

Sua Myong

List of Publications by Year in descending order

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Version: 2024-02-01

84
papers

7,056
citations

76326

40
h-index

71685

76
g-index

89
all docs

89
docs citations

89
times ranked

7966
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-molecule and ensemble methods to probe RNP nucleation and condensate properties. <i>Methods</i> , 2022, 197, 74-81.	3.8	11
2	Next generation single-molecule techniques: Imaging, labeling, and manipulation in vitro and in cellulo. <i>Molecular Cell</i> , 2022, 82, 304-314.	9.7	17
3	Single molecule probing of disordered RNA binding proteins. <i>STAR Protocols</i> , 2022, 3, 101131.	1.2	1
4	Protocol for generation and regeneration of PEG-passivated slides for single-molecule measurements. <i>STAR Protocols</i> , 2022, 3, 101152.	1.2	11
5	TDP-43 represses cryptic exon inclusion in the FTD-ALS gene UNC13A. <i>Nature</i> , 2022, 603, 124-130.	27.8	193
6	Poly(ADP-ribose) drives condensation of FUS via a transient interaction. <i>Molecular Cell</i> , 2022, 82, 969-985.e11.	9.7	41
7	Helicase mediated vectorial folding of telomere G-quadruplex. <i>Methods in Enzymology</i> , 2022, , 283-297.	1.0	0
8	Vectorial folding of telomere overhang promotes higher accessibility. <i>Nucleic Acids Research</i> , 2022, 50, 6271-6283.	14.5	4
9	FRET-based dynamic structural biology: Challenges, perspectives and an appeal for open-science practices. <i>ELife</i> , 2021, 10, .	6.0	152
10	Is transcriptional regulation just going through a phase?. <i>Molecular Cell</i> , 2021, 81, 1579-1585.	9.7	27
11	A Helicase Unwinds Hexanucleotide Repeat RNA G-Quadruplexes and Facilitates Repeat-Associated Non-AUG Translation. <i>Journal of the American Chemical Society</i> , 2021, 143, 7368-7379.	13.7	43
12	Regeneration of PEG slide for multiple rounds of single-molecule measurements. <i>Biophysical Journal</i> , 2021, 120, 1788-1799.	0.5	19
13	Probing steps in DNA transcription using single-molecule methods. <i>Journal of Biological Chemistry</i> , 2021, 297, 101086.	3.4	6
14	TRF2 promotes dynamic and stepwise looping of POT1 bound telomeric overhang. <i>Nucleic Acids Research</i> , 2021, 49, 12377-12393.	14.5	8
15	Loss of Dynamic RNA Interaction and Aberrant Phase Separation Induced by Two Distinct Types of ALS/FTD-Linked FUS Mutations. <i>Molecular Cell</i> , 2020, 77, 82-94.e4.	9.7	119
16	ALS/FTLD-Linked Mutations in FUS Glycine Residues Cause Accelerated Gelation and Reduced Interactions with Wild-Type FUS. <i>Molecular Cell</i> , 2020, 80, 666-681.e8.	9.7	62
17	Methods to Study Phase-Separated Condensates and the Underlying Molecular Interactions. <i>Trends in Biochemical Sciences</i> , 2020, 45, 1004-1005.	7.5	28
18	R-loop induced G-quadruplex in non-template promotes transcription by successive R-loop formation. <i>Nature Communications</i> , 2020, 11, 3392.	12.8	71

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19	Position-Dependent Effect of Guanine Base Damage and Mutations on Telomeric G-Quadruplex and Telomerase Extension. <i>Biochemistry</i> , 2020, 59, 2627-2639.	2.5	21
20	RNA Droplets. <i>Annual Review of Biophysics</i> , 2020, 49, 247-265.	10.0	102
21	Antigenic Variation in <i>Neisseria gonorrhoeae</i> Occurs Independently of RecQ-Mediated Unwinding of the <i>pilE</i> G Quadruplex. <i>Journal of Bacteriology</i> , 2020, 202, .	2.2	9
22	Just Took a DNA Test, Turns Out 100% Not That Phase. <i>Molecular Cell</i> , 2020, 78, 193-194.	9.7	10
23	The yeast Hrq1 helicase stimulates Pso2 translesion nuclease activity and thereby promotes DNA interstrand crosslink repair. <i>Journal of Biological Chemistry</i> , 2020, 295, 8945-8957.	3.4	12
24	<i>E. coli</i> Rep helicase and RecA recombinase unwind G4 DNA and are important for resistance to G4-stabilizing ligands. <i>Nucleic Acids Research</i> , 2020, 48, 6640-6653.	14.5	24
25	RNA promotes phase separation of glycolysis enzymes into yeast G bodies in hypoxia. <i>ELife</i> , 2020, 9, .	6.0	70
26	Editorial overview: Advances and future prospects of molecular imaging for studying and quantifying biological processes. <i>Current Opinion in Chemical Biology</i> , 2019, 51, A4-A5.	6.1	1
27	G-Quadruplex and Protein Binding by Single-Molecule FRET Microscopy. <i>Methods in Molecular Biology</i> , 2019, 2035, 309-322.	0.9	15
28	RNA G-quadruplex is resolved by repetitive and ATP-dependent mechanism of DHX36. <i>Nature Communications</i> , 2019, 10, 1855.	12.8	56
29	Evolving Methods in Defining the Role of RNA in RNP Assembly. <i>Biological and Medical Physics Series</i> , 2019, , 39-55.	0.4	0
30	FUS Regulates Activity of MicroRNA-Mediated Gene Silencing. <i>Molecular Cell</i> , 2018, 69, 787-801.e8.	9.7	76
31	mRNA structure determines specificity of a polyQ-driven phase separation. <i>Science</i> , 2018, 360, 922-927.	12.6	421
32	Ubiquilin 2 modulates ALS/FTD-linked FUS-RNA complex dynamics and stress granule formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E11485-E11494.	7.1	100
33	A guanine-flipping and sequestration mechanism for G-quadruplex unwinding by RecQ helicases. <i>Nature Communications</i> , 2018, 9, 4201.	12.8	46
34	Detection of Methylation on dsDNA at Single-Molecule Level using Solid-State Nanopores. <i>Biophysical Journal</i> , 2018, 114, 216a.	0.5	1
35	Single-Molecule and Ensemble Methods to Probe Initial Stages of RNP Granule Assembly. <i>Methods in Molecular Biology</i> , 2018, 1814, 325-338.	0.9	8
36	Single-Cell Imaging Approaches for Studying Small-RNA-Induced Gene Regulation. <i>Biophysical Journal</i> , 2018, 115, 203-208.	0.5	2

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37	Probing Dynamic Assembly and Disassembly of Rad51 Tuned by Srs2 Using smFRET. <i>Methods in Enzymology</i> , 2018, 600, 321-345.	1.0	1
38	Structural basis of G-quadruplex unfolding by the DEAH/RHA helicase DHX36. <i>Nature</i> , 2018, 558, 465-469.	27.8	224
39	RNA Scanning of a Molecular Machine with a Built-in Ruler. <i>Journal of the American Chemical Society</i> , 2017, 139, 262-268.	13.7	10
40	The Chd1 Chromatin Remodeler Shifts Nucleosomal DNA Bidirectionally as a Monomer. <i>Molecular Cell</i> , 2017, 68, 76-88.e6.	9.7	50
41	Molecular mechanisms by which oxidative DNA damage promotes telomerase activity. <i>Nucleic Acids Research</i> , 2017, 45, 11752-11765.	14.5	58
42	RNA stem structure governs coupling of dicing and gene silencing in RNA interference. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E10349-E10358.	7.1	5
43	Single molecule probing by fluorescence and force detection. <i>Methods</i> , 2016, 105, 1-2.	3.8	0
44	Visualizing repetitive diffusion activity of double-strand RNA binding proteins by single molecule fluorescence assays. <i>Methods</i> , 2016, 105, 109-118.	3.8	6
45	Quantitative analysis and prediction of G-quadruplex forming sequences in double-stranded DNA. <i>Nucleic Acids Research</i> , 2016, 44, 4807-4817.	14.5	20
46	Single-molecule imaging reveals a common mechanism shared by G-quadruplex-resolving helicases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 8448-8453.	7.1	85
47	RNA Remodeling Activity of DEAD Box Proteins Tuned by Protein Concentration, RNA Length, and ATP. <i>Molecular Cell</i> , 2016, 63, 865-876.	9.7	51
48	Single-Molecule Imaging With One Color Fluorescence. <i>Methods in Enzymology</i> , 2016, 581, 33-51.	1.0	6
49	Oxidative guanine base damage regulates human telomerase activity. <i>Nature Structural and Molecular Biology</i> , 2016, 23, 1092-1100.	8.2	134
50	Systematic and Quantitative Analysis of G-Quadruplex DNA Folding. <i>Biophysical Journal</i> , 2016, 110, 565a.	0.5	0
51	Molecular Mechanism of Resolving Trinucleotide Repeat Hairpin by Helicases. <i>Structure</i> , 2015, 23, 1018-1027.	3.3	6
52	The disordered P granule protein LAF-1 drives phase separation into droplets with tunable viscosity and dynamics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 7189-7194.	7.1	986
53	Nanopore-Based Assay for Detection of Methylation in Double-Stranded DNA Fragments. <i>ACS Nano</i> , 2015, 9, 290-300.	14.6	73
54	Quantitative Analysis of RNA Interference by mRNA Counting at Single-Cell Level. <i>Biophysical Journal</i> , 2015, 108, 364a.	0.5	0

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55	The transcription factor GABP selectively binds and activates the mutant TERT promoter in cancer. <i>Science</i> , 2015, 348, 1036-1039.	12.6	451
56	Dynamic profiling of double-stranded RNA binding proteins. <i>Nucleic Acids Research</i> , 2015, 43, 7566-7576.	14.5	53
57	G-quadruplex formation in double strand DNA probed by NMM and CV fluorescence. <i>Nucleic Acids Research</i> , 2015, 43, 7961-7970.	14.5	74
58	Repetitive RNA unwinding by RNA helicase A facilitates RNA annealing. <i>Nucleic Acids Research</i> , 2014, 42, 8556-8564.	14.5	39
59	G-quadruplex conformation and dynamics are determined by loop length and sequence. <i>Nucleic Acids Research</i> , 2014, 42, 8106-8114.	14.5	142
60	Single Molecule Imaging of Proteins That Recognize and Repair DNA Damages. <i>IEEE Journal of Selected Topics in Quantum Electronics</i> , 2014, 20, 223-231.	2.9	0
61	Protein induced fluorescence enhancement (PIFE) for probing protein-nucleic acid interactions. <i>Chemical Society Reviews</i> , 2014, 43, 1221-1229.	38.1	195
62	Telomeric Overhang Length Determines Structural Dynamics and Accessibility to Telomerase and ALT-Associated Proteins. <i>Structure</i> , 2014, 22, 842-853.	3.3	62
63	Context-Dependent Remodeling of Rad51-DNA Complexes by Srs2 Is Mediated by a Specific Protein-Protein Interaction. <i>Journal of Molecular Biology</i> , 2014, 426, 1883-1897.	4.2	8
64	Single-molecule real-time detection of telomerase extension activity. <i>Scientific Reports</i> , 2014, 4, 6391.	3.3	37
65	ATP-independent diffusion of double-stranded RNA binding proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 151-156.	7.1	62
66	Srs2 prevents Rad51 filament formation by repetitive motion on DNA. <i>Nature Communications</i> , 2013, 4, 2281.	12.8	86
67	ATP Hydrolysis Enhances RNA Recognition and Antiviral Signal Transduction by the Innate Immune Sensor, Laboratory of Genetics and Physiology 2 (LGP2). <i>Journal of Biological Chemistry</i> , 2013, 288, 938-946.	3.4	74
68	POT1-TPP1 Regulates Telomeric Overhang Structural Dynamics. <i>Structure</i> , 2012, 20, 1872-1880.	3.3	115
69	Protein induced fluorescence enhancement as a single molecule assay with short distance sensitivity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 7414-7418.	7.1	224
70	Single-Molecule Nanopositioning: Structural Transitions of a Helicase-DNA Complex during ATP Hydrolysis. <i>Biophysical Journal</i> , 2011, 101, 976-984.	0.5	11
71	Rotations of the 2B Sub-domain of E. coli UvrD Helicase/Translocase Coupled to Nucleotide and DNA Binding. <i>Journal of Molecular Biology</i> , 2011, 411, 633-648.	4.2	57
72	Single Molecule Detection of One, Two and Multiplex Proteins Involved in DNA/RNA Transaction. <i>Cellular and Molecular Bioengineering</i> , 2011, 4, 125-137.	2.1	0

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73	Stepwise translocation of nucleic acid motors. <i>Current Opinion in Structural Biology</i> , 2010, 20, 121-127.	5.7	47
74	PcrA Helicase Dismantles RecA Filaments by Reeling in DNA in Uniform Steps. <i>Cell</i> , 2010, 142, 544-555.	28.9	156
75	Characterization of Single Molecule Protein Induced Fluorescence Enhancement (PIFE) as an Alternative to SmFRET. <i>Biophysical Journal</i> , 2010, 98, 590a.	0.5	0
76	Cytosolic Viral Sensor RIG-I Is a 5'-Triphosphate-Dependent Translocase on Double-Stranded RNA. <i>Science</i> , 2009, 323, 1070-1074.	12.6	325
77	Single Molecule Nanocontainers Made Porous Using a Bacterial Toxin. <i>Journal of the American Chemical Society</i> , 2009, 131, 14844-14849.	13.7	52
78	Roles of RIG-I N-terminal tandem CARD and splice variant in TRIM25-mediated antiviral signal transduction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 16743-16748.	7.1	219
79	Spring-Loaded Mechanism of DNA Unwinding by Hepatitis C Virus NS3 Helicase. <i>Science</i> , 2007, 317, 513-516.	12.6	281
80	Real-Time Observation of RecA Filament Dynamics with Single Monomer Resolution. <i>Cell</i> , 2006, 126, 515-527.	28.9	285
81	Bridging Conformational Dynamics and Function Using Single-Molecule Spectroscopy. <i>Structure</i> , 2006, 14, 633-643.	3.3	53
82	Unraveling helicase mechanisms one molecule at a time. <i>Nucleic Acids Research</i> , 2006, 34, 4225-4231.	14.5	28
83	Repetitive shuttling of a motor protein on DNA. <i>Nature</i> , 2005, 437, 1321-1325.	27.8	254
84	DNA-binding Orientation and Domain Conformation of the E.coli Rep Helicase Monomer Bound to a Partial Duplex Junction: Single-molecule Studies of Fluorescently Labeled Enzymes. <i>Journal of Molecular Biology</i> , 2004, 336, 395-408.	4.2	159