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List of Publications by Year in descending order

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687335 477281 46 995 13 29 g-index citations h-index papers 50 50 50 1716 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Epithelial Cells in 2D and 3D Cultures Exhibit Large Differences in Higher-order Genomic Interactions. Genomics, Proteomics and Bioinformatics, 2022, 20, 101-109.	6.9	4
2	SARS-CoV-2 infection and oxidative stress: Pathophysiological insight into thrombosis and therapeutic opportunities. Cytokine and Growth Factor Reviews, 2022, 63, 44-57.	7.2	41
3	Robust Acquisition of Spatial Transcriptional Programs in Tissues With Immunofluorescence-Guided Laser Capture Microdissection. Frontiers in Cell and Developmental Biology, 2022, 10, 853188.	3.7	3
4	Single-Molecule Micromanipulation and Super-Resolution Imaging Resolve Nanodomains Underlying Chromatin Folding in Mitotic Chromosomes. ACS Nano, 2022, 16, 8030-8039.	14.6	7
5	Threeâ€Dimensional Quantitative Imaging of Native Microbiota Distribution in the Gut. Angewandte Chemie - International Edition, 2021, 60, 3055-3061.	13.8	31
6	Threeâ€Dimensional Quantitative Imaging of Native Microbiota Distribution in the Gut. Angewandte Chemie, 2021, 133, 3092-3098.	2.0	1
7	Efficient and Fast Immuno-labeling of Clarified Tissues Using Low-Field Enhanced Diffusion. IEEE Transactions on Biomedical Engineering, 2021, 68, 1-1.	4.2	O
8	High-resolution single-cell 3D-models of chromatin ensembles during Drosophila embryogenesis. Nature Communications, 2021, 12, 205.	12.8	17
9	The role of vitamin D in reducing SARS-CoV-2 infection: An update. International Immunopharmacology, 2021, 97, 107686.	3.8	31
10	Monocytic THP-1 cells diverge significantly from their primary counterparts: a comparative examination of the chromosomal conformations and transcriptomes. Hereditas, 2021, 158, 43.	1.4	8
11	Controlling Water Flow through a Synthetic Nanopore with Permeable Cations. ACS Central Science, 2021, 7, 2092-2098.	11.3	8
12	Evidence for heightened genetic instability in precancerous spasmolytic polypeptide expressing gastric glands. Journal of Medical Genetics, 2020, 57, 385-388.	3.2	6
13	Q-Nuc: a bioinformatics pipeline for the quantitative analysis of nucleosomal profiles. Interdisciplinary Sciences, Computational Life Sciences, 2020, 12, 69-81.	3.6	6
14	Significant improvement in data quality with simplified SCRB-seq. Acta Biochimica Et Biophysica Sinica, 2020, 52, 457-459.	2.0	5
15	Ultrasensitive liposome-based assay for the quantification of fundamental ion channel properties. Analytica Chimica Acta, 2020, 1112, 8-15.	5.4	7
16	Massive reorganization of the genome during primary monocyte differentiation into macrophage. Acta Biochimica Et Biophysica Sinica, 2020, 52, 546-553.	2.0	4
17	Unconventional Atomic Structure of Graphene Sheets on Solid Substrates. Small, 2019, 15, 1902637.	10.0	2
18	Atomic force microscopy-based single-molecule force spectroscopy detects DNA base mismatches. Nanoscale, 2019, 11, 17206-17210.	5.6	13

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19	Interplay between the bacterial protein deacetylase CobB and the second messenger câ€di― <scp>GMP</scp> . EMBO Journal, 2019, 38, e100948.	7.8	28
20	Discovery and genetic characterization of intestinal metaplasia in the Helicobacter felis-infected mouse model of gastric cancer. Acta Biochimica Et Biophysica Sinica, 2019, 51, 219-222.	2.0	1
21	Identification of Serine 119 as an Effective Inhibitor Binding Site of M. tuberculosis Ubiquitin-like Protein Ligase PafA Using Purified Proteins and M. smegmatis. EBioMedicine, 2018, 30, 225-236.	6.1	9
22	Sub-kb Hi-C in D. melanogaster reveals conserved characteristics of TADs between insect and mammalian cells. Nature Communications, 2018, 9, 188.	12.8	126
23	Super-resolution Imaging of Individual Human Subchromosomal Regions <i>in Situ</i> Reveals Nanoscopic Building Blocks of Higher-Order Structure. ACS Nano, 2018, 12, 4909-4918.	14.6	41
24	abLIM1 constructs non-erythroid cortical actin networks to prevent mechanical tension-induced blebbing. Cell Discovery, 2018, 4, 42.	6.7	10
25	Helix $\hat{l}\pm -3$ inter-molecular salt bridges and conformational changes are essential for toxicity of Bacillus thuringiensis 3D-Cry toxin family. Scientific Reports, 2018, 8, 10331.	3.3	13
26	Cell Lysate Microarray for Mapping the Network of Genetic Regulators for Histone Marks. Molecular and Cellular Proteomics, 2018, 17, 1720-1736.	3.8	1
27	Glycan-independent binding and internalization of human IgM to FCMR, its cognate cellular receptor. Scientific Reports, 2017, 7, 42989.	3.3	20
28	The Ser/Thr Protein Kinase Protein-Protein Interaction Map of M. tuberculosis*. Molecular and Cellular Proteomics, 2017, 16, 1491-1506.	3.8	39
29	Improved clearing of lipid droplet-rich tissues for three-dimensional structural elucidation. Acta Biochimica Et Biophysica Sinica, 2017, 49, 465-467.	2.0	9
30	Toward the development of magnetic tweezers for high-throughput measurement of protein–protein interactions. Acta Biochimica Et Biophysica Sinica, 2017, 49, 468-470.	2.0	0
31	Ultra-deep sequencing of ribosome-associated poly-adenylated RNA in early <i>Drosophila</i> embryos reveals hundreds of conserved translated sORFs. DNA Research, 2016, 23, 571-580.	3.4	14
32	Complex clonal mosaicism within microdissected intestinal metaplastic glands without concurrent gastric cancer. Journal of Medical Genetics, 2016, 53, 643-646.	3.2	6
33	Nanoscopic characterization of the water vapor-salt interfacial layer reveals a unique biphasic adsorption process. Scientific Reports, 2016, 6, 31688.	3.3	3
34	Development of a low-noise cryogenic atomic force microscope for high resolution imaging of large biological complexes. Acta Biochimica Et Biophysica Sinica, 2016, 48, 859-861.	2.0	1
35	The non-coding RNA composition of the mitotic chromosome by 5′-tag sequencing. Nucleic Acids Research, 2016, 44, 4934-4946.	14.5	16
36	Fast immuno-labeling by electrophoretically driven infiltration for intact tissue imaging. Scientific Reports, 2015, 5, 10640.	3.3	40

3

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37	Developing the IVIG biomimetic, Hexa-Fc, for drug and vaccine applications. Scientific Reports, 2015, 5, 9526.	3.3	33
38	Identification of the minimal binding region of a Plasmodium falciparum IgM binding PfEMP1 domain. Molecular and Biochemical Parasitology, 2015, 201, 76-82.	1.1	14
39	Systematic identification of arsenic-binding proteins reveals that hexokinase-2 is inhibited by arsenic. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 15084-15089.	7.1	126
40	Illuminated up close: near-field optical microscopy of cell surfaces. Nanomedicine: Nanotechnology, Biology, and Medicine, 2015, 11, 119-125.	3.3	4
41	YcgC represents a new protein deacetylase family in prokaryotes. ELife, 2015, 4, .	6.0	52
42	Single molecule compression reveals intra-protein forces drive cytotoxin pore formation. ELife, 2015, 4, e08421.	6.0	16
43	Re-evaluation of the widely applied force-frequency relation for frequency-modulation AFM under solution. Journal of Shanghai Jiaotong University (Science), 2014, 19, 612-616.	0.9	0
44	Direct resolution of the pitch of DNA on positively charged lipid bilayers by frequency-modulation AFM. Journal of Shanghai Jiaotong University (Science), 2014, 19, 565-568.	0.9	1
45	Novel experimental strategy for high resolution AFM imaging of membrane-associated bacterial toxins. Journal of Shanghai Jiaotong University (Science), 2014, 19, 569-573.	0.9	1
46	Self-assembling subnanometer pores with unusual mass-transport properties. Nature Communications, 2012, 3, 949.	12.8	174