

Srirupa Chakraborty

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

522
citations

1039880

9
h-index

1058333

14
g-index

28
all docs

28
docs citations

28
times ranked

1156
citing authors

#	ARTICLE	IF	CITATIONS
1	A neutralizing antibody target in early HIV-1 infection was recapitulated in rhesus macaques immunized with the transmitted/founder envelope sequence. <i>PLoS Pathogens</i> , 2022, 18, e1010488.	2.1	3
2	Development of Martini 2.2 parameters for N-glycans: a case study of the HIV-1 Env glycoprotein dynamics. <i>Glycobiology</i> , 2021, 31, 787-799.	1.3	7
3	The SARS-CoV-2 Spike variant D614G favors an open conformational state. <i>Science Advances</i> , 2021, 7, .	4.7	156
4	Exploring the Role of Glycans in the Interaction of SARS-CoV-2 RBD and Human Receptor ACE2. <i>Viruses</i> , 2021, 13, 927.	1.5	29
5	HIV-1 and SARS-CoV-2: Patterns in the evolution of two pandemic pathogens. <i>Cell Host and Microbe</i> , 2021, 29, 1093-1110.	5.1	73
6	Visualization of the HIV-1 Env glycan shield across scales. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 28014-28025.	3.3	57
7	Quantification of the Resilience and Vulnerability of HIV-1 Native Glycan Shield at Atomistic Detail. <i>IScience</i> , 2020, 23, 101836.	1.9	11
8	Graph-Directed Approach for Downselecting Toxins for Experimental Structure Determination. <i>Marine Drugs</i> , 2020, 18, 256.	2.2	4
9	Acetylcholine Receptor Gating: Click-Twist-Tilt-Rip-Pop. <i>Biophysical Journal</i> , 2017, 112, 552a.	0.2	0
10	A mechanism for acetylcholine receptor gating based on structure, coupling, phi, and flip. <i>Journal of General Physiology</i> , 2017, 149, 85-103.	0.9	47
11	Emergence of Alternative Structures in Amyloid Beta 1-42 Monomeric Landscape by N-terminal Hexapeptide Amyloid Inhibitors. <i>Scientific Reports</i> , 2017, 7, 9941.	1.6	23
12	Structural correlates of affinity in fetal versus adult endplate nicotinic receptors. <i>Nature Communications</i> , 2016, 7, 11352.	5.8	14
13	Simulations of Endplate AChRs: Agonist Site β -Sheet and M1 α -Helix. <i>Biophysical Journal</i> , 2016, 110, 603a-604a.	0.2	0
14	Between the Sheets: Inter-Subunit Backbone Interactions at AChR Neurotransmitter Binding Sites. <i>Biophysical Journal</i> , 2016, 110, 604a.	0.2	0
15	Molecular Simulations of Muscle AChR Agonist Binding Sites. <i>Biophysical Journal</i> , 2015, 108, 429a.	0.2	0
16	Decrypting the Structural, Dynamic and Energetic Basis of Kinesin Interacting with Tubulin Dimer in Three ATPase States by All-Atom Molecular Dynamics Simulation. <i>Biophysical Journal</i> , 2015, 108, 134a.	0.2	0
17	Function of the M1 α -helix in endplate receptor activation and desensitization. <i>Journal of Physiology</i> , 2015, 593, 2851-2866.	1.3	10
18	Decrypting the Structural, Dynamic, and Energetic Basis of a Monomeric Kinesin Interacting with a Tubulin Dimer in Three ATPase States by All-Atom Molecular Dynamics Simulation. <i>Biochemistry</i> , 2015, 54, 859-869.	1.2	14

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19	Functional differences between neurotransmitter binding sites of muscle acetylcholine receptors. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 17660-17665.	3.3	22
20	A Comparative Study of the Major Biochemical States of Kinesin-MT Complex using Computational Techniques and All-Atom Structural Models. Biophysical Journal, 2014, 106, 443a.	0.2	0
21	Quantification of the Resilience and Vulnerability of HIV-1 Native Glycan Shield at Atomistic Detail. SSRN Electronic Journal, 0, , .	0.4	4