

Christopher T Lee

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

27
papers

484
citations

9
h-index

22
g-index

32
ext. papers

632
ext. citations

8.4
avg, IF

3.91
L-index

#	Paper	IF	Citations
27	Emerging Computational Methods for the Rational Discovery of Allosteric Drugs. <i>Chemical Reviews</i> , 2016 , 116, 6370-90	68.1	140
26	Simulation-Based Approaches for Determining Membrane Permeability of Small Compounds. <i>Journal of Chemical Information and Modeling</i> , 2016 , 56, 721-33	6.1	127
25	On the Application of Molecular-Dynamics Based Markov State Models to Functional Proteins. <i>Journal of Chemical Theory and Computation</i> , 2014 , 10, 2648-2657	6.4	57
24	Known structure, unknown function: An inquiry-based undergraduate biochemistry laboratory course. <i>Biochemistry and Molecular Biology Education</i> , 2015 , 43, 245-62	1.3	29
23	Two Relations to Estimate Membrane Permeability Using Milestoning. <i>Journal of Physical Chemistry B</i> , 2016 , 120, 8606-16	3.4	26
22	Quantitative Ranking of Ligand Binding Kinetics with a Multiscale Milestoning Simulation Approach. <i>Journal of Physical Chemistry Letters</i> , 2018 , 9, 4941-4948	6.4	24
21	3D mesh processing using GAMer 2 to enable reaction-diffusion simulations in realistic cellular geometries. <i>PLoS Computational Biology</i> , 2020 , 16, e1007756	5	19
20	Structural basis for ligand modulation of the CCR2 conformational landscape. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 8131-8136	11.5	16
19	Exascale Computing: A New Dawn for Computational Biology. <i>Computing in Science and Engineering</i> , 2018 , 20, 18-25	1.5	13
18	An Open-Source Mesh Generation Platform for Biophysical Modeling Using Realistic Cellular Geometries. <i>Biophysical Journal</i> , 2020 , 118, 1003-1008	2.9	7
17	The Implementation of the Colored Abstract Simplicial Complex and its Application to Mesh Generation. <i>ACM Transactions on Mathematical Software</i> , 2019 , 45,	2.3	6
16	Applications and Challenges of Machine Learning to Enable Realistic Cellular Simulations. <i>Frontiers in Physics</i> , 2020 , 7,	3.9	5
15	GAMer 2: A System for 3D Mesh Processing of Cellular Electron Micrographs		4
14	Independent Markov decomposition: Toward modeling kinetics of biomolecular complexes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	3
13	Independent Markov Decomposition: Towards modeling kinetics of biomolecular complexes		2
12	Value of models for membrane budding. <i>Current Opinion in Cell Biology</i> , 2021 , 71, 38-45	9	2
11	Morphological principles of neuronal mitochondria. <i>Journal of Comparative Neurology</i> , 2021 ,	3.4	1

10	Structural basis for ligand modulation of the CCR2 conformational landscape	1
9	Benchmarking ensemble docking methods as a scientific outreach project	1
8	An Open Source Mesh Generation Platform for Biophysical Modeling Using Realistic Cellular Geometries	1
7	Benchmarking ensemble docking methods in D3R Grand Challenge 4.. <i>Journal of Computer-Aided Molecular Design</i> , 2022 , 36, 87	4.2
6	3D mesh processing using GAMer 2 to enable reaction-diffusion simulations in realistic cellular geometries 2020 , 16, e1007756	
5	3D mesh processing using GAMer 2 to enable reaction-diffusion simulations in realistic cellular geometries 2020 , 16, e1007756	
4	3D mesh processing using GAMer 2 to enable reaction-diffusion simulations in realistic cellular geometries 2020 , 16, e1007756	
3	3D mesh processing using GAMer 2 to enable reaction-diffusion simulations in realistic cellular geometries 2020 , 16, e1007756	
2	3D mesh processing using GAMer 2 to enable reaction-diffusion simulations in realistic cellular geometries 2020 , 16, e1007756	
1	3D mesh processing using GAMer 2 to enable reaction-diffusion simulations in realistic cellular geometries 2020 , 16, e1007756	