

R Gregor Weir

List of Publications by Year in descending order

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papers

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1307594

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#	ARTICLE	IF	CITATIONS
1	Relative free-energy calculations for scaffold hopping-type transformations with an automated RE-EDS sampling procedure. <i>Journal of Computer-Aided Molecular Design</i> , 2022, 36, 117-130.	2.9	10
2	Glycanâ€‘protein interactions determine kinetics of <i>N</i>-glycan remodeling. <i>RSC Chemical Biology</i> , 2021, 2, 917-931.	4.1	16
3	Volume-scaled common nearest neighbor clustering algorithm with free-energy hierarchy. <i>Journal of Chemical Physics</i> , 2021, 154, 084106.	3.0	8
4	N-Glycosylation Enhances Conformational Flexibility of Protein Disulfide Isomerase Revealed by Microsecond Molecular Dynamics and Markov State Modeling. <i>Journal of Physical Chemistry B</i> , 2021, 125, 9467-9479.	2.6	16
5	Connecting dynamic reweighting Algorithms: Derivation of the dynamic reweighting family tree. <i>Journal of Chemical Physics</i> , 2020, 153, 234106.	3.0	5
6	Variational implicit-solvent predictions of the dryâ€‘wet transition pathways for ligandâ€‘receptor binding and unbinding kinetics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 14989-14994.	7.1	12
7	Affinity, kinetics, and pathways of anisotropic ligands binding to hydrophobic model pockets. <i>Journal of Chemical Physics</i> , 2018, 149, 094902.	3.0	1
8	Principles for Tuning Hydrophobic Ligandâ€‘Receptor Binding Kinetics. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 3012-3019.	5.3	13
9	Confined Water Determines Transport Properties of Guest Molecules in Narrow Pores. <i>ACS Nano</i> , 2016, 10, 7646-7656.	14.6	66
10	Solvent Fluctuations Induce Non-Markovian Kinetics in Hydrophobic Pocket-Ligand Binding. <i>Journal of Physical Chemistry B</i> , 2016, 120, 8127-8136.	2.6	8
11	Curvature Dependence of Hydrophobic Hydration Dynamics. <i>Physical Review Letters</i> , 2015, 114, 187802.	7.8	17