Steve P Meisburger

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/755016/publications.pdf

Version: 2024-02-01

30 papers 1,444 citations

471509 17 h-index 25 g-index

32 all docs 32 docs citations

times ranked

32

 $\begin{array}{c} 1765 \\ \text{citing authors} \end{array}$

#	Article	IF	CITATIONS
1	<i>REGALS</i> : a general method to deconvolve X-ray scattering data from evolving mixtures. IUCrJ, 2021, 8, 225-237.	2.2	23
2	Correlated Motions in Structural Biology. Biochemistry, 2021, 60, 2331-2340.	2.5	18
3	Diffuse X-ray scattering from correlated motions in a protein crystal. Nature Communications, 2020, 11, 1271.	12.8	37
4	The phenylketonuria-associated substitution R68S converts phenylalanine hydroxylase to a constitutively active enzyme but reduces its stability. Journal of Biological Chemistry, 2019, 294, 4359-4367.	3.4	8
5	An endogenous dAMP ligand in <i>Bacillus subtilis</i> class Ib RNR promotes assembly of a noncanonical dimer for regulation by dATP. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E4594-E4603.	7.1	18
6	Visualizing single-stranded nucleic acids in solution. Nucleic Acids Research, 2017, 45, gkw1297.	14.5	25
7	The impact of base stacking on the conformations and electrostatics of single-stranded DNA. Nucleic Acids Research, 2017, 45, 3932-3943.	14.5	47
8	X-ray Scattering Studies of Protein Structural Dynamics. Chemical Reviews, 2017, 117, 7615-7672.	47.7	83
9	Asymmetric DNA Unwrapping Drives Sequential Dimer Release in Nucleosomes. Biophysical Journal, 2017, 112, 370a-371a.	0.5	O
10	Conformations of Single-Stranded Nucleic Acids in Solution. Biophysical Journal, 2017, 112, 473a-474a.	0.5	0
10	Conformations of Single-Stranded Nucleic Acids in Solution. Biophysical Journal, 2017, 112, 473a-474a. Correlated Motions from Crystallography beyond Diffraction. Accounts of Chemical Research, 2017, 50, 580-583.	0.5	0
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11	Correlated Motions from Crystallography beyond Diffraction. Accounts of Chemical Research, 2017, 50, 580-583. Asymmetric unwrapping of nucleosomal DNA propagates asymmetric opening and dissociation of the histone core. Proceedings of the National Academy of Sciences of the United States of America, 2017,	15.6	11
11 12	Correlated Motions from Crystallography beyond Diffraction. Accounts of Chemical Research, 2017, 50, 580-583. Asymmetric unwrapping of nucleosomal DNA propagates asymmetric opening and dissociation of the histone core. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 334-339. Full-length model of the human galectin-4 and insights into dynamics of inter-domain communication.	15.6 7.1	11 89
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11 12 13	Correlated Motions from Crystallography beyond Diffraction. Accounts of Chemical Research, 2017, 50, 580-583. Asymmetric unwrapping of nucleosomal DNA propagates asymmetric opening and dissociation of the histone core. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 334-339. Full-length model of the human galectin-4 and insights into dynamics of inter-domain communication. Scientific Reports, 2016, 6, 33633. Domain Movements upon Activation of Phenylalanine Hydroxylase Characterized by Crystallography and Chromatography-Coupled Small-Angle X-ray Scattering. Journal of the American Chemical Society, 2016, 138, 6506-6516. A microfabricated fixed path length silicon sample holder improves background subtraction for	15.6 7.1 3.3 13.7	11 89 15 100
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19	Revealing transient structures of nucleosomes as DNA unwinds. Nucleic Acids Research, 2014, 42, 8767-8776.	14.5	73
20	Breaking the Radiation Damage Limit with Cryo-SAXS. Biophysical Journal, 2013, 104, 227-236.	0.5	53
21	Introducing Cryo-SAXS for Measuring Low Resolution Macromolecular Structure without Radiation Damage. Biophysical Journal, 2013, 104, 502a.	0.5	0
22	Polyelectrolyte properties of single stranded DNA measured using SAXS and singleâ€molecule FRET: Beyond the wormlike chain model. Biopolymers, 2013, 99, 1032-1045.	2.4	34
23	lonic strength-dependent persistence lengths of single-stranded RNA and DNA. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 799-804.	7.1	322
24	Small-Angle X-ray Scattering and Single-Molecule FRET Spectroscopy Produce Highly Divergent Views of the Low-Denaturant Unfolded State. Journal of Molecular Biology, 2012, 418, 226-236.	4.2	92
25	RNA and Its Ionic Cloud: Solution Scattering Experiments and Atomically Detailed Simulations. Biophysical Journal, 2012, 102, 819-828.	0.5	89
26	Effects of a Protecting Osmolyte on the Ion Atmosphere Surrounding DNA Duplexes. Biochemistry, 2011, 50, 8540-8547.	2.5	16
27	Double-Stranded RNA Resists Condensation. Physical Review Letters, 2011, 106, 108101.	7.8	47
28	Counting Ions around DNA with Anomalous Small-Angle X-ray Scattering. Journal of the American Chemical Society, 2010, 132, 16334-16336.	13.7	83
29	The Role of Helix Topology and Counterion Distributions in RNA Interactions. Biophysical Journal, 2010, 98, 471a.	0.5	0
30	Both helix topology and counterion distribution contribute to the more effective charge screening in dsRNA compared with dsDNA. Nucleic Acids Research, 2009, 37, 3887-3896.	14.5	72