Xiaorong Liu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6074438/publications.pdf Version: 2024-02-01



XIAORONG LIU

#	Article	IF	CITATIONS
1	Accelerating atomistic simulations of proteins using multiscale enhanced sampling with independent tempering. Journal of Computational Chemistry, 2021, 42, 358-364.	3.3	2
2	EGCC binds intrinsically disordered N-terminal domain of p53 and disrupts p53-MDM2 interaction. Nature Communications, 2021, 12, 986.	12.8	77
3	Cancer-Associated Mutations Perturb the Disordered Ensemble and Interactions of the Intrinsically Disordered p53 Transactivation Domain. Journal of Molecular Biology, 2021, 433, 167048.	4.2	14
4	Accelerating the Generalized Born with Molecular Volume and Solvent Accessible Surface Area Implicit Solvent Model Using Graphics Processing Units. Journal of Computational Chemistry, 2020, 41, 830-838.	3.3	9
5	Targeting Intrinsically Disordered Proteins through Dynamic Interactions. Biomolecules, 2020, 10, 743.	4.0	34
6	Power generation from ambient humidity using protein nanowires. Nature, 2020, 578, 550-554.	27.8	398
7	Bioinspired bio-voltage memristors. Nature Communications, 2020, 11, 1861.	12.8	144
8	Modulation of p53 Transactivation Domain Conformations by Ligand Binding and Cancer-Associated Mutations. Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing, 2020, 25, 195-206.	0.7	4
9	Residual Structures and Transient Long-Range Interactions of p53 Transactivation Domain: Assessment of Explicit Solvent Protein Force Fields. Journal of Chemical Theory and Computation, 2019, 15, 4708-4720.	5.3	32
10	Residual Structure Accelerates Binding of Intrinsically Disordered ACTR by Promoting Efficient Folding upon Encounter. Journal of Molecular Biology, 2019, 431, 422-432.	4.2	24
11	Expression and Characterization of <i>Manduca sexta</i> Stress Responsive Peptide-1; An Inducer of Antimicrobial Peptide Synthesis. Biochemistry and Molecular Biology, 2019, 4, 42.	0.4	1
12	Modulation of p53 Transactivation Domain Conformations by Ligand Binding and Cancer-Associated Mutations. , 2019, , .		4
13	Atomistic Peptide Folding Simulations Reveal Interplay of Entropy and Long-Range Interactions in Folding Cooperativity. Scientific Reports, 2018, 8, 13668.	3.3	7
14	Enhanced Sampling of Intrinsic Structural Heterogeneity of the BH3-Only Protein Binding Interface of Bcl-xL. Journal of Physical Chemistry B, 2017, 121, 9160-9168.	2.6	17
15	HyRes: a coarse-grained model for multi-scale enhanced sampling of disordered protein conformations. Physical Chemistry Chemical Physics, 2017, 19, 32421-32432.	2.8	19
16	Dynamics of the BH3-Only Protein Binding Interface of Bcl-xL. Biophysical Journal, 2015, 109, 1049-1057.	0.5	11
17	Translocation Thermodynamics of Linear and Cyclic Nonaarginine into Model DPPC Bilayer via Coarse-Grained Molecular Dynamics Simulation: Implications of Pore Formation and Nonadditivity. Journal of Physical Chemistry B, 2014, 118, 2670-2682.	2.6	31
18	Toxicity of nano zinc oxide to mitochondria. Toxicology Research, 2012, 1, 137.	2.1	70

XIAORONG LIU

#	Article	IF	CITATIONS
19	Conformation and Thermodynamic Properties of the Binding of Vitamin C to Human Serum Albumin. Journal of Solution Chemistry, 2012, 41, 351-366.	1.2	23
20	Mitochondria as target of Quantum dots toxicity. Journal of Hazardous Materials, 2011, 194, 440-444.	12.4	63
21	Biophysical Studies on the Interactions of a Classic Mitochondrial Uncoupler with Bovine Serum Albumin by Spectroscopic, Isothermal Titration Calorimetric and Molecular Modeling Methods. Journal of Fluorescence, 2011, 21, 475-485.	2.5	52
22	Spectroscopic and Microscopic Studies on the Mechanisms of Mitochondrial Toxicity Induced by Different Concentrations of Cadmium. Journal of Membrane Biology, 2011, 241, 39-49.	2.1	38
23	Mitochondrial Permeability Transition Induced by Different Concentrations of Zinc. Journal of Membrane Biology, 2011, 244, 105-112.	2.1	21
24	Microcalorimetric, spectroscopic and microscopic investigation on the toxic effects of CdTe quantum dots on <i>Halobacterium halobium</i> R1. Nanotechnology, 2010, 21, 475102.	2.6	20
25	A simple and sensitive method for l-cysteine detection based on the fluorescence intensity increment of quantum dots. Analytica Chimica Acta, 2009, 645, 73-78.	5.4	96
26	Direct observation of the binding process between protein and quantum dots byin situsurface plasmon resonance measurements. Nanotechnology, 2009, 20, 325101.	2.6	23