

Jan Lipfert

List of Publications by Year in descending order

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104
papers

5,541
citations

76294

40
h-index

88593

70
g-index

118
all docs

118
docs citations

118
times ranked

5720
citing authors

#	ARTICLE	IF	CITATIONS
1	Understanding Nucleic Acid-Ion Interactions. Annual Review of Biochemistry, 2014, 83, 813-841.	5.0	358
2	Small-Angle X-Ray Scattering from RNA, Proteins, and Protein Complexes. Annual Review of Biophysics and Biomolecular Structure, 2007, 36, 307-327.	18.3	270
3	Magnetic torque tweezers: measuring torsional stiffness in DNA and RecA-DNA filaments. Nature Methods, 2010, 7, 977-980.	9.0	263
4	Quantitative and Comprehensive Decomposition of the Ion Atmosphere around Nucleic Acids. Journal of the American Chemical Society, 2007, 129, 14981-14988.	6.6	255
5	Size and Shape of Detergent Micelles Determined by Small-Angle X-ray Scattering. Journal of Physical Chemistry B, 2007, 111, 12427-12438.	1.2	219
6	Cellular and Molecular Probing of Intact Human Organs. Cell, 2020, 180, 796-812.e19.	13.5	187
7	Dependence of Micelle Size and Shape on Detergent Alkyl Chain Length and Head Group. PLoS ONE, 2013, 8, e62488.	1.1	182
8	Quantitative Modeling and Optimization of Magnetic Tweezers. Biophysical Journal, 2009, 96, 5040-5049.	0.2	171
9	Double-stranded RNA under force and torque: Similarities to and striking differences from double-stranded DNA. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 15408-15413.	3.3	162
10	Evaluation of Ion Binding to DNA Duplexes Using a Size-Modified Poisson-Boltzmann Theory. Biophysical Journal, 2007, 93, 3202-3209.	0.2	134
11	Salt dependence of the radius of gyration and flexibility of single-stranded DNA in solution probed by small-angle x-ray scattering. Physical Review E, 2012, 86, 021901.	0.8	131
12	Unusual compactness of a polyproline type II structure. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 11698-11703.	3.3	124
13	Freely orbiting magnetic tweezers to directly monitor changes in the twist of nucleic acids. Nature Communications, 2011, 2, 439.	5.8	120
14	Structural Transitions and Thermodynamics of a Glycine-Dependent Riboswitch from Vibrio cholerae. Journal of Molecular Biology, 2007, 365, 1393-1406.	2.0	116
15	Long single α -helical tail domains bridge the gap between structure and function of myosin VI. Nature Structural and Molecular Biology, 2008, 15, 591-597.	3.6	109
16	Quantitative Guidelines for Force Calibration through Spectral Analysis of Magnetic Tweezers Data. Biophysical Journal, 2010, 99, 1292-1302.	0.2	97
17	Torsional sensing of small-molecule binding using magnetic tweezers. Nucleic Acids Research, 2010, 38, 7122-7132.	6.5	92
18	Studying genomic processes at the single-molecule level: introducing the tools and applications. Nature Reviews Genetics, 2013, 14, 9-22.	7.7	83

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19	A repulsive field: advances in the electrostatics of the ion atmosphere. <i>Current Opinion in Chemical Biology</i> , 2008, 12, 619-625.	2.8	80
20	Mixing and Matching Detergents for Membrane Protein NMR Structure Determination. <i>Journal of the American Chemical Society</i> , 2009, 131, 7320-7326.	6.6	79
21	Twist-Bend Coupling and the Torsional Response of Double-Stranded DNA. <i>Physical Review Letters</i> , 2017, 118, 217801.	2.9	79
22	The Complete VS Ribozyme in Solution Studied by Small-Angle X-Ray Scattering. <i>Structure</i> , 2008, 16, 1357-1367.	1.6	78
23	High Spatiotemporal-Resolution Magnetic Tweezers: Calibration and Applications for DNA Dynamics. <i>Biophysical Journal</i> , 2015, 109, 2113-2125.	0.2	77
24	Probing the mechanical properties, conformational changes, and interactions of nucleic acids with magnetic tweezers. <i>Journal of Structural Biology</i> , 2017, 197, 26-36.	1.3	77
25	Critical Assessment of Nucleic Acid Electrostatics via Experimental and Computational Investigation of an Unfolded State Ensemble. <i>Journal of the American Chemical Society</i> , 2008, 130, 12334-12341.	6.6	74
26	Electromagnetic Torque Tweezers: A Versatile Approach for Measurement of Single-Molecule Twist and Torque. <i>Nano Letters</i> , 2012, 12, 3634-3639.	4.5	70
27	Multiplexed protein force spectroscopy reveals equilibrium protein folding dynamics and the low-force response of von Willebrand factor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 18798-18807.	3.3	69
28	The Ligand-Free State of the TPP Riboswitch: A Partially Folded RNA Structure. <i>Journal of Molecular Biology</i> , 2010, 396, 153-165.	2.0	67
29	Torque Spectroscopy for the Study of Rotary Motion in Biological Systems. <i>Chemical Reviews</i> , 2015, 115, 1449-1474.	23.0	65
30	Cation-Anion Interactions within the Nucleic Acid Ion Atmosphere Revealed by Ion Counting. <i>Journal of the American Chemical Society</i> , 2015, 137, 14705-14715.	6.6	65
31	Protein Misfolding and Amyloid Formation for the Peptide GNNQQNY from Yeast Prion Protein Sup35: Simulation by Reaction Path Annealing. <i>Journal of Molecular Biology</i> , 2005, 349, 648-658.	2.0	64
32	A force calibration standard for magnetic tweezers. <i>Review of Scientific Instruments</i> , 2014, 85, 123114.	0.6	63
33	Biological Magnetometry: Torque on Superparamagnetic Beads in Magnetic Fields. <i>Physical Review Letters</i> , 2015, 114, 218301.	2.9	61
34	Probing the salt dependence of the torsional stiffness of DNA by multiplexed magnetic torque tweezers. <i>Nucleic Acids Research</i> , 2017, 45, 5920-5929.	6.5	60
35	Dissecting electrostatic screening, specific ion binding, and ligand binding in an energetic model for glycine riboswitch folding. <i>Rna</i> , 2010, 16, 708-719.	1.6	57
36	The temperature dependence of the helical twist of DNA. <i>Nucleic Acids Research</i> , 2018, 46, 7998-8009.	6.5	55

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37	Do conformational biases of simple helical junctions influence RNA folding stability and specificity?. <i>Rna</i> , 2009, 15, 2195-2205.	1.6	53
38	Force sensing by the vascular protein von Willebrand factor is tuned by a strong intermonomer interaction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 1208-1213.	3.3	51
39	Explaining the striking difference in twist-stretch coupling between DNA and RNA: A comparative molecular dynamics analysis. <i>Nucleic Acids Research</i> , 2015, 43, gkv1028.	6.5	50
40	Nucleosome Assembly Dynamics Involve Spontaneous Fluctuations in the Handedness of Tetrasomes. <i>Cell Reports</i> , 2015, 10, 216-225.	2.9	48
41	Expression, purification, and characterization of <i>Thermotoga maritima</i> membrane proteins for structure determination. <i>Protein Science</i> , 2006, 15, 961-975.	3.1	46
42	Absolute Intramolecular Distance Measurements with Angstrom-Resolution Using Anomalous Small-Angle X-ray Scattering. <i>Nano Letters</i> , 2016, 16, 5353-5357.	4.5	42
43	Time-Resolved Small-Angle X-ray Scattering Reveals Millisecond Transitions of a DNA Origami Switch. <i>Nano Letters</i> , 2018, 18, 2672-2676.	4.5	42
44	Free Energy Landscape and Dynamics of Supercoiled DNA by High-Speed Atomic Force Microscopy. <i>ACS Nano</i> , 2018, 12, 11907-11916.	7.3	39
45	A tethered ligand assay to probe SARS-CoV-2:ACE2 interactions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2114397119.	3.3	38
46	Sample holder for small-angle x-ray scattering static and flow cell measurements. <i>Review of Scientific Instruments</i> , 2006, 77, 046108.	0.6	37
47	Low-resolution models for nucleic acids from small-angle X-ray scattering with applications to electrostatic modeling. <i>Journal of Applied Crystallography</i> , 2007, 40, s229-s234.	1.9	37
48	Blind Predictions of DNA and RNA Tweezers Experiments with Force and Torque. <i>PLoS Computational Biology</i> , 2014, 10, e1003756.	1.5	36
49	A method to track rotational motion for use in single-molecule biophysics. <i>Review of Scientific Instruments</i> , 2011, 82, 103707.	0.6	35
50	Single-Molecule Magnetic Tweezers Studies of Type IB Topoisomerases. <i>Methods in Molecular Biology</i> , 2009, 582, 71-89.	0.4	34
51	Twisting DNA by salt. <i>Nucleic Acids Research</i> , 2022, 50, 5726-5738.	6.5	34
52	Conformational Changes and Flexibility of DNA Devices Observed by Small-Angle X-ray Scattering. <i>Nano Letters</i> , 2016, 16, 4871-4879.	4.5	33
53	Molecular structure, DNA binding mode, photophysical properties and recommendations for use of SYBR Gold. <i>Nucleic Acids Research</i> , 2021, 49, 5143-5158.	6.5	31
54	Mis-translation of a Computationally Designed Protein Yields an Exceptionally Stable Homodimer: Implications for Protein Engineering and Evolution. <i>Journal of Molecular Biology</i> , 2006, 362, 1004-1024.	2.0	29

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55	Riboswitch Conformations Revealed by Small-Angle X-Ray Scattering. <i>Methods in Molecular Biology</i> , 2009, 540, 141-159.	0.4	26
56	Structural and torsional properties of the RAD51-dsDNA nucleoprotein filament. <i>Nucleic Acids Research</i> , 2013, 41, 7023-7030.	6.5	25
57	Structure, mechanics, and binding mode heterogeneity of LEDGF/p75â€“DNA nucleoprotein complexes revealed by scanning force microscopy. <i>Nanoscale</i> , 2014, 6, 4611-4619.	2.8	24
58	High-throughput AFM analysis reveals unwrapping pathways of H3 and CENP-A nucleosomes. <i>Nanoscale</i> , 2021, 13, 5435-5447.	2.8	24
59	Dynamics and energy landscape of DNA plectoneme nucleation. <i>Physical Review E</i> , 2018, 98, .	0.8	21
60	Tuning Micelle Dimensions and Properties with Binary Surfactant Mixtures. <i>Langmuir</i> , 2014, 30, 13353-13361.	1.6	20
61	Dynamics of the Buckling Transition in Double-Stranded DNA and RNA. <i>Biophysical Journal</i> , 2020, 118, 1690-1701.	0.2	20
62	pH-Dependent Interactions in Dimers Govern the Mechanics and Structure of von Willebrand Factor. <i>Biophysical Journal</i> , 2016, 111, 312-322.	0.2	18
63	Stretching and Heating Single DNA Molecules with Optically Trapped Goldâ€“Silica Janus Particles. <i>ACS Photonics</i> , 2017, 4, 2843-2851.	3.2	18
64	Quantitative evaluation of statistical errors in small-angle X-ray scattering measurements. <i>Journal of Applied Crystallography</i> , 2017, 50, 621-630.	1.9	18
65	Designed anchoring geometries determine lifetimes of biotinâ€“streptavidin bonds under constant load and enable ultra-stable coupling. <i>Nanoscale</i> , 2020, 12, 21131-21137.	2.8	18
66	Analysis of small-angle X-ray scattering data of proteinâ€“detergent complexes by singular value decomposition. <i>Journal of Applied Crystallography</i> , 2007, 40, s235-s239.	1.9	17
67	The free energy landscape of retroviral integration. <i>Nature Communications</i> , 2019, 10, 4738.	5.8	17
68	Compensatory Mechanisms in Temperature Dependence of DNA Double Helical Structure: Bending and Elongation. <i>Journal of Chemical Theory and Computation</i> , 2020, 16, 2857-2863.	2.3	16
69	A Mo-anode-based in-house source for small-angle X-ray scattering measurements of biological macromolecules. <i>Review of Scientific Instruments</i> , 2016, 87, 025103.	0.6	15
70	Applying torque to the Escherichia coli flagellar motor using magnetic tweezers. <i>Scientific Reports</i> , 2017, 7, 43285.	1.6	15
71	Temperatureâ€“Dependent Atomic Models of Detergent Micelles Refined against Smallâ€“Angle Xâ€“Ray Scattering Data. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 5635-5639.	7.2	15
72	Use of Small Angle X-ray Scattering (SAXS) to Characterize Conformational States of Functional RNAs. <i>Methods in Enzymology</i> , 2009, 469, 237-251.	0.4	14

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73	Radiation-Induced Damage in Serine Phosphate Insights into a Mechanism for Direct DNA Strand Breakage. <i>Journal of Physical Chemistry B</i> , 2004, 108, 8036-8042.	1.2	12
74	Structural Architecture of the Nucleosome Remodeler ISWI Determined from Cross-Linking, Mass Spectrometry, SAXS, and Modeling. <i>Structure</i> , 2018, 26, 282-294.e6.	1.6	11
75	Quantifying the Precision of Single-Molecule Torque and Twist Measurements Using Allan Variance. <i>Biophysical Journal</i> , 2018, 114, 1970-1979.	0.2	11
76	Gold nanocrystal labels provide a sequence-agnostic 3D structure map in SAXS reconstructions. <i>Science Advances</i> , 2018, 4, eaar4418.	4.7	11
77	Measuring Single-Molecule Twist and Torque in Multiplexed Magnetic Tweezers. <i>Methods in Molecular Biology</i> , 2018, 1814, 75-98.	0.4	11
78	The structural ensemble of a Holliday junction determined by X-ray scattering interference. <i>Nucleic Acids Research</i> , 2020, 48, 8090-8098.	6.5	10
79	Quantifying the influence of the ion cloud on SAXS profiles of charged proteins. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 26351-26361.	1.3	9
80	The Dissociation Rate of Acetylacetonate Ligands Governs the Size of Ferrimagnetic Zinc Ferrite Nanocubes. <i>ACS Applied Materials & Interfaces</i> , 2020, 12, 217-226.	4.0	9
81	Quantifying epigenetic modulation of nucleosome breathing by high-throughput AFM imaging. <i>Biophysical Journal</i> , 2022, 121, 841-851.	0.2	9
82	Magnetic Tweezers for the Measurement of Twist and Torque. <i>Journal of Visualized Experiments</i> , 2014, , .	0.2	8
83	Recording and Analyzing Nucleic Acid Distance Distributions with X-ray Scattering Interferometry (XSI). <i>Current Protocols in Nucleic Acid Chemistry</i> , 2018, 73, e54.	0.5	8
84	Ru(TAP)32+ uses multivalent binding to accelerate and constrain photo-adduct formation on DNA. <i>Chemical Communications</i> , 2019, 55, 8764-8767.	2.2	8
85	Deciphering the Gene Regulatory Landscape Encoded in DNA Biophysical Features. <i>IScience</i> , 2019, 21, 638-649.	1.9	7
86	A benchmark data set for the mechanical properties of double-stranded DNA and RNA under torsional constraint. <i>Data in Brief</i> , 2020, 30, 105404.	0.5	6
87	A High-throughput Pipeline to Determine DNA and Nucleosome Conformations by AFM Imaging. <i>Bio-protocol</i> , 2021, 11, e4180.	0.2	4
88	Optical Investigation of Individual Red Blood Cells for Determining Cell Count and Cellular Hemoglobin Concentration in a Microfluidic Channel. <i>Micromachines</i> , 2021, 12, 358.	1.4	4
89	Mixing and Matching Detergents for Membrane Protein NMR Structure Determination. <i>Biophysical Journal</i> , 2009, 96, 195a.	0.2	1
90	A Gentle Twist on DNA. <i>Physics Magazine</i> , 0, 14, .	0.1	1

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91	Scaling Behavior of Single Stranded DNA Measured by Small Angle X-ray Scattering. Biophysical Journal, 2009, 96, 347a.	0.2	0
92	Small-Molecule Binding to DNA under Tension and Twist Studied by Magnetic Tweezers. Biophysical Journal, 2009, 96, 216a.	0.2	0
93	Supercoiling Double-Stranded RNA. Biophysical Journal, 2010, 98, 468a.	0.2	0
94	Topoisomerase IB Activity Investigated By Single Molecule Magnetic Tweezers: Mechanisms of Cytotoxicity. Biophysical Journal, 2010, 98, 63a.	0.2	0
95	Quantitative Guidelines for Force Calibration Through Spectral Analysis of Magnetic Tweezers Data. Biophysical Journal, 2011, 100, 481a.	0.2	0
96	Characterization of Supercoiled Double-Stranded RNA, and Comparison to Double-Stranded DNA. Biophysical Journal, 2011, 100, 236a.	0.2	0
97	RNA Structure, Function, and (Thermo-) Dynamics: A SAXS and Single-Molecule Perspective. Biophysical Journal, 2011, 100, 1a-2a.	0.2	0
98	Introducing the Electromagnetic Torque Tweezers. Biophysical Journal, 2012, 102, 386a.	0.2	0
99	Modulating the Physical Properties of Micelles for Membrane Protein Investigations. Biophysical Journal, 2013, 104, 44a.	0.2	0
100	Quantifying the Resolution of Single-Molecule Torque Measurements by Allan Variance. Biophysical Journal, 2014, 106, 450a.	0.2	0
101	Nucleosome Dynamics Involve Spontaneous Fluctuations in the Handedness of Tetrasomes. Biophysical Journal, 2014, 106, 74a.	0.2	0
102	Magnetization Properties of Superparamagnetic Beads. Biophysical Journal, 2014, 106, 393a.	0.2	0
103	Nucleosome Assembly Dynamics Involve Spontaneous Fluctuations in the Handedness of Tetrasomes. Biophysical Journal, 2015, 108, 205a.	0.2	0
104	Temperature-Dependent Atomic Models of Detergent Micelles Refined against Small-Angle X-Ray Scattering Data. Angewandte Chemie, 2018, 130, 5737-5741.	1.6	0