

Alexey Krushelnitsky

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

742
citations

516710

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610901

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docs citations

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times ranked

681
citing authors

#	ARTICLE	IF	CITATIONS
1	Trajectory-Based Approach for the Analysis of CODEX Solid-State Exchange Experiments in the Slow and Intermediate Motion Regime: Comparison of Experiment, Simulation, and Analytical Treatment. <i>Journal of Physical Chemistry C</i> , 2021, 125, 6839-6850.	3.1	1
2	Relaxation-induced dipolar exchange with recoupling (RIDER) distortions in CODEX experiments. <i>Magnetic Resonance</i> , 2020, 1, 247-259.	1.9	2
3	Microsecond motions probed by near-rotary-resonance ^1H - ^{15}N MAS NMR experiments: the model case of protein overall-rocking in crystals. <i>Journal of Biomolecular NMR</i> , 2018, 71, 53-67.	2.8	34
4	Quantitative NMR study of heat-induced aggregation of eye-lens crystallin proteins under crowding conditions. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2018, 1866, 1055-1061.	2.3	3
5	Chapter 6. CODEX-based Methods for Studying Slow Dynamics. <i>New Developments in NMR</i> , 2018, , 161-192.	0.1	2
6	Coupling and Decoupling of Rotational and Translational Diffusion of Proteins under Crowding Conditions. <i>Journal of the American Chemical Society</i> , 2016, 138, 10365-10372.	13.7	86
7	NMR-Detected Brownian Dynamics of β -Crystallin over a Wide Range of Concentrations. <i>Biophysical Journal</i> , 2015, 108, 98-106.	0.5	21
8	The τ_{long} of the protein tumbling correlation function: observation by ^1H NMR relaxometry in a wide frequency and concentration range. <i>Journal of Biomolecular NMR</i> , 2015, 63, 403-415.	2.8	19
9	Slow motions in microcrystalline proteins as observed by MAS-dependent ^{15}N rotating-frame NMR relaxation. <i>Journal of Magnetic Resonance</i> , 2014, 248, 8-12.	2.1	41
10	Internal protein dynamics on ps to μs timescales as studied by multi-frequency ^{15}N solid-state NMR relaxation. <i>Journal of Biomolecular NMR</i> , 2013, 57, 219-235.	2.8	37
11	Solid-State NMR Approaches to Internal Dynamics of Proteins: From Picoseconds to Microseconds and Seconds. <i>Accounts of Chemical Research</i> , 2013, 46, 2028-2036.	15.6	72
12	The relation of the X-ray B-factor to protein dynamics: insights from recent dynamic solid-state NMR data. <i>Journal of Biomolecular Structure and Dynamics</i> , 2012, 30, 617-627.	3.5	16
13	The trehalose coating effect on the internal protein dynamics. <i>Physical Chemistry Chemical Physics</i> , 2012, 14, 2727.	2.8	23
14	The nuclear magnetic resonance relaxation data analysis in solids: General equations and the model-free approach. <i>Journal of Chemical Physics</i> , 2011, 135, 184104.	3.0	58
15	Microsecond Time Scale Mobility in a Solid Protein As Studied by the ^{15}N Site-Specific NMR Relaxation Rates. <i>Journal of the American Chemical Society</i> , 2010, 132, 11850-11853.	13.7	57
16	Direct Observation of Millisecond to Second Motions in Proteins by Dipolar CODEX NMR Spectroscopy. <i>Journal of the American Chemical Society</i> , 2009, 131, 12097-12099.	13.7	45
17	Intermolecular electrostatic interactions and Brownian tumbling in protein solutions. <i>Physical Chemistry Chemical Physics</i> , 2006, 8, 2117.	2.8	19
18	Comparison of the internal dynamics of globular proteins in the microcrystalline and rehydrated lyophilized states. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2006, 1764, 1639-1645.	2.3	10

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19	15N spin diffusion rate in solid-state NMR of totally enriched proteins: The magic angle spinning frequency effect. Journal of Magnetic Resonance, 2006, 182, 339-342.	2.1	33
20	Complex 1H, 13C-NMR relaxation and computer simulation study of side-chain dynamics in solid polylysine. Biopolymers, 2005, 78, 129-139.	2.4	5
21	Solid-state NMR and protein dynamics. Progress in Nuclear Magnetic Resonance Spectroscopy, 2005, 47, 1-25.	7.5	98
22	Hydration dependence of backbone and side chain polylysine dynamics: A 13C solid-state NMR and IR spectroscopy study. Biopolymers, 2004, 73, 1-15.	2.4	18
23	Expanding the Frequency Range of the Solid-State T1 ρ-Experiment for Heteronuclear Dipolar Relaxation. Solid State Nuclear Magnetic Resonance, 2002, 22, 423-438.	2.3	41