

Fangqiang Zhu

List of Publications by Year in descending order

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32
papers

1,739
citations

516710

16
h-index

454955

30
g-index

32
all docs

32
docs citations

32
times ranked

2112
citing authors

#	ARTICLE	IF	CITATIONS
1	Discovery of Small-Molecule Inhibitors of SARS-CoV-2 Proteins Using a Computational and Experimental Pipeline. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 678701.	3.5	22
2	Toward Convergence in Free Energy Calculations for Protein Conformational Changes: A Case Study on the Thin Gate of Mhp1 Transporter. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 6583-6596.	5.3	3
3	High-throughput virtual screening of small molecule inhibitors for SARS-CoV-2 protein targets with deep fusion models. , 2021, , .		7
4	Assessing the Perturbing Effects of Drugs on Lipid Bilayers Using Gramicidin Channel-Based <i>In Silico</i> and <i>In Vitro</i> Assays. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 11809-11818.	6.4	10
5	Binding Affinity Prediction by Pairwise Function Based on Neural Network. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 2766-2772.	5.4	39
6	Calculating Single-Channel Permeability and Conductance from Transition Paths. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 777-785.	5.4	10
7	Kinetic mechanism for water in vibrating carbon nanotubes. <i>Physical Review E</i> , 2018, 98, .	2.1	5
8	Vitamin E Has Reduced Affinity for a Polyunsaturated Phospholipid: An Umbrella Sampling Molecular Dynamics Simulations Study. <i>Journal of Physical Chemistry B</i> , 2018, 122, 8351-8358.	2.6	9
9	Structural Model of the Tubular Assembly of the Rous Sarcoma Virus Capsid Protein. <i>Journal of the American Chemical Society</i> , 2017, 139, 2006-2013.	13.7	10
10	Parameter Optimization for Interaction between C-Terminal Domains of HIV-1 Capsid Protein. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 1134-1141.	5.4	3
11	Calculating transition rates from durations of transition paths. <i>Journal of Chemical Physics</i> , 2017, 146, 124128.	3.0	5
12	Thermodynamics of Protein Folding Studied by Umbrella Sampling along a Reaction Coordinate of Native Contacts. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 2086-2097.	5.3	20
13	Finite Temperature String Method with Umbrella Sampling: Application on a Side Chain Flipping in Mhp1 Transporter. <i>Journal of Physical Chemistry B</i> , 2017, 121, 3376-3386.	2.6	14
14	Construction of a novel coarse grain model for simulations of HIV capsid assembly to capture the backbone structure and inter-domain motions in solution. <i>Data in Brief</i> , 2015, 5, 506-512.	1.0	9
15	Conformational Changes in Two Inter-Helical Loops of Mhp1 Membrane Transporter. <i>PLoS ONE</i> , 2015, 10, e0133388.	2.5	11
16	Mechanism of polymorphism and curvature of HIV capsid assemblies probed by 3D simulations with a novel coarse grain model. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2015, 1850, 2353-2367.	2.4	32
17	Interface dipole enhancement effect and enhanced Rayleigh scattering. <i>Nano Research</i> , 2015, 8, 303-319.	10.4	12
18	True-color real-time imaging and spectroscopy of carbon nanotubes on substrates using enhanced Rayleigh scattering. <i>Nano Research</i> , 2015, 8, 2721-2732.	10.4	34

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19	Monte Carlo Simulations of HIV Capsid Protein Homodimer. <i>Journal of Chemical Information and Modeling</i> , 2015, 55, 1361-1368.	5.4	11
20	Evaluating membrane affinity by integrating protein orientations. <i>Journal of Molecular Graphics and Modelling</i> , 2014, 54, 141-147.	2.4	0
21	How does Water Pass through a Sugar Transporter?. <i>Biophysical Journal</i> , 2014, 106, 1229-1230.	0.5	4
22	Collective Diffusion Model for Ion Conduction through Microscopic Channels. <i>Biophysical Journal</i> , 2013, 104, 368-376.	0.5	18
23	Conformational Dynamics of a Ligand-Free Adenylate Kinase. <i>PLoS ONE</i> , 2013, 8, e68023.	2.5	27
24	Drying Transition in the Hydrophobic Gate of the GLIC Channel Blocks Ion Conduction. <i>Biophysical Journal</i> , 2012, 103, 219-227.	0.5	94
25	Theory and Simulation of Ion Conduction in the Pentameric GLIC Channel. <i>Journal of Chemical Theory and Computation</i> , 2012, 8, 3759-3768.	5.3	62
26	Convergence and error estimation in free energy calculations using the weighted histogram analysis method. <i>Journal of Computational Chemistry</i> , 2012, 33, 453-465.	3.3	187
27	Water Transport through Nanotubes with Varying Interaction Strength between Tube Wall and Water. <i>Journal of Physical Chemistry Letters</i> , 2011, 2, 2978-2983.	4.6	90
28	Pore opening and closing of a pentameric ligand-gated ion channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 19814-19819.	7.1	145
29	Gating Transition of Pentameric Ligand-Gated Ion Channels. <i>Biophysical Journal</i> , 2009, 97, 2456-2463.	0.5	35
30	Collective Diffusion Model for Water Permeation through Microscopic Channels. <i>Physical Review Letters</i> , 2004, 93, 224501.	7.8	204
31	Theory and Simulation of Water Permeation in Aquaporin-1. <i>Biophysical Journal</i> , 2004, 86, 50-57.	0.5	332
32	Water and Proton Conduction through Carbon Nanotubes as Models for Biological Channels. <i>Biophysical Journal</i> , 2003, 85, 236-244.	0.5	275