## M Jordan Rowley

List of Publications by Year in descending order

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M IOPDAN ROWLEY

#	Article	IF	CITATIONS
1	Organizational principles of 3D genome architecture. Nature Reviews Genetics, 2018, 19, 789-800.	16.3	832
2	Evolutionarily Conserved Principles Predict 3D Chromatin Organization. Molecular Cell, 2017, 67, 837-852.e7.	9.7	458
3	Widespread long-range cis-regulatory elements in the maize genome. Nature Plants, 2019, 5, 1237-1249.	9.3	250
4	A SWI/SNF Chromatin-Remodeling Complex Acts in Noncoding RNA-Mediated Transcriptional Silencing. Molecular Cell, 2013, 49, 298-309.	9.7	178
5	Spatial and functional relationships among Pol V-associated loci, Pol IV-dependent siRNAs, and cytosine methylation in the <i>Arabidopsis</i> epigenome. Genes and Development, 2012, 26, 1825-1836.	5.9	137
6	A Dicer-Independent Route for Biogenesis of siRNAs that Direct DNA Methylation in Arabidopsis. Molecular Cell, 2016, 61, 222-235.	9.7	134
7	DifferentÂenhancer classes in Drosophila bind distinct architectural proteins and mediate unique chromatin interactions and 3D architecture. Nucleic Acids Research, 2017, 45, 1714-1730.	14.5	133
8	Immediate and deferred epigenomic signatures of in vivo neuronal activation in mouse hippocampus. Nature Neuroscience, 2019, 22, 1718-1730.	14.8	114
9	The three-dimensional genome: principles and roles of long-distance interactions. Current Opinion in Cell Biology, 2016, 40, 8-14.	5.4	113
10	Condensin II Counteracts Cohesin and RNA Polymerase II in the Establishment of 3D Chromatin Organization. Cell Reports, 2019, 26, 2890-2903.e3.	6.4	97
11	<scp>RNA</scp> â€directed <scp>DNA</scp> methylation requires stepwise binding of silencing factors to long nonâ€coding <scp>RNA</scp> . Plant Journal, 2014, 79, 181-191.	5.7	83
12	Maintenance of CTCF- and Transcription Factor-Mediated Interactions from the Gametes to the Early Mouse Embryo. Molecular Cell, 2019, 75, 154-171.e5.	9.7	81
13	Long non-coding RNA produced by RNA polymerase V determines boundaries of heterochromatin. ELife, 2016, 5, .	6.0	76
14	Analysis of Hi-C data using SIP effectively identifies loops in organisms from <i>C. elegans</i> to mammals. Genome Research, 2020, 30, 447-458.	5.5	70
15	Epigenetic regulation of Plasmodium falciparum clonally variant gene expression during development in Anopheles gambiae. Scientific Reports, 2017, 7, 40655.	3.3	69
16	Independent Chromatin Binding of ARGONAUTE4 and SPT5L/KTF1 Mediates Transcriptional Gene Silencing. PLoS Genetics, 2011, 7, e1002120.	3.5	62
17	Architectural Proteins and Pluripotency Factors Cooperate to Orchestrate the Transcriptional Response of hESCs to Temperature Stress. Molecular Cell, 2018, 71, 940-955.e7.	9.7	62
18	<scp>RNA</scp> polymerase <scp>V</scp> targets transcriptional silencing components to promoters of proteinâ€coding genes. Plant Journal, 2013, 73, 179-189.	5.7	61

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19	Long-range control of gene expression via RNA-directed DNA methylation. PLoS Genetics, 2017, 13, e1006749.	3.5	33
20	Ecdysone-Induced 3D Chromatin Reorganization Involves Active Enhancers Bound by Pipsqueak and Polycomb. Cell Reports, 2019, 28, 2715-2727.e5.	6.4	32
21	Analysis of long non-coding RNAs produced by a specialized RNA polymerase in Arabidopsis thaliana. Methods, 2013, 63, 160-169.	3.8	31
22	The inhibition of LSD1 via sequestration contributes to tau-mediated neurodegeneration. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 29133-29143.	7.1	24
23	Sex-specific multi-level 3D genome dynamics in the mouse brain. Nature Communications, 2022, 13, .	12.8	15
24	Evolutionary History and Activity of RNase H1-Like Proteins in <i>Arabidopsis thaliana</i> . Plant and Cell Physiology, 2020, 61, 1107-1119.	3.1	12
25	Implications of Dosage Deficiencies in CTCF and Cohesin on Genome Organization, Gene Expression, and Human Neurodevelopment. Genes, 2022, 13, 583.	2.4	10
26	Ecdysoneless Protein Regulates Viral and Cellular mRNA Splicing to Promote Cervical Oncogenesis. Molecular Cancer Research, 2022, 20, 305-318.	3.4	6
27	Capturing native interactions: intrinsicÂmethods to study chromatinÂconformation. Molecular Systems Biology, 2016, 12, 897.	7.2	4
28	Minute-Made Data Analysis: Tools for Rapid Interrogation of Hi-C Contacts. Molecular Cell, 2016, 64, 9-11.	9.7	4
29	Elevating SOX2 Downregulates MYC through a SOX2:MYC Signaling Axis and Induces a Slowly Cycling Proliferative State in Human Tumor Cells. Cancers, 2022, 14, 1946.	3.7	4
30	Analysis of Chromatin Interactions Mediated by Specific Architectural Proteins in Drosophila Cells. Methods in Molecular Biology, 2018, 1766, 239-256.	0.9	1
31	Somatic Diversification of Rearranged Antibody Gene Segments by Intra- and Interchromosomal Templated Mutagenesis. Journal of Immunology, 2022, , ji2100434.	0.8	0