

Jianping Hu

List of Publications by Year in descending order

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Version: 2024-02-01

48
papers

630
citations

687335

13
h-index

642715

23
g-index

49
all docs

49
docs citations

49
times ranked

981
citing authors

#	ARTICLE	IF	CITATIONS
1	COVID-19 Docking Server: a meta server for docking small molecules, peptides and antibodies against potential targets of COVID-19. <i>Bioinformatics</i> , 2020, 36, 5109-5111.	4.1	122
2	Long Non-coding RNAs: Emerging Roles in the Immunosuppressive Tumor Microenvironment. <i>Frontiers in Oncology</i> , 2020, 10, 48.	2.8	63
3	MAPK-Targeted Drug Delivered by a pH-Sensitive MSNP Nanocarrier Synergizes with PD-1 Blockade in Melanoma without T-Cell Suppression. <i>Advanced Functional Materials</i> , 2019, 29, 1806916.	14.9	34
4	PD-L1 Nanobody Competitively Inhibits the Formation of the PD-1/PD-L1 Complex: Comparative Molecular Dynamics Simulations. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1984.	4.1	31
5	Difference and Influence of Inactive and Active States of Cannabinoid Receptor Subtype CB2: From Conformation to Drug Discovery. <i>Journal of Chemical Information and Modeling</i> , 2016, 56, 1152-1163.	5.4	26
6	Combination of MAPK inhibition with photothermal therapy synergistically augments the anti-tumor efficacy of immune checkpoint blockade. <i>Journal of Controlled Release</i> , 2021, 332, 194-209.	9.9	25
7	Study on the molecular mechanism of inhibiting HIV-1 integrase by EBR28 peptide via molecular modeling approach. <i>Biophysical Chemistry</i> , 2008, 132, 69-80.	2.8	21
8	Metal binding mediated conformational change of XPA protein: a potential cytotoxic mechanism of nickel in the nucleotide excision repair. <i>Journal of Molecular Modeling</i> , 2016, 22, 156.	1.8	18
9	Study on the interactions between diketo-acid inhibitors and prototype foamy virus integrase-DNA complex via molecular docking and comparative molecular dynamics simulation methods. <i>Journal of Biomolecular Structure and Dynamics</i> , 2013, 31, 734-747.	3.5	17
10	Combined 3D-QSAR, molecular docking, and molecular dynamics study of tacrine derivatives as potential acetylcholinesterase (AChE) inhibitors of Alzheimer's disease. <i>Journal of Molecular Modeling</i> , 2015, 21, 277.	1.8	16
11	Integrin-Src-YAP1 signaling mediates the melanoma acquired resistance to MAPK and PI3K/mTOR dual targeted therapy. <i>Molecular Biomedicine</i> , 2020, 1, 12.	4.4	16
12	Efficient whole-cell catalysis for 5-aminovalerate production from L-lysine by using engineered <i>Escherichia coli</i> with ethanol pretreatment. <i>Scientific Reports</i> , 2020, 10, 990.	3.3	16
13	3D-QSAR, Molecular Docking and Molecular Dynamics Simulation of <i>Pseudomonas aeruginosa</i> LpxC Inhibitors. <i>International Journal of Molecular Sciences</i> , 2017, 18, 761.	4.1	15
14	Inhibition of programmed cell death protein ligand-1 (PD-L1) by benzyl ether derivatives: analyses of conformational change, molecular recognition and binding free energy. <i>Journal of Biomolecular Structure and Dynamics</i> , 2019, 37, 4801-4812.	3.5	15
15	Specifically targeting cancer proliferation and metastasis processes: the development of matriptase inhibitors. <i>Cancer and Metastasis Reviews</i> , 2019, 38, 507-524.	5.9	14
16	3D-QSAR and molecular recognition of <i>Klebsiella pneumoniae</i> NDM-1 inhibitors. <i>Molecular Simulation</i> , 2019, 45, 694-705.	2.0	13
17	Concepts and Application of DNA Origami and DNA Self-Assembly: A Systematic Review. <i>Applied Bionics and Biomechanics</i> , 2021, 2021, 1-15.	1.1	12
18	Omicron-included mutation-induced changes in epitopes of SARS-CoV-2 spike protein and effectiveness assessments of current antibodies. <i>Molecular Biomedicine</i> , 2022, 3, 12.	4.4	12

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19	Molecular Dynamics Simulation of HIV-1 Integrase Dimer Complexed with Viral DNA. <i>Chinese Journal of Chemistry</i> , 2010, 28, 33-40.	4.9	11
20	The Evolution of Acquired Resistance to BRAFV600E Kinase inhibitor Is Sustained by IGF1-Driven Tumor Vascular Remodeling. <i>Journal of Investigative Dermatology</i> , 2022, 142, 445-458.	0.7	11
21	Probing the Behaviour of Cas1-Cas2 upon Protospacer Binding in CRISPR-Cas Systems using Molecular Dynamics Simulations. <i>Scientific Reports</i> , 2019, 9, 3188.	3.3	10
22	Structural characterization of the Plasmodium falciparum lactate transporter PffNT alone and in complex with antimalarial compound MMV007839 reveals its inhibition mechanism. <i>PLoS Biology</i> , 2021, 19, e3001386.	5.6	10
23	Substrate Recognition and Motion Mode Analyses of PFV Integrase in Complex with Viral DNA via Coarse-Grained Models. <i>PLoS ONE</i> , 2013, 8, e54929.	2.5	9
24	The global motion affecting electron transfer in <i>Plasmodium falciparum</i> type II NADH dehydrogenases: a novel non-competitive mechanism for quinoline ketone derivative inhibitors. <i>Physical Chemistry Chemical Physics</i> , 2019, 21, 18105-18118.	2.8	9
25	Applications of Molecular Simulation in the Discovery of Antituberculosis Drugs: A Review. <i>Protein and Peptide Letters</i> , 2019, 26, 648-663.	0.9	9
26	BP[dG]-induced distortions to DNA polymerase and DNA duplex: A detailed mechanism of BP adducts blocking replication. <i>Food and Chemical Toxicology</i> , 2020, 140, 111325.	3.6	8
27	Theoretical insight into the photodeactivation pathway of the tetradentate Pt(II) complex: The π -conjugation effect. <i>Applied Organometallic Chemistry</i> , 2018, 32, e4220.	3.5	7
28	Theoretical insight into the photodeactivation pathway of the tetradentate Pt (II) complex with different inductive substituents. <i>Applied Organometallic Chemistry</i> , 2019, 33, e4879.	3.5	7
29	Classification and Design of HIV-1 Integrase Inhibitors Based on Machine Learning. <i>Computational and Mathematical Methods in Medicine</i> , 2021, 2021, 1-11.	1.3	6
30	C- and N-Adducts of N-alkenyl substituted Arduengo carbene and N-alkyl substituted imidazole with PF5: synthesis and structural investigation. <i>Russian Chemical Bulletin</i> , 2014, 63, 2668-2674.	1.5	5
31	Inhibition Mechanism of Indoleamine 2, 3-Dioxygenase 1 (IDO1) by Amidoxime Derivatives and Its Revelation in Drug Design: Comparative Molecular Dynamics Simulations. <i>Frontiers in Molecular Biosciences</i> , 2019, 6, 164.	3.5	5
32	Research progress of human AP endonuclease 1: structure, catalytic mechanism and inhibitors. <i>Current Protein and Peptide Science</i> , 2022, 23, .	1.4	5
33	Design of stable platinum(II) complexes exhibited various colors via auxiliary ligand and electron-donating/withdrawing groups: A theoretical investigation. <i>Organic Electronics</i> , 2019, 71, 251-257.	2.6	4
34	Specifically targeting Mtb cell-wall and TMM transporter: the development of MmpL3 inhibitors. <i>Current Protein and Peptide Science</i> , 2021, 22, 290-303.	1.4	4
35	Allosteric and transport modulation of human concentrative nucleoside transporter 3 at the atomic scale. <i>Physical Chemistry Chemical Physics</i> , 2021, 23, 25401-25413.	2.8	4
36	An effective HIV-1 integrase inhibitor screening platform: Rationality validation of drug screening, conformational mobility and molecular recognition analysis for PFV integrase complex with viral DNA. <i>Journal of Molecular Graphics and Modelling</i> , 2017, 78, 96-109.	2.4	3

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37	Non-synonymous substitution of evolutionarily conserved residue in Tau class glutathione transferases alters structural and catalytic features. <i>International Journal of Biological Macromolecules</i> , 2022, 197, 39-48.	7.5	3
38	A potential strategy used for controlling the phosphorescence quantum yield of cyclometalated (C [†] C*) platinum(II) NHC complexes: The theoretical insight. <i>Organic Electronics</i> , 2018, 57, 367-376.	2.6	2
39	Influence of restricted rotation of small-sized substituent on phosphorescence efficiency for Pt(II) complexes: A theoretical investigation. <i>Organic Electronics</i> , 2018, 61, 25-34.	2.6	2
40	Potential strategy used for controlling the phosphorescent properties in tetradentate Pt(II) complexes: Effect of azole ligand. <i>Applied Organometallic Chemistry</i> , 2019, 33, e5125.	3.5	2
41	An Overview of Computational Tools of Nucleic Acid Binding Site Prediction for Site-specific Proteins and Nucleases. <i>Protein and Peptide Letters</i> , 2020, 27, 370-384.	0.9	2
42	In vitro selection and identification of a cold-tolerant variant in pineapple (<i>Ananas comosus</i>). <i>Horticulture Environment and Biotechnology</i> , 2022, 63, 275.	2.1	2
43	A novel ligand swing-mediated active site coordination change of human apurinic/aprimidinic endonuclease 1: A potential cytotoxic mechanism of nickel ion in the base excision repair. <i>Chemical Physics</i> , 2022, 555, 111456.	1.9	2
44	Molecular basis for substrate recognition by the bacterial nucleoside transporter NupG. <i>Journal of Biological Chemistry</i> , 2021, 296, 100479.	3.4	1
45	A promising strategy for increasing phosphorescent quantum yield: The ligand 10- β -cyclic chelate of the tetradentate Pt(II) complex. <i>Applied Organometallic Chemistry</i> , 2022, 36, .	3.5	1
46	Immunotherapy: MAPK β -targeted Drug Delivered by a pH-sensitive MSNP Nanocarrier Synergizes with PD-1 Blockade in Melanoma without T-cell Suppression (<i>Adv. Funct. Mater.</i> 12/2019). <i>Advanced Functional Materials</i> , 2019, 29, 1970079.	14.9	0
47	Phosphine ligand-coated Cu nanoparticle-catalyzed selective semihydrogenation of alkynes: electronic or hindrance effects of the ligand?. <i>Physical Chemistry Chemical Physics</i> , 2020, 22, 16905-16913.	2.8	0
48	DFT insight into Hashmi phenol synthesis catalyzed by Au single-walled nanotubes: mechanism and charge effect. <i>Theoretical Chemistry Accounts</i> , 2021, 140, 1.	1.4	0