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List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4737770/publications.pdf

Version: 2024-02-01

20 1,984 14 20 papers citations h-index g-index

25 25 25 25 2502

25 25 25 2502 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Single-nucleus Hi-C reveals unique chromatin reorganization at oocyte-to-zygote transition. Nature, 2017, 544, 110-114.	13.7	604
2	Active chromatin and transcription play a key role in chromosome partitioning into topologically associating domains. Genome Research, 2016, 26, 70-84.	2.4	311
3	A mechanism of cohesinâ€dependent loop extrusion organizes zygotic genome architecture. EMBO Journal, 2017, 36, 3600-3618.	3.5	291
4	Developmentally regulated <i>Shh</i> expression is robust to TAD perturbations. Development (Cambridge), 2019, 146, .	1.2	111
5	<i>Coolpup.py:</i> versatile pile-up analysis of Hi-C data. Bioinformatics, 2020, 36, 2980-2985.	1.8	111
6	A central role for canonical PRC1 in shaping the 3D nuclear landscape. Genes and Development, 2020, 34, 931-949.	2.7	100
7	Nuclear lamina integrity is required for proper spatial organization of chromatin in Drosophila. Nature Communications, 2019, 10, 1176.	5.8	83
8	DNA Methylation Directs Polycomb-Dependent 3D Genome Re-organization in Naive Pluripotency. Cell Reports, 2019, 29, 1974-1985.e6.	2.9	76
9	MCM complexes are barriers that restrict cohesin-mediated loop extrusion. Nature, 2022, 606, 197-203.	13.7	58
10	Order and stochasticity in the folding of individual Drosophila genomes. Nature Communications, 2021, 12, 41.	5.8	49
11	Bivalent promoter hypermethylation in cancer is linked to the H327me3/H3K4me3 ratio in embryonic stem cells. BMC Biology, 2020, 18, 25.	1.7	35
12	Nuclear organisation and replication timing are coupled through RIF1–PP1 interaction. Nature Communications, 2021, 12, 2910.	5.8	29
13	Activation of the alpha-globin gene expression correlates with dramatic upregulation of nearby non-globin genes and changes in local and large-scale chromatin spatial structure. Epigenetics and Chromatin, 2017, 10, 35.	1.8	19
14	Quantitative differences in TAD border strength underly the TAD hierarchy in <i>Drosophila</i> chromosomes. Journal of Cellular Biochemistry, 2019, 120, 4494-4503.	1.2	17
15	C-TALE, a new cost-effective method for targeted enrichment of Hi-C/3C-seq libraries. Methods, 2020, 170, 48-60.	1.9	13
16	Evolution of the Genome 3D Organization: Comparison of Fused and Segregated Globin Gene Clusters. Molecular Biology and Evolution, 2017, 34, 1492-1504.	3.5	10
17	Single-nucleus Hi-C of mammalian oocytes and zygotes. Methods in Cell Biology, 2018, 144, 389-407.	0.5	9
18	From bedside to bench: regulation of host factors in SARS-CoV-2 infection. Experimental and Molecular Medicine, 2021, 53, 483-494.	3.2	6

#	Article	IF	CITATION
19	A Mechanism of Cohesin-Dependent Loop Extrusion Organizes Mammalian Chromatin Structure in the Developing Embryo. Biophysical Journal, 2018, 114, 255a.	0.2	4
20	Low Level of Expression of C-Terminally Truncated Human FUS Causes Extensive Changes in the Spinal Cord Transcriptome of Asymptomatic Transgenic Mice. Neurochemical Research, 2020, 45, 1168-1179.	1.6	3