Erich R Kuechler

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Distinct Features of Stress Granule Proteins Predict Localization in Membraneless Organelles. Journal of Molecular Biology, 2020, 432, 2349-2368.	4.2	57
2	A Variational Linear-Scaling Framework to Build Practical, Efficient Next-Generation Orbital-Based Quantum Force Fields. Journal of Chemical Theory and Computation, 2013, 9, 1417-1427.	5.3	55
3	Protein–Protein Interactions Mediated by Intrinsically Disordered Protein Regions Are Enriched in Missense Mutations. Biomolecules, 2020, 10, 1097.	4.0	22
4	Charge-dependent many-body exchange and dispersion interactions in combined QM/MM simulations. Journal of Chemical Physics, 2015, 143, 234111.	3.0	20
5	Quantum mechanical study of solvent effects in a prototype S <i>N</i> 2 reaction in solution: Clâ' attack on CH3Cl. Journal of Chemical Physics, 2014, 140, 054109.	3.0	16
6	Multiscale Methods for Computational RNA Enzymology. Methods in Enzymology, 2015, 553, 335-374.	1.0	16
7	Proteomic analysis reveals the direct recruitment of intrinsically disordered regions to stress granules. Journal of Cell Science, 2020, 133, .	2.0	15
8	GraPES: The Granule Protein Enrichment Server for prediction of biological condensate constituents. Nucleic Acids Research, 2022, 50, W384-W391.	14.5	10
9	VR-SCOSMO: A smooth conductor-like screening model with charge-dependent radii for modeling chemical reactions. Journal of Chemical Physics, 2016, 144, 164115.	3.0	3
10	Protein feature analysis of heat shock induced ubiquitination sites reveals preferential modification site localization. Journal of Proteomics, 2021, 239, 104182.	2.4	3