

Soohyung Park

List of Publications by Year in descending order

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33
papers

1,149
citations

489802

18
h-index

466096

32
g-index

34
all docs

34
docs citations

34
times ranked

1828
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative Molecular Dynamics Simulation Studies of Realistic Eukaryotic, Prokaryotic, and Archaeal Membranes. <i>Journal of Chemical Information and Modeling</i> , 2022, 62, 1036-1051.	2.5	32
2	Evolutionary balance between foldability and functionality of a glucose transporter. <i>Nature Chemical Biology</i> , 2022, 18, 713-723.	3.9	13
3	CHARMM-GUI Polymer Builder for Modeling and Simulation of Synthetic Polymers. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 2431-2443.	2.3	58
4	CHARMM-GUI Membrane Builder for Lipid Nanoparticles with Ionizable Cationic Lipids and PEGylated Lipids. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 5192-5202.	2.5	25
5	Experimentally Guided Computational Methods Yield Highly Accurate Insights into Transmembrane Interactions within the T Cell Receptor Complex. <i>Journal of Physical Chemistry B</i> , 2020, 124, 10303-10310.	1.2	1
6	Quantitative Characterization of Protein-Lipid Interactions by Free Energy Simulation between Binary Bilayers. <i>Journal of Chemical Theory and Computation</i> , 2019, 15, 6491-6503.	2.3	7
7	U-shaped caveolin-1 conformations are tightly regulated by hydrogen bonds with lipids. <i>Journal of Computational Chemistry</i> , 2019, 40, 1570-1577.	1.5	8
8	Analysis of Lipid Order States and Domains in Lipid Bilayer Simulations. <i>Journal of Chemical Theory and Computation</i> , 2019, 15, 688-697.	2.3	12
9	Structural Conservation and Effects of Alterations in T Cell Receptor Transmembrane Interfaces. <i>Biophysical Journal</i> , 2018, 114, 1030-1035.	0.2	8
10	Quantitative Characterization of Cholesterol Partitioning between Binary Bilayers. <i>Journal of Chemical Theory and Computation</i> , 2018, 14, 2829-2833.	2.3	14
11	Transmembrane features governing Fc receptor CD16A assembly with CD16A signaling adaptor molecules. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E5645-E5654.	3.3	32
12	CHARMM-GUI 10 years for biomolecular modeling and simulation. <i>Journal of Computational Chemistry</i> , 2017, 38, 1114-1124.	1.5	224
13	A conserved $\beta\beta$ transmembrane interface forms the core of a compact T-cell receptor-CD3 structure within the membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E6649-E6658.	3.3	40
14	Bilayer Properties of Lipid A from Various Gram-Negative Bacteria. <i>Biophysical Journal</i> , 2016, 111, 1750-1760.	0.2	88
15	Influence of Ganglioside GM1 Concentration on Lipid Clustering and Membrane Properties and Curvature. <i>Biophysical Journal</i> , 2016, 111, 1987-1999.	0.2	41
16	How Tolerant are Membrane Simulations with Mismatch in Area per Lipid between Leaflets?. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 3466-3477.	2.3	64
17	CHARMM-GUI HMMM Builder for Membrane Simulations with the Highly Mobile Membrane-Mimetic Model. <i>Biophysical Journal</i> , 2015, 109, 2012-2022.	0.2	89
18	Insight into Early-Stage Unfolding of GPI-Anchored Human Prion Protein. <i>Biophysical Journal</i> , 2015, 109, 2090-2100.	0.2	18

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19	Transmembrane Complexes of DAP12 Crystallized in Lipid Membranes Provide Insights into Control of Oligomerization in Immunoreceptor Assembly. <i>Cell Reports</i> , 2015, 11, 1184-1192.	2.9	20
20	Lipid-Linked Oligosaccharides in Membranes Sample Conformations That Facilitate Binding to Oligosaccharyltransferase. <i>Biophysical Journal</i> , 2014, 107, 1885-1895.	0.2	21
21	Theory of Adaptive Optimization for Umbrella Sampling. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 2719-2728.	2.3	21
22	Two Dimensional Window Exchange Umbrella Sampling for Transmembrane Helix Assembly. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 13-17.	2.3	38
23	Transmembrane Helix Assembly by Window Exchange Umbrella Sampling. <i>Physical Review Letters</i> , 2012, 108, 108102.	2.9	61
24	Atomistic Simulations of Dilute Polyelectrolyte Solutions. <i>Journal of Physical Chemistry B</i> , 2012, 116, 4319-4327.	1.2	25
25	Multisite reversible geminate reaction. <i>Journal of Chemical Physics</i> , 2009, 130, 074507.	1.2	16
26	Distance-Dependent Proton Transfer along Water Wires Connecting Acid-Base Pairs. <i>Journal of Physical Chemistry A</i> , 2009, 113, 6599-6606.	1.1	77
27	Diffusion-Influenced Excited-State Reversible Geminate ABCD Reaction in the Presence of an External Field. <i>Chemistry - an Asian Journal</i> , 2008, 3, 1266-1276.	1.7	5
28	Concentration Profiles near an Activated Enzyme. <i>Journal of Physical Chemistry B</i> , 2008, 112, 12104-12114.	1.2	10
29	Diffusion-Influenced Reversible Geminate ABCD Reaction in the Presence of an External Field. <i>Journal of Physical Chemistry B</i> , 2008, 112, 6241-6249.	1.2	9
30	Theory and Simulation of Diffusion-Controlled Michaelis-Menten Kinetics for a Static Enzyme in Solution. <i>Journal of Physical Chemistry B</i> , 2008, 112, 5977-5987.	1.2	39
31	Diffusion-Influenced Reversible Trapping Problem in the Presence of an External Field. <i>Chemistry - an Asian Journal</i> , 2006, 1, 216-223.	1.7	7
32	Diffusion-influenced excited-state reversible transfer reactions, $A^* + B \rightleftharpoons C^* + D$, with two different lifetimes: Theories and simulations. <i>Journal of Chemical Physics</i> , 2005, 123, 034507.	1.2	15
33	Exact solution of the excited-state geminate $A^* + B \rightleftharpoons C^* + D$ reaction with two different lifetimes and quenching. <i>Journal of Chemical Physics</i> , 2004, 121, 868-876.	1.2	10