

Julija Zavadlav

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

481
citations

623734

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839539

18
g-index

19
all docs

19
docs citations

19
times ranked

329
citing authors

#	ARTICLE	IF	CITATIONS
1	Adaptive resolution simulation of an atomistic protein in MARTINI water. <i>Journal of Chemical Physics</i> , 2014, 140, 054114.	3.0	74
2	Adaptive Resolution Simulation of MARTINI Solvents. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 2591-2598.	5.3	46
3	Adaptive Resolution Simulation of a DNA Molecule in Salt Solution. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 5035-5044.	5.3	46
4	Adaptive resolution simulation of polarizable supramolecular coarse-grained water models. <i>Journal of Chemical Physics</i> , 2015, 142, 244118.	3.0	39
5	Learning neural network potentials from experimental data via Differentiable Trajectory Reweighting. <i>Nature Communications</i> , 2021, 12, 6884.	12.8	36
6	Adaptive Resolution Simulation of Supramolecular Water: The Concurrent Making, Breaking, and Remaking of Water Bundles. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 4138-4145.	5.3	30
7	Order and interactions in DNA arrays: Multiscale molecular dynamics simulation. <i>Scientific Reports</i> , 2017, 7, 4775.	3.3	27
8	Adaptive resolution simulation of an atomistic DNA molecule in MARTINI salt solution. <i>European Physical Journal: Special Topics</i> , 2016, 225, 1595-1607.	2.6	25
9	Adaptive resolution simulations coupling atomistic water to dissipative particle dynamics. <i>Journal of Chemical Physics</i> , 2017, 147, 114110.	3.0	22
10	Open-Boundary Molecular Dynamics of a DNA Molecule in a Hybrid Explicit/Implicit Salt Solution. <i>Biophysical Journal</i> , 2018, 114, 2352-2362.	0.5	22
11	Adaptive resolution simulations of biomolecular systems. <i>European Biophysics Journal</i> , 2017, 46, 821-835.	2.2	20
12	Multiscale Simulation of Protein Hydration Using the SWINGER Dynamical Clustering Algorithm. <i>Journal of Chemical Theory and Computation</i> , 2018, 14, 1754-1761.	5.3	19
13	Bayesian selection for coarse-grained models of liquid water. <i>Scientific Reports</i> , 2019, 9, 99.	3.3	18
14	Accelerated Simulations of Molecular Systems through Learning of Effective Dynamics. <i>Journal of Chemical Theory and Computation</i> , 2022, 18, 538-549.	5.3	17
15	Molecular Dynamics Simulation of High Density DNA Arrays. <i>Computation</i> , 2018, 6, 3.	2.0	12
16	SWINGER: a clustering algorithm for concurrent coupling of atomistic and supramolecular liquids. <i>Interface Focus</i> , 2019, 9, 20180075.	3.0	11
17	Tuning the Dielectric Response of Water in Nanoconfinement through Surface Wettability. <i>ACS Nano</i> , 2021, 15, 20311-20318.	14.6	10
18	Back-mapping augmented adaptive resolution simulation. <i>Journal of Chemical Physics</i> , 2020, 153, 164118.	3.0	7

#	ARTICLE	IF	CITATIONS
19	Extending the Adress Multiscale Scheme for Protein and Bilayer Applications. Biophysical Journal, 2016, 110, 643a-644a.	0.5	0