## Ursula Kahler

List of Publications by Year in descending order

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1039880 1281743 11 266 9 11 citations h-index g-index papers 11 11 11 351 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Solvation Thermodynamics in Different Solvents: Water–Chloroform Partition Coefficients from Grid Inhomogeneous Solvation Theory. Journal of Chemical Information and Modeling, 2020, 60, 3843-3853.	2.5	11
2	Polarizable and non-polarizable force fields: Protein folding, unfolding, and misfolding. Journal of Chemical Physics, 2020, 153, 185102.	1.2	26
3	Macrocycle Cell Permeability Measured by Solvation Free Energies in Polar and Apolar Environments. Journal of Chemical Information and Modeling, 2020, 60, 3508-3517.	2.5	15
4	Protein-Protein Binding as a Two-Step Mechanism: Preselection of Encounter Poses during the Binding of BPTI and Trypsin. Biophysical Journal, 2020, 119, 652-666.	0.2	22
5	Catalytic Site p <i>K</i> <sub>a</sub> Values of Aspartic, Cysteine, and Serine Proteases: Constant pH MD Simulations. Journal of Chemical Information and Modeling, 2020, 60, 3030-3042.	2.5	44
6	Sodium-induced population shift drives activation of thrombin. Scientific Reports, 2020, 10, 1086.	1.6	8
7	An unexpected switch in peptide binding mode: from simulation to substrate specificity. Journal of Biomolecular Structure and Dynamics, 2018, 36, 4072-4084.	2.0	7
8	Electrostatic recognition in substrate binding to serine proteases. Journal of Molecular Recognition, 2018, 31, e2727.	1.1	13
9	Characterizing the Diversity of the CDR-H3 Loop Conformational Ensembles in Relationship to Antibody Binding Properties. Frontiers in Immunology, 2018, 9, 3065.	2.2	73
10	Localization of Millisecond Dynamics: Dihedral Entropy from Accelerated MD. Journal of Chemical Theory and Computation, 2016, 12, 3449-3455.	2.3	23
11	Dynamics Govern Specificity of a Protein-Protein Interface: Substrate Recognition by Thrombin. PLoS ONE, 2015, 10, e0140713.	1.1	24