

Wenhui Xi

List of Publications by Year in descending order

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papers

971
citations

623574

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docs citations

30
times ranked

1422
citing authors

#	ARTICLE	IF	CITATIONS
1	Protein Ensembles: How Does Nature Harness Thermodynamic Fluctuations for Life? The Diverse Functional Roles of Conformational Ensembles in the Cell. <i>Chemical Reviews</i> , 2016, 116, 6516-6551.	23.0	302
2	The molecular mechanism of fullerene-inhibited aggregation of Alzheimer's β -amyloid peptide fragment. <i>Nanoscale</i> , 2014, 6, 9752-9762.	2.8	135
3	Tunable assembly of amyloid-forming peptides into nanosheets as a retrovirus carrier. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 2996-3001.	3.3	123
4	Interactions of a Water-Soluble Fullerene Derivative with Amyloid- β Protofibrils: Dynamics, Binding Mechanism, and the Resulting Salt-Bridge Disruption. <i>Journal of Physical Chemistry B</i> , 2014, 118, 6733-6741.	1.2	50
5	Synergistic Inhibitory Effect of Peptide-Organic Coassemblies on Amyloid Aggregation. <i>ACS Nano</i> , 2016, 10, 4143-4153.	7.3	47
6	Atomic-Level Study of the Effects of O4 Molecules on the Structural Properties of Protofibrillar β -Trimer: β -Sheet Stabilization, Salt Bridge Protection, and Binding Mechanism. <i>Journal of Physical Chemistry B</i> , 2015, 119, 2786-2794.	1.2	40
7	Simulating Protein Fold Switching by Replica Exchange with Tunneling. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 5656-5666.	2.3	31
8	Fibril-Barrel Transitions in Cylindrin Amyloids. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 3936-3944.	2.3	28
9	Ring-like N-fold Models of β 42 fibrils. <i>Scientific Reports</i> , 2017, 7, 6588.	1.6	26
10	On the Conformational Dynamics of β -Amyloid Forming Peptides: A Computational Perspective. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020, 8, 532.	2.0	23
11	Out-of-Register β 42 Assemblies as Models for Neurotoxic Oligomers and Fibrils. <i>Journal of Chemical Theory and Computation</i> , 2018, 14, 1099-1110.	2.3	22
12	Stability of a Recently Found Triple- β -Stranded β 1-42 Fibril Motif. <i>Journal of Physical Chemistry B</i> , 2016, 120, 4548-4557.	1.2	21
13	Evaluation of residue-residue contact prediction methods: From retrospective to prospective. <i>PLoS Computational Biology</i> , 2021, 17, e1009027.	1.5	19
14	Template Induced Conformational Change of Amyloid- β Monomer. <i>Journal of Physical Chemistry B</i> , 2012, 116, 7398-7405.	1.2	17
15	The effect of retro-inverse D-amino acid β -peptides on β -fibril formation. <i>Journal of Chemical Physics</i> , 2019, 150, 095101.	1.2	14
16	Bioinformatics Screening of Potential Biomarkers from mRNA Expression Profiles to Discover Drug Targets and Agents for Cervical Cancer. <i>International Journal of Molecular Sciences</i> , 2022, 23, 3968.	1.8	13
17	Inhibitory effect of hydrophobic fullerenes on the β -sheet-rich oligomers of a hydrophilic GNNQQNY peptide revealed by atomistic simulations. <i>RSC Advances</i> , 2017, 7, 13947-13956.	1.7	12
18	Stability of the N-Terminal Helix and Its Role in Amyloid Formation of Serum Amyloid A. <i>ACS Omega</i> , 2018, 3, 16184-16190.	1.6	10

#	ARTICLE	IF	CITATIONS
19	Large fatty acid-derived A β 242 oligomers form ring-like assemblies. <i>Journal of Chemical Physics</i> , 2019, 150, 075101.	1.2	9
20	Identification of Autism spectrum disorder based on a novel feature selection method and Variational Autoencoder. <i>Computers in Biology and Medicine</i> , 2022, 148, 105854.	3.9	9
21	Protein Residue Contact Prediction Based on Deep Learning and Massive Statistical Features from Multi-Sequence Alignment. <i>Tsinghua Science and Technology</i> , 2022, 27, 843-854.	4.1	8
22	Conversion between parallel and antiparallel β -sheets in wild-type and Iowa mutant A β 40 fibrils. <i>Journal of Chemical Physics</i> , 2018, 148, 045103.	1.2	6
23	Stability of A β fibril fragments in the presence of fatty acids. <i>Protein Science</i> , 2019, 28, 1973-1981.	3.1	3
24	Inter-Residue Distance Prediction From Duet Deep Learning Models. <i>Frontiers in Genetics</i> , 2022, 13, .	1.1	3
25	STRUCTURAL INSIGHT INTO THE POLYMORPHISM OF NNQNTF PROTOFIBRIL: IMPORTANCE OF INTERFACIAL WATER, POLAR AND AROMATIC RESIDUES. <i>Journal of Theoretical and Computational Chemistry</i> , 2013, 12, 1341012.	1.8	0