

# Luming Meng

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/731421/publications.pdf>

Version: 2024-02-01

10  
papers

213  
citations

1478505

6  
h-index

1474206

9  
g-index

12  
all docs

12  
docs citations

12  
times ranked

333  
citing authors

#	ARTICLE	IF	CITATIONS
1	From 1D sequence to 3D chromatin dynamics and cellular functions: a phase separation perspective. <i>Nucleic Acids Research</i> , 2018, 46, 9367-9383.	14.5	51
2	Structural Modeling of Chromatin Integrates Genome Features and Reveals Chromosome Folding Principle. <i>Scientific Reports</i> , 2017, 7, 2818.	3.3	42
3	DNA Methylation Landscape Reflects the Spatial Organization of Chromatin in Different Cells. <i>Biophysical Journal</i> , 2017, 113, 1395-1404.	0.5	39
4	Superresolution imaging reveals spatiotemporal propagation of human replication foci mediated by CTCF-organized chromatin structures. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 15036-15046.	7.1	27
5	Unexpected aggregation induced circular dichroism, circular polarized luminescence and helical assembly from achiral hexaphenylsilole (HPS). <i>RSC Advances</i> , 2017, 7, 24841-24847.	3.6	25
6	DeepAntigen: a novel method for neoantigen prioritization via 3D genome and deep sparse learning. <i>Bioinformatics</i> , 2020, 36, 4894-4901.	4.1	17
7	Si-C is a method for inferring super-resolution intact genome structure from single-cell Hi-C data. <i>Nature Communications</i> , 2021, 12, 4369.	12.8	9
8	A novel neoantigen discovery approach based on chromatin high order conformation. <i>BMC Medical Genomics</i> , 2020, 13, 62.	1.5	2
9	3D genome assisted protein-protein interaction prediction. <i>Future Generation Computer Systems</i> , 2022, 137, 87-96.	7.5	1
10	Improving Protein-protein Interaction Prediction by Incorporating 3D Genome Information. <i>Lecture Notes in Computer Science</i> , 2021, , 511-520.	1.3	0