Karen L Reddy

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	vrille, Pdp1, and dClock Form a Second Feedback Loop in the Drosophila Circadian Clock. Cell, 2003, 112, 329-341.	28.9	474
2	DNA Sequence-Dependent Compartmentalization and Silencing of Chromatin at the Nuclear Lamina. Cell, 2012, 149, 1474-1487.	28.9	405
3	Directed targeting of chromatin to the nuclear lamina is mediated by chromatin state and A-type lamins. Journal of Cell Biology, 2015, 208, 33-52.	5.2	266
4	Regulation of B cell fate commitment and immunoglobulin heavy-chain gene rearrangements by Ikaros. Nature Immunology, 2008, 9, 927-936.	14.5	228
5	Assembling a Gene Regulatory Network for Specification of the B Cell Fate. Developmental Cell, 2004, 7, 607-617.	7.0	212
6	Regulation of interleukin 7–dependent immunoglobulin heavy-chain variable gene rearrangements by transcription factor STAT5. Nature Immunology, 2005, 6, 836-843.	14.5	131
7	Altered Chromosomal Positioning, Compaction, and Gene Expression with a Lamin A/C Gene Mutation. PLoS ONE, 2010, 5, e14342.	2.5	111
8	Higher order chromatin organization in cancer. Seminars in Cancer Biology, 2013, 23, 109-115.	9.6	83
9	BioSITe: A Method for Direct Detection and Quantitation of Site-Specific Biotinylation. Journal of Proteome Research, 2018, 17, 759-769.	3.7	70
10	Initiation of allelic exclusion by stochastic interaction of Tcrb alleles with repressive nuclear compartments. Nature Immunology, 2008, 9, 802-809.	14.5	68
11	Genome regulation at the peripheral zone: lamina associated domains in development and disease. Current Opinion in Genetics and Development, 2014, 25, 50-61.	3.3	66
12	NET gains and losses: the role of changing nuclear envelope proteomes in genome regulation. Current Opinion in Cell Biology, 2014, 28, 105-120.	5.4	60
13	Methylation of histone H3K23 blocks DNA damage in pericentric heterochromatin during meiosis. ELife, 2014, 3, e02996.	6.0	51
14	The Drosophila PAR Domain Protein 1 (Pdp1) Gene Encodes Multiple Differentially Expressed mRNAs and Proteins through the Use of Multiple Enhancers and Promoters. Developmental Biology, 2000, 224, 401-414.	2.0	42
15	A Lamina-Associated Domain Border Governs Nuclear Lamina Interactions, Transcription, and Recombination of the Tcrb Locus. Cell Reports, 2018, 25, 1729-1740.e6.	6.4	37
16	Nuclear lamin isoforms differentially contribute to LINC complex-dependent nucleocytoskeletal coupling and whole-cell mechanics. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2121816119.	7.1	33
17	The Nuclear Lamina. Cold Spring Harbor Perspectives in Biology, 2022, 14, a040113.	5.5	28
18	An Accessible Proteogenomics Informatics Resource for Cancer Researchers. Cancer Research, 2017, 77, e43-e46.	0.9	27

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19	The High Mobility Group A1 (HMGA1) gene is highly overexpressed in human uterine serous carcinomas and carcinosarcomas and drives Matrix Metalloproteinase-2 (MMP-2) in a subset of tumors. Gynecologic Oncology, 2016, 141, 580-587.	1.4	26
20	Mapping the micro-proteome of the nuclear lamina and lamina-associated domains. Life Science Alliance, 2021, 4, e202000774.	2.8	26
21	The shifting shape of genomes: dynamics of heterochromatin interactions at the nuclear lamina. Current Opinion in Genetics and Development, 2021, 67, 163-173.	3.3	25
22	Lamin C is required to establish genome organization after mitosis. Genome Biology, 2021, 22, 305.	8.8	24
23	HMGA1 chromatin regulators induce transcriptional networks involved in GATA2 and proliferation during MPN progression. Blood, 2022, 139, 2797-2815.	1.4	20
24	The Drosophila Par domain protein I gene, Pdp1, is a regulator of larval growth, mitosis and endoreplication. Developmental Biology, 2006, 289, 100-114.	2.0	16
25	Molecular Pathways and Mechanisms Regulating the Recombination of Immunoglobulin Genes during B-Lymphocyte Development. Advances in Experimental Medicine and Biology, 2009, 650, 133-147.	1.6	14
26	Prediction of Gene Activity in Early B Cell Development Based on an Integrative Multi-Omics Analysis. Journal of Proteomics and Bioinformatics, 2014, 07, .	0.4	13
27	Tagged Chromosomal Insertion Site System. Methods in Enzymology, 2016, 569, 433-453.	1.0	6
28	Finding the Middlemen in Genome Organization. Developmental Cell, 2015, 35, 670-671.	7.0	4
29	The Nuclear Lamina and Genome Organization. , 2018, , 321-343.		2
30	Differential Signaling through p190 and p210 Forms of BCR-ABL Fusion Proteins Revealed By Proteomic Analysis. Blood, 2015, 126, 3651-3651.	1.4	1
31	Abstract 2414: HMGA1 induces <i>FGF19</i> to drive tumor progression and recruit cancer associated fibroblasts in pancreatic adenocarcinoma. Cancer Research, 2021, 81, 2414-2414.	0.9	0
32	Abstract 2666: HMGA1: An epigenetic switch required for MPN progression by inducingGATA-2and cell cycle progression through enhancer rewiring. , 2021, , .		0
33	Proteomic/Transcriptomic Signatures of Infant MLL-r Rearranged B-ALL at Diagnosis and Relapse Reveal Lineage Plasticity and Diagnostic Heterogeneity. Blood, 2016, 128, 2697-2697.	1.4	0
34	HMGA1 Chromatin Regulators Drive Progression in Myeloproliferative Neoplasms through Epigenetic Rewiring to Induce Networks Involved in GATA2 and Proliferation. Blood, 2021, 138, 625-625.	1.4	0