and 8-Substituted Derivatives. <i>Journal of Medicinal Chemistry</i> , 2004, 47, 5674-5682.	2.0	51
81	Solid-Phase Synthesis of Dipeptide-Conjugated Nucleosides and Their Interaction with RNA. <i>Helvetica Chimica Acta</i> , 2003, 86, 3516-3524.	1.0	8
82	Synthesis of 2-amino-2-deoxy- β -D-glycosyl-(1 \rightarrow 5)-nucleosides and the interaction with RNA. <i>Bioorganic and Medicinal Chemistry</i> , 2003, 11, 3273-3278.	1.4	10
83	Synthesis Of Spin Labeled Conjugate Of Peptide and Peptide Nucleic Acid. <i>Synthetic Communications</i> , 1999, 29, 1519-1525.	1.1	3
84	Synthesis and Duplex Stabilization of Oligonucleotides Consisting of Isonucleosides. <i>Helvetica Chimica Acta</i> , 1999, 82, 2037-2043.	1.0	19
85	8-chloroadenosine induces apoptosis in human MOLT-4 cell line. <i>Science Bulletin</i> , 1997, 42, 592-597.	1.7	0
86	Assignments of Nonexchangeable Proton Resonances and the Solution Structures of d ⁵ TGGGT. <i>Spectroscopy Letters</i> , 1993, 26, 1537-1546.	0.5	1