

Marcelo Nollmann

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8779107/publications.pdf>

Version: 2024-02-01

64
papers

4,529
citations

136950

32
h-index

118850

62
g-index

73
all docs

73
docs citations

73
times ranked

5989
citing authors

#	ARTICLE	IF	CITATIONS
1	RNA imaging in bacteria. <i>FEMS Microbiology Reviews</i> , 2021, 45, .	8.6	1
2	Biology across scales: from atomic processes to bacterial communities through the lens of the microscope. <i>FEMS Microbiology Reviews</i> , 2021, 45, .	8.6	1
3	Single-particle tracking photoactivated localization microscopy of membrane proteins in living plant tissues. <i>Nature Protocols</i> , 2021, 16, 1600-1628.	12.0	28
4	Cis-regulatory chromatin loops arise before TADs and gene activation, and are independent of cell fate during early <i>Drosophila</i> development. <i>Nature Genetics</i> , 2021, 53, 477-486.	21.4	111
5	The Impact of Space and Time on the Functional Output of the Genome. <i>Cold Spring Harbor Perspectives in Biology</i> , 2021, , a040378.	5.5	10
6	Perspectives on Chromosome Organization. <i>Journal of Molecular Biology</i> , 2020, 432, 635-637.	4.2	0
7	TADs or no TADs: Lessons From Single-cell Imaging of Chromosome Architecture. <i>Journal of Molecular Biology</i> , 2020, 432, 682-693.	4.2	9
8	ATP-Driven Separation of Liquid Phase Condensates in Bacteria. <i>Molecular Cell</i> , 2020, 79, 293-303.e4.	9.7	107
9	LifeTime and improving European healthcare through cell-based interceptive medicine. <i>Nature</i> , 2020, 587, 377-386.	27.8	108
10	Direct and simultaneous observation of transcription and chromosome architecture in single cells with Hi-M. <i>Nature Protocols</i> , 2020, 15, 840-876.	12.0	23
11	G1/S transcription factors assemble in increasing numbers of discrete clusters through G1 phase. <i>Journal of Cell Biology</i> , 2020, 219, .	5.2	8
12	Osmotic Stress Activates Two Reactive Oxygen Species Pathways with Distinct Effects on Protein Nanodomains and Diffusion. <i>Plant Physiology</i> , 2019, 179, 1581-1593.	4.8	62
13	Nanoscale organization of tetraspanins during HIV-1 budding by correlative dSTORM/AFM. <i>Nanoscale</i> , 2019, 11, 6036-6044.	5.6	35
14	Developmental control of plant Rho GTPase nano-organization by the lipid phosphatidylserine. <i>Science</i> , 2019, 364, 57-62.	12.6	182
15	Microscopy-Based Chromosome Conformation Capture Enables Simultaneous Visualization of Genome Organization and Transcription in Intact Organisms. <i>Molecular Cell</i> , 2019, 74, 212-222.e5.	9.7	183
16	TADs are 3D structural units of higher-order chromosome organization in <i>Drosophila</i> . <i>Science Advances</i> , 2018, 4, eaar8082.	10.3	237
17	Sequence-dependent catalytic regulation of the SpoIIIE motor activity ensures directionality of DNA translocation. <i>Scientific Reports</i> , 2018, 8, 5254.	3.3	3
18	Super Resolution Imaging of Start Transcription Factors in Yeast. <i>Biophysical Journal</i> , 2018, 114, 547a.	0.5	1

#	ARTICLE	IF	CITATIONS
19	Challenges and guidelines toward 4D nucleome data and model standards. <i>Nature Genetics</i> , 2018, 50, 1352-1358.	21.4	47
20	DNA Organization and Superresolved Segregation. <i>Methods in Molecular Biology</i> , 2018, 1805, 271-289.	0.9	1
21	Imaging of Bacterial Chromosome Organization by 3D Super-Resolution Microscopy. <i>Methods in Molecular Biology</i> , 2017, 1624, 253-268.	0.9	7
22	Highly efficient multicolor multifocus microscopy by optimal design of diffraction binary gratings. <i>Scientific Reports</i> , 2017, 7, 5284.	3.3	19
23	Angular reconstitution-based 3D reconstructions of nanomolecular structures from superresolution light-microscopy images. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 9273-9278.	7.1	36
24	Single-cell absolute contact probability detection reveals chromosomes are organized by multiple low-frequency yet specific interactions. <i>Nature Communications</i> , 2017, 8, 1753.	12.8	137
25	New insights into the function of a versatile class of membrane molecular motors from studies of <i>Myxococcus xanthus</i> surface (gliding) motility. <i>Microbial Cell</i> , 2017, 4, 98-100.	3.2	10
26	Astigmatic multifocus microscopy enables deep 3D super-resolved imaging. <i>Biomedical Optics Express</i> , 2016, 7, 2163.	2.9	22
27	Nanometer resolved single-molecule colocalization of nuclear factors by two-color super resolution microscopy imaging. <i>Methods</i> , 2016, 105, 44-55.	3.8	32
28	The mechanism of force transmission at bacterial focal adhesion complexes. <i>Nature</i> , 2016, 539, 530-535.	27.8	120
29	Bacterial partition complexes segregate within the volume of the nucleoid. <i>Nature Communications</i> , 2016, 7, 12107.	12.8	105
30	Multifocus microscopy with precise color multi-phase diffractive optics applied in functional neuronal imaging. <i>Biomedical Optics Express</i> , 2016, 7, 855.	2.9	47
31	Direct observation of the translocation mechanism of transcription termination factor Rho. <i>Nucleic Acids Research</i> , 2015, 43, 2367-2377.	14.5	27
32	Stochastic Self-Assembly of ParB Proteins Builds the Bacterial DNA Segregation Apparatus. <i>Cell Systems</i> , 2015, 1, 163-173.	6.2	118
33	The fluorescence properties and binding mechanism of SYTOX green, a bright, low photo-damage DNA intercalating agent. <i>European Biophysics Journal</i> , 2015, 44, 337-348.	2.2	50
34	A matter of scale: how emerging technologies are redefining our view of chromosome architecture. <i>Trends in Genetics</i> , 2015, 31, 454-464.	6.7	20
35	Roles of chromatin insulators in the formation of long-range contacts. <i>Nucleus</i> , 2015, 6, 118-122.	2.2	6
36	Condensin- and Replication-Mediated Bacterial Chromosome Folding and Origin Condensation Revealed by Hi-C and Super-resolution Imaging. <i>Molecular Cell</i> , 2015, 59, 588-602.	9.7	245

#	ARTICLE	IF	CITATIONS
37	Constructing a Magnetic Tweezers to Monitor RNA Translocation at the Single-Molecule Level. <i>Methods in Molecular Biology</i> , 2015, 1259, 257-273.	0.9	4
38	Chromatin Insulator Factors Involved in Long-Range DNA Interactions and Their Role in the Folding of the <i>Drosophila</i> Genome. <i>PLoS Genetics</i> , 2014, 10, e1004544.	3.5	101
39	Structure and DNA-binding properties of the <i>Bacillus subtilis</i> SpoIIIE DNA translocase revealed by single-molecule and electron microscopies. <i>Nucleic Acids Research</i> , 2014, 42, 2624-2636.	14.5	22
40	Chromatin Immunoprecipitation Indirect Peaks Highlight Long-Range Interactions of Insulator Proteins and Pol II Pausing. <i>Molecular Cell</i> , 2014, 53, 672-681.	9.7	102
41	Chromosome Organization: Original Condensins. <i>Current Biology</i> , 2014, 24, R111-R113.	3.9	4
42	Recruitment, Assembly, and Molecular Architecture of the SpoIIIE DNA Pump Revealed by Superresolution Microscopy. <i>PLoS Biology</i> , 2013, 11, e1001557.	5.6	71
43	SpoIIIE mechanism of directional translocation involves target search coupled to sequence-independent motor stimulation. <i>EMBO Reports</i> , 2013, 14, 473-479.	4.5	25
44	Super-Resolution Imaging of Bacteria in a Microfluidics Device. <i>PLoS ONE</i> , 2013, 8, e76268.	2.5	35
45	Single-molecule super-resolution imaging in bacteria. <i>Current Opinion in Microbiology</i> , 2012, 15, 758-763.	5.1	26
46	Sequence-directed DNA export guides chromosome translocation during sporulation in <i>Bacillus subtilis</i> . <i>Nature Structural and Molecular Biology</i> , 2008, 15, 485-493.	8.2	91
47	Dynamics of Neutrophil Migration in Lymph Nodes during Infection. <i>Immunity</i> , 2008, 29, 487-496.	14.3	366
48	SpoIIIE strips proteins off the DNA during chromosome translocation. <i>Genes and Development</i> , 2008, 22, 1786-1795.	5.9	63
49	Thirty years of <i>Escherichia coli</i> DNA gyrase: From in vivo function to single-molecule mechanism. <i>Biochimie</i> , 2007, 89, 490-499.	2.6	103
50	Multiple modes of <i>Escherichia coli</i> DNA gyrase activity revealed by force and torque. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 264-271.	8.2	101
51	Giant proteins that move DNA: bullies of the genomic playground. <i>Nature Reviews Molecular Cell Biology</i> , 2006, 7, 580-588.	37.0	44
52	Identification of the FtsK sequence-recognition domain. <i>Nature Structural and Molecular Biology</i> , 2006, 13, 1023-1025.	8.2	52
53	Mechanochemical analysis of DNA gyrase using rotor bead tracking. <i>Nature</i> , 2006, 439, 100-104.	27.8	172
54	DNA overwinds when stretched. <i>Nature</i> , 2006, 442, 836-839.	27.8	358

#	ARTICLE	IF	CITATIONS
55	SOMO (Solution MOdeler). Structure, 2005, 13, 723-734.	3.3	101
56	A global multi-technique approach to study low-resolution solution structures. Journal of Applied Crystallography, 2005, 38, 874-887.	4.5	10
57	Behavior of Tn3 Resolvase in Solution and Its Interaction with res. Biophysical Journal, 2005, 89, 1920-1931.	0.5	12
58	The Solution Structure and Oligomerization Behavior of Two Bacterial Toxins: Pneumolysin and Perfringolysin O. Biophysical Journal, 2004, 87, 540-552.	0.5	48
59	Solution Structure of the Tn3 Resolvase-Crossover Site Synaptic Complex. Molecular Cell, 2004, 16, 127-137.	9.7	44
60	Low-Resolution Reconstruction of a Synthetic DNA Holliday Junction. Biophysical Journal, 2004, 86, 3060-3069.	0.5	18
61	The Role of Cholesterol in the Activity of Pneumolysin, a Bacterial Protein Toxin. Biophysical Journal, 2004, 86, 3141-3151.	0.5	51
62	Heat does not come in different colours: entropyâ€“enthalpy compensation, free energy windows, quantum confinement, pressure perturbation calorimetry, solvation and the multiple causes of heat capacity effects in biomolecular interactions. Biophysical Chemistry, 2001, 93, 215-230.	2.8	308
63	Microscopy-Based Chromosome Conformation Capture Enables Simultaneous Visualization of Genome Organization and Transcription in Intact Organisms. SSRN Electronic Journal, 0, , .	0.4	2
64	Qudi-HiM: an open-source acquisition software package for highly multiplexed sequential and combinatorial optical imaging. Open Research Europe, 0, 2, 46.	2.0	2