

Dan-feng Shi

List of Publications by Year in descending order

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times ranked

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#	ARTICLE	IF	CITATIONS
1	Two new iridoid glycosides from the fruit of <i>Gardenia jasminoides</i> . <i>Natural Product Research</i> , 2022, 36, 186-192.	1.8	9
2	Identifying the molecular basis of Jinhong tablets against chronic superficial gastritis via chemical profile identification and symptom-guided network pharmacology analysis. <i>Journal of Pharmaceutical Analysis</i> , 2022, 12, 65-76.	5.3	8
3	Three new cycloart-7-ene triterpenoid glycosides from <i>Cimicifuga dahurica</i> and their anti-inflammatory effects. <i>Natural Product Research</i> , 2021, 35, 3634-3643.	1.8	7
4	A new nitrogen-containing iridoid glycoside from <i>Lonicera macranthoides</i> . <i>Natural Product Research</i> , 2021, 35, 3432-3438.	1.8	3
5	Metabolic profiles of Jinhong tablets in rats by ultra-performance liquid chromatography coupled with quadrupole time-of-flight tandem mass spectrometry. <i>Biomedical Chromatography</i> , 2021, 35, e5072.	1.7	2
6	Discovery of novel IDO1 inhibitors via structure-based virtual screening and biological assays. <i>Journal of Computer-Aided Molecular Design</i> , 2021, 35, 679-694.	2.9	2
7	Systematically identifying the anti-inflammatory constituents of <i>Cimicifuga dahurica</i> by UPLC-Q/TOF-MS combined with network pharmacology analysis. <i>Biomedical Chromatography</i> , 2021, 35, e5177.	1.7	4
8	Arteannoides U: Six undescribed sesquiterpenoids with anti-inflammatory activities from the aerial parts of <i>Artemisia annua</i> (Qinghao). <i>FÄ-toterapÄ-Äç</i> , 2021, 154, 105002.	2.2	4
9	Structural Modification of Aminophenylarsenoxides Generates Candidates for Leukemia Treatment <i>via</i> Thioredoxin Reductase Inhibition. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 16132-16146.	6.4	16
10	4-Hydroxy Pyridones from Heterologous Expression and Cultivation of the Native Host. <i>Journal of Natural Products</i> , 2020, 83, 3338-3346.	3.0	19
11	Computational Insight Into the Small Molecule Intervening PD-L1 Dimerization and the Potential Structure-Activity Relationship. <i>Frontiers in Chemistry</i> , 2019, 7, 764.	3.6	33
12	Fluorophore-Dependent Cleavage of Disulfide Bond Leading to a Highly Selective Fluorescent Probe of Thioredoxin. <i>Analytical Chemistry</i> , 2019, 91, 8524-8531.	6.5	26
13	Deciphering the Allosteric Effect of Antagonist Vismodegib on Smoothed Receptor Deactivation Using Metadynamics Simulation. <i>Frontiers in Chemistry</i> , 2019, 7, 406.	3.6	2
14	pH-Induced Misfolding Mechanism of Prion Protein: Insights from Microsecond-Accelerated Molecular Dynamics Simulations. <i>ACS Chemical Neuroscience</i> , 2019, 10, 2718-2729.	3.5	13
15	Virtual screening-guided discovery of thioredoxin reductase inhibitors. <i>Toxicology and Applied Pharmacology</i> , 2019, 370, 106-116.	2.8	15
16	Phenolic acids and their glycosides from the rhizomes of <i>Cimicifuga dahurica</i> . <i>FÄ-toterapÄ-Äç</i> , 2019, 134, 485-492.	2.2	8
17	Targeting Thioredoxin Reductase by Ibrutinib Promotes Apoptosis of SMMC-7721 Cells. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2019, 369, 212-222.	2.5	10
18	How Does Agonist and Antagonist Binding Lead to Different Conformational Ensemble Equilibria of the μ -Opioid Receptor: Insight from Long-Time Gaussian Accelerated Molecular Dynamics Simulation. <i>ACS Chemical Neuroscience</i> , 2019, 10, 1575-1584.	3.5	9

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19	Computational study on the selective inhibition mechanism of MS402 to the first and second bromodomains of BRD4. <i>Proteins: Structure, Function and Bioinformatics</i> , 2019, 87, 3-11.	2.6	16
20	Computational studies on horseshoe shape pocket of human orexin receptor type 2 and boat conformation of suvorexant by molecular dynamics simulations. <i>Chemical Biology and Drug Design</i> , 2018, 92, 1221-1231.	3.2	4
21	Understanding the structural and energetic basis of PD-1 and monoclonal antibodies bound to PD-L1: A molecular modeling perspective. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2018, 1862, 576-588.	2.4	21
22	Influence of EGCG on α -synuclein (α S) aggregation and identification of their possible binding mode: A computational study using molecular dynamics simulation. <i>Chemical Biology and Drug Design</i> , 2018, 91, 162-171.	3.2	24
23	Molecular dynamics simulation, binding free energy calculation and unbinding pathway analysis on selectivity difference between FKBP51 and FKBP52: Insight into the molecular mechanism of isoform selectivity. <i>Proteins: Structure, Function and Bioinformatics</i> , 2018, 86, 43-56.	2.6	35
24	Molecular dynamics simulations and novel drug discovery. <i>Expert Opinion on Drug Discovery</i> , 2018, 13, 23-37.	5.0	292
25	Investigate the mechanisms of Chinese medicine Fuzhengkangai towards EGFR mutation-positive lung adenocarcinomas by network pharmacology. <i>BMC Complementary and Alternative Medicine</i> , 2018, 18, 293.	3.7	37
26	Xanthatin Promotes Apoptosis via Inhibiting Thioredoxin Reductase and Eliciting Oxidative Stress. <i>Molecular Pharmaceutics</i> , 2018, 15, 3285-3296.	4.6	34
27	Identification of a new pyruvate kinase M2 isoform (PKM2) activator for the treatment of non-small-cell lung cancer (NSCLC). <i>Chemical Biology and Drug Design</i> , 2018, 92, 1851-1858.	3.2	17
28	Revealing inhibition difference between PFI-2 enantiomers against SETD7 by molecular dynamics simulations, binding free energy calculations and unbinding pathway analysis. <i>Scientific Reports</i> , 2017, 7, 46547.	3.3	25
29	Synthesis of naphthazarin derivatives and identification of novel thioredoxin reductase inhibitor as potential anticancer agent. <i>European Journal of Medicinal Chemistry</i> , 2017, 140, 435-447.	5.5	23
30	Protective V127 prion variant prevents prion disease by interrupting the formation of dimer and fibril from molecular dynamics simulations. <i>Scientific Reports</i> , 2016, 6, 21804.	3.3	21
31	Inhibition of Monoamine Oxidase by Stilbenes from <i>Rheum palmatum</i> . <i>Iranian Journal of Pharmaceutical Research</i> , 2016, 15, 885-892.	0.5	6
32	Influence of Chirality of Crizotinib on Its MTH1 Protein Inhibitory Activity: Insight from Molecular Dynamics Simulations and Binding Free Energy Calculations. <i>PLoS ONE</i> , 2015, 10, e0145219.	2.5	20
33	Synthesis and quantitative structure-activity relationship (QSAR) study of C7-oxime ester derivatives of obacunone as insecticidal agents. <i>RSC Advances</i> , 2015, 5, 31700-31707.	3.6	21
34	Ligand induced change of β 2-adrenergic receptor from active to inactive conformation and its implication for the closed/open state of the water channel: insight from molecular dynamics simulation, free energy calculation and Markov state model analysis. <i>Physical Chemistry Chemical Physics</i> , 2014, 16, 15874.	2.8	35
35	Desmodeganine, a new alkaloid from the leaves of <i>Desmodium elegans</i> as a potential monoamine oxidase inhibitor. <i>FÄ-toterapÄ-c</i> , 2014, 98, 160-165.	2.2	28