

Chenyi Liao

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2747858/publications.pdf>

Version: 2024-02-01

35
papers

855
citations

567281

15
h-index

501196

28
g-index

36
all docs

36
docs citations

36
times ranked

1567
citing authors

#	ARTICLE	IF	CITATIONS
1	Stiffening Effect of Ceramide on Lipid Membranes Provides Non-Sacrificial Protection against Potent Chemical Damage. <i>Langmuir</i> , 2022, 38, 3522-3529.	3.5	1
2	A double bilayer to study the nonequilibrium environmental response of GIRK2 in complex states. <i>Physical Chemistry Chemical Physics</i> , 2021, 23, 15784-15795.	2.8	1
3	Molecular basis for the adaptive evolution of environment-sensing by H-NS proteins. <i>ELife</i> , 2021, 10, .	6.0	9
4	Phosphorylation of SNX27 by MAPK11/14 links cellular stress signaling pathways with endocytic recycling. <i>Journal of Cell Biology</i> , 2021, 220, .	5.2	30
5	Molecular Basis of Class B GPCR Selectivity for the Neuropeptides PACAP and VIP. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 644644.	3.5	15
6	Aggregation State of Synergistic Antimicrobial Peptides. <i>Journal of Physical Chemistry Letters</i> , 2020, 11, 9501-9506.	4.6	13
7	Higher Accuracy Achieved for Protein-Ligand Binding Pose Prediction by Elastic Network Model-Based Ensemble Docking. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 2939-2950.	5.4	13
8	PAC1 Receptors: Shapeshifters in Motion. <i>Journal of Molecular Neuroscience</i> , 2019, 68, 331-339.	2.3	18
9	Targeting the PAC1 Receptor for Neurological and Metabolic Disorders. <i>Current Topics in Medicinal Chemistry</i> , 2019, 19, 1399-1417.	2.1	43
10	Top-down Multiscale Approach To Simulate Peptide Self-Assembly from Monomers. <i>Journal of Chemical Theory and Computation</i> , 2019, 15, 1514-1522.	5.3	10
11	Assessment of Conformational State Transitions of Class B GPCRs Using Molecular Dynamics. <i>Methods in Molecular Biology</i> , 2019, 1947, 3-19.	0.9	3
12	H-NS uses an autoinhibitory conformational switch for environment-controlled gene silencing. <i>Nucleic Acids Research</i> , 2019, 47, 2666-2680.	14.5	45
13	Printing DNA Strand Patterns on Small Molecules with Control of Valency, Directionality, and Sequence. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 3042-3047.	13.8	14
14	DNA-imprinted polymer nanoparticles with monodispersity and prescribed DNA-strand patterns. <i>Nature Chemistry</i> , 2018, 10, 184-192.	13.6	80
15	Direct cysteine sulfenylation drives activation of the Src kinase. <i>Nature Communications</i> , 2018, 9, 4522.	12.8	87
16	Image-Matching Based Identification of Store Signage Using Web-Crawled Information. <i>IEEE Access</i> , 2018, 6, 45590-45605.	4.2	3
17	Capturing the multiscale dynamics of membrane protein complexes with all-atom, mixed-resolution, and coarse-grained models. <i>Physical Chemistry Chemical Physics</i> , 2017, 19, 9181-9188.	2.8	13
18	A New Mixed All-Atom/Coarse-Grained Model: Application to Melittin Aggregation in Aqueous Solution. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 3881-3897.	5.3	16

#	ARTICLE	IF	CITATIONS
19	Studying Peptide Aggregation using Mixed All-Atom/Coarse Grain Molecular Dynamics Simulations. <i>Biophysical Journal</i> , 2017, 112, 198a.	0.5	1
20	Conformational Transitions of the Pituitary Adenylate Cyclase-Activating Polypeptide Receptor, a Human Class B GPCR. <i>Scientific Reports</i> , 2017, 7, 5427.	3.3	19
21	Targeting the apoptotic Mcl-1-PUMA interface with a dual-acting compound. <i>Oncotarget</i> , 2017, 8, 54236-54242.	1.8	16
22	Event.Locky: System of Event-Data Extraction from Webpages based on Web Mining. <i>Journal of Information Processing</i> , 2017, 25, 321-330.	0.4	1
23	Conformational Heterogeneity of Bax Helix 9 Dimer for Apoptotic Pore Formation. <i>Scientific Reports</i> , 2016, 6, 29502.	3.3	18
24	<sc>BH</sc> 3â€¢inâ€¢groove dimerization initiates and helix 9 dimerization expands Bax pore assembly in membranes. <i>EMBO Journal</i> , 2016, 35, 208-236.	7.8	81
25	Controlled Self-Assembly inside C-Shaped Polyaromatic Strips. <i>Synlett</i> , 2016, 27, 2145-2149.	1.8	8
26	Melittin Aggregation in Aqueous Solutions: Insight from Molecular Dynamics Simulations. <i>Journal of Physical Chemistry B</i> , 2015, 119, 10390-10398.	2.6	38
27	Design and Implementation of Event Information Summarization System. , 2014, , .		1
28	Computer simulations of fibronectin adsorption on hydroxyapatite surfaces. <i>RSC Advances</i> , 2014, 4, 15759.	3.6	49
29	Replica-Exchange Molecular Dynamics Simulation of Basic Fibroblast Growth Factor Adsorption on Hydroxyapatite. <i>Journal of Physical Chemistry B</i> , 2014, 118, 5843-5852.	2.6	48
30	Molecular simulations of myoglobin adsorbed on rutile (110) and (001) surfaces. <i>Fluid Phase Equilibria</i> , 2014, 362, 349-354.	2.5	19
31	Computer Simulations of Fibronectin Adsorption on Graphene Modified Titanium Dioxide Surfaces. <i>Acta Chimica Sinica</i> , 2014, 72, 401.	1.4	2
32	Multiscale Simulations of Protein G B1 Adsorbed on Charged Self-Assembled Monolayers. <i>Langmuir</i> , 2013, 29, 11366-11374.	3.5	54
33	Effects of external electric fields on lysozyme adsorption by molecular dynamics simulations. <i>Biophysical Chemistry</i> , 2013, 179, 26-34.	2.8	58
34	Replica Exchange Molecular Dynamics Simulations on the Folding of Trpzip4 Î²-Hairpin. <i>Acta Chimica Sinica</i> , 2013, 71, 593.	1.4	3
35	Two-dimensional self-assembly of esters with different configurations at the liquidâ€“solid interface. <i>Applied Surface Science</i> , 2011, 257, 4559-4565.	6.1	10