

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Nutritional control regulates symbiont proliferation and life history in coral-dinoflagellate symbiosis. BMC Biology, 2022, 20, 103.	3.8	17
2	Genetic and spatial organization of the unusual chromosomes of the dinoflagellate Symbiodinium microadriaticum. Nature Genetics, 2021, 53, 618-629.	21.4	54
3	Chromosome-Level Assembly of the Atlantic Silverside Genome Reveals Extreme Levels of Sequence Diversity and Structural Genetic Variation. Genome Biology and Evolution, 2021, 13, .	2.5	20
4	Symbiodinium microadriaticum (coral microalgal endosymbiont). Trends in Genetics, 2021, 37, 1044-1045.	6.7	3
5	Transcriptional Silencers in Drosophila Serve a Dual Role as Transcriptional Enhancers in Alternate Cellular Contexts. Molecular Cell, 2020, 77, 324-337.e8.	9.7	85
6	SPEN integrates transcriptional and epigenetic control of X-inactivation. Nature, 2020, 578, 455-460.	27.8	146
7	Measuring the reproducibility and quality of Hi-C data. Genome Biology, 2019, 20, 57.	8.8	125
8	The non-canonical SMC protein SmcHD1 antagonises TAD formation and compartmentalisation on the inactive X chromosome. Nature Communications, 2019, 10, 30.	12.8	87
9	Integrative detection and analysis of structural variation in cancer genomes. Nature Genetics, 2018, 50, 1388-1398.	21.4	268
10	The <i>HoxD</i> cluster is a dynamic and resilient TAD boundary controlling the segregation of antagonistic regulatory landscapes. Genes and Development, 2017, 31, 2264-2281.	5.9	155
11	Structural organization of the inactive X chromosome in the mouse. Nature, 2016, 535, 575-579.	27.8	369
12	High-Affinity Sites Form an Interaction Network to Facilitate Spreading of the MSL Complex across the X Chromosome in Drosophila. Molecular Cell, 2015, 60, 146-162.	9.7	70
13	Genome-wide Maps of Nuclear Lamina Interactions in Single Human Cells. Cell, 2015, 163, 134-147.	28.9	399
14	Organization of the Mitotic Chromosome. Science, 2013, 342, 948-953.	12.6	894
15	Hi–C: A comprehensive technique to capture the conformation of genomes. Methods, 2012, 58, 268-276.	3.8	792
16	Mediator and cohesin connect gene expression and chromatin architecture. Nature, 2010, 467, 430-435.	27.8	1,707