

Silvina Matysiak

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

1,034
citations

516215

16
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414034

32
g-index

41
all docs

41
docs citations

41
times ranked

1263
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamics of Polymer Translocation through Nanopores: Theory Meets Experiment. <i>Physical Review Letters</i> , 2006, 96, 118103.	2.9	119
2	Balancing energy and entropy: A minimalist model for the characterization of protein folding landscapes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 10141-10146.	3.3	96
3	Mesoporous Encapsulated Chiral Nanogold for Use in Enantioselective Reactions. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 16791-16795.	7.2	91
4	Adaptive resolution simulation of liquid water. <i>Journal of Physics Condensed Matter</i> , 2007, 19, 292201.	0.7	85
5	Effect of pH on chitosan hydrogel polymer network structure. <i>Chemical Communications</i> , 2017, 53, 7373-7376.	2.2	69
6	Optimal Combination of Theory and Experiment for the Characterization of the Protein Folding Landscape of S6: How Far Can a Minimalist Model Go?. <i>Journal of Molecular Biology</i> , 2004, 343, 235-248.	2.0	64
7	Minimalist Protein Model as a Diagnostic Tool for Misfolding and Aggregation. <i>Journal of Molecular Biology</i> , 2006, 363, 297-308.	2.0	56
8	Acetylation within the First 17 Residues of Huntingtin Exon 1 Alters Aggregation and Lipid Binding. <i>Biophysical Journal</i> , 2016, 111, 349-362.	0.2	55
9	Role of Hydrophobic Hydration in Protein Stability: A 3D Water-Explicit Protein Model Exhibiting Cold and Heat Denaturation. <i>Journal of Physical Chemistry B</i> , 2012, 116, 8095-8104.	1.2	52
10	Looking at the Disordered Proteins through the Computational Microscope. <i>ACS Central Science</i> , 2018, 4, 534-542.	5.3	46
11	Dissecting the Energetics of Hydrophobic Hydration of Polypeptides. <i>Journal of Physical Chemistry B</i> , 2011, 115, 14859-14865.	1.2	34
12	Direct Characterization of Hydrophobic Hydration during Cold and Pressure Denaturation. <i>Journal of Physical Chemistry B</i> , 2012, 116, 5342-5348.	1.2	25
13	The Effects of Flanking Sequences in the Interaction of Polyglutamine Peptides with a Membrane Bilayer. <i>Journal of Physical Chemistry B</i> , 2014, 118, 6368-6379.	1.2	25
14	Multiscale Simulation of Liquid Water Using a Four-to-One Mapping for Coarse-Graining. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 5168-5175.	2.3	19
15	Effect of lipid head group interactions on membrane properties and membrane-induced cationic β -hairpin folding. <i>Physical Chemistry Chemical Physics</i> , 2016, 18, 17836-17850.	1.3	19
16	Mapping folding energy landscapes with theory and experiment. <i>Archives of Biochemistry and Biophysics</i> , 2008, 469, 29-33.	1.4	17
17	Dual mechanism of ionic liquid-induced protein unfolding. <i>Physical Chemistry Chemical Physics</i> , 2020, 22, 19779-19786.	1.3	17
18	Interplay between the hydrophobic effect and dipole interactions in peptide aggregation at interfaces. <i>Physical Chemistry Chemical Physics</i> , 2016, 18, 2449-2458.	1.3	16

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19	Pathways of amyloid-beta absorption and aggregation in a membranous environment. <i>Physical Chemistry Chemical Physics</i> , 2019, 21, 8559-8568.	1.3	16
20	Mesoporous Encapsulated Chiral Nanogold for Use in Enantioselective Reactions. <i>Angewandte Chemie</i> , 2018, 130, 17033-17037.	1.6	14
21	Computational insights into lipid assisted peptide misfolding and aggregation in neurodegeneration. <i>Physical Chemistry Chemical Physics</i> , 2019, 21, 22679-22694.	1.3	14
22	Long Distance Modulation of Disorder-to-Order Transitions in Protein Allostery. <i>Biochemistry</i> , 2017, 56, 4478-4488.	1.2	12
23	Influence of Monovalent Cation Size on Nanodomain Formation in Anionic-Zwitterionic Mixed Bilayers. <i>Journal of Physical Chemistry B</i> , 2017, 121, 787-799.	1.2	11
24	Effects of Sequence and Solvation on the Temperature-Pressure Conformational Landscape of Proteinlike Heteropolymers. <i>Physical Review Letters</i> , 2013, 111, 058103.	2.9	9
25	Interplay of Dynamical Properties between Ionic Liquids and Ionic Surfactants: Mechanism and Aggregation. <i>Journal of Physical Chemistry B</i> , 2015, 119, 9925-9932.	1.2	8
26	How Hydrophobic Hydration Destabilizes Surfactant Micelles at Low Temperature: A Coarse-Grained Simulation Study. <i>Langmuir</i> , 2018, 34, 12590-12599.	1.6	7
27	Load-Bearing Nanostructures in Composites of Chitosan with Anionic Surfactants: Implications for Programmable Mechanomaterials. <i>ACS Applied Nano Materials</i> , 2022, 5, 6463-6473.	2.4	7
28	Superrepression through Altered Corepressor-Activated Protein:Protein Interactions. <i>Biochemistry</i> , 2018, 57, 1119-1129.	1.2	6
29	Microscopic Picture of Calcium-Assisted Lipid Demixing and Membrane Remodeling Using Multiscale Simulations. <i>Journal of Physical Chemistry B</i> , 2020, 124, 7327-7335.	1.2	6
30	Tuning Allostery through Integration of Disorder to Order with a Residue Network. <i>Biochemistry</i> , 2020, 59, 790-801.	1.2	6
31	Effects of applied surface-tension on membrane-assisted A β aggregation. <i>Physical Chemistry Chemical Physics</i> , 2021, 23, 20627-20633.	1.3	5
32	Phase Behavior of a Lattice Hydrophobic Oligomer in Explicit Water. <i>Journal of Physical Chemistry B</i> , 2012, 116, 9540-9548.	1.2	3
33	Interplay between Conformational Heterogeneity and Hydration in the Folding Landscape of a Designed Three-Helix Bundle. <i>Journal of Physical Chemistry B</i> , 2017, 121, 2731-2738.	1.2	3
34	Recovery of enzyme structure and activity following rehydration from ionic liquid. <i>Physical Chemistry Chemical Physics</i> , 2022, , .	1.3	2
35	Thermal Stability of Hydrophobic Helical Oligomers: A Lattice Simulation Study in Explicit Water. <i>Journal of Physical Chemistry B</i> , 2012, 116, 9963-9970.	1.2	0
36	Length and Sequence Dependence in the Association of Htt Protein with Lipid Bilayers. <i>Biophysical Journal</i> , 2013, 104, 431a.	0.2	0

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37	Quantifying the Relationship between Monovalent Cation Size and Lipid Domain Formation in Anionic-Zwitterionic Mixed Lipid Bilayers. <i>Biophysical Journal</i> , 2017, 112, 520a-521a.	0.2	0
38	Protein:Protein Interactions Control Sensitivity of a Transcription Response to Input Signal. <i>Biophysical Journal</i> , 2018, 114, 69a.	0.2	0
39	Dual Mechanism of Ionic Liquid-Induced Protein Unfolding. <i>Biophysical Journal</i> , 2021, 120, 201a.	0.2	0
40	Assessment of physiological environment on neurodegenerative peptide aggregation. <i>Biophysical Journal</i> , 2022, 121, 352a.	0.2	0