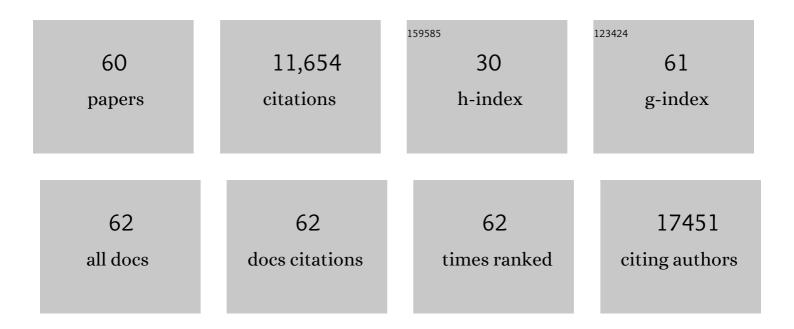


List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	High-Resolution Profiling of Histone Methylations in the Human Genome. Cell, 2007, 129, 823-837.	28.9	6,036
2	Dynamic Regulation of Nucleosome Positioning in the Human Genome. Cell, 2008, 132, 887-898.	28.9	1,211
3	Global Mapping of H3K4me3 and H3K27me3 Reveals Specificity and Plasticity in Lineage Fate Determination of Differentiating CD4+ T Cells. Immunity, 2009, 30, 155-167.	14.3	1,005
4	H3.3/H2A.Z double variant–containing nucleosomes mark 'nucleosome-free regions' of active promoters and other regulatory regions. Nature Genetics, 2009, 41, 941-945.	21.4	679
5	Genome-wide Analyses of Transcription Factor GATA3-Mediated Gene Regulation in Distinct T Cell Types. Immunity, 2011, 35, 299-311.	14.3	293
6	A transcriptomic analysis of superhybrid rice <i>LYP9</i> and its parents. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7695-7701.	7.1	184
7	Detection of single nucleotide variations in expressed exons of the human genome using RNA-Seq. Nucleic Acids Research, 2009, 37, e106-e106.	14.5	152
8	Rutin ameliorates obesity through brown fat activation. FASEB Journal, 2017, 31, 333-345.	0.5	151
9	Brown adipose tissue transplantation ameliorates polycystic ovary syndrome. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 2708-2713.	7.1	141
10	Critical role of histone demethylase Jmjd3 in the regulation of CD4+ T-cell differentiation. Nature Communications, 2014, 5, 5780.	12.8	136
11	Iron-dependent histone 3 lysine 9 demethylation controls B cell proliferation and humoral immune responses. Nature Communications, 2019, 10, 2935.	12.8	107
12	The Nuclear Matrix Protein SAFB Cooperates with Major Satellite RNAs to Stabilize Heterochromatin Architecture Partially through Phase Separation. Molecular Cell, 2020, 77, 368-383.e7.	9.7	104
13	Identification of Integrator-PP2A complex (INTAC), an RNA polymerase II phosphatase. Science, 2020, 370,	12.6	104
14	Enhanced PAPSS2/VCAN sulfation axis is essential for Snail-mediated breast cancer cell migration and metastasis. Cell Death and Differentiation, 2019, 26, 565-579.	11.2	91
15	MiR-17-5p enhances pancreatic cancer proliferation by altering cell cycle profiles via disruption of RBL2/E2F4-repressing complexes. Cancer Letters, 2018, 412, 59-68.	7.2	75
16	USP21 prevents the generation of T-helper-1-like Treg cells. Nature Communications, 2016, 7, 13559.	12.8	67
17	Foxp1/2/4 regulate endochondral ossification as a suppresser complex. Developmental Biology, 2015, 398, 242-254.	2.0	62
18	Global histone modification profiling reveals the epigenomic dynamics during malignant transformation in a four-stage breast cancer model. Clinical Epigenetics, 2016, 8, 34.	4.1	61

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19	SPOP-containing complex regulates SETD2 stability and H3K36me3-coupled alternative splicing. Nucleic Acids Research, 2017, 45, 92-105.	14.5	60
20	Brown adipose tissue activation by rutin ameliorates polycystic ovary syndrome in rat. Journal of Nutritional Biochemistry, 2017, 47, 21-28.	4.2	59
21	Differential stem cell aging kinetics in Hutchinson-Gilford progeria syndrome and Werner syndrome. Protein and Cell, 2018, 9, 333-350.	11.0	56
22	Improved nucleosome-positioning algorithm iNPS for accurate nucleosome positioning from sequencing data. Nature Communications, 2014, 5, 4909.	12.8	55
23	Myricetin-induced brown adipose tissue activation prevents obesity and insulin resistance in db/db mice. European Journal of Nutrition, 2018, 57, 391-403.	3.9	52
24	GFRα2 prompts cell growth and chemoresistance through down-regulating tumor suppressor gene PTEN via Mir-17-5p in pancreatic cancer. Cancer Letters, 2016, 380, 434-441.	7.2	51
25	Stage-Dependent and Locus-Specific Role of Histone Demethylase Jumonji D3 (JMJD3) in the Embryonic Stages of Lung Development. PLoS Genetics, 2014, 10, e1004524.	3.5	50
26	The chromatin remodeler Chd4 maintains embryonic stem cell identity by controlling pluripotency- and differentiation-associated genes. Journal of Biological Chemistry, 2017, 292, 8507-8519.	3.4	46
27	The thermogenic activity of adjacent adipocytes fuels the progression of ccRCC and compromises anti-tumor therapeutic efficacy. Cell Metabolism, 2021, 33, 2021-2039.e8.	16.2	45
28	Distinct Gene Expression and Epigenetic Signatures in Hepatocyte-like Cells Produced by Different Strategies from the Same Donor. Stem Cell Reports, 2017, 9, 1813-1824.	4.8	37
29	Deletion of Macrophage Mineralocorticoid Receptor Protects Hepatic Steatosis and Insulin Resistance Through ERI±/HGF/Met Pathway. Diabetes, 2017, 66, 1535-1547.	0.6	36
30	SH3RF3 promotes breast cancer stem-like properties via JNK activation and PTX3 upregulation. Nature Communications, 2020, 11, 2487.	12.8	35
31	Large-scale chromatin reorganization reactivates placenta-specific genes that drive cellular aging. Developmental Cell, 2022, 57, 1347-1368.e12.	7.0	32
32	Coordinated regulation of infection-related morphogenesis by the KMT2-Cre1-Hyd4 regulatory pathway to facilitate fungal infection. Science Advances, 2020, 6, eaaz1659.	10.3	31
33	The Selective Activation of p53 Target Genes Regulated by SMYD2 in BIX-01294 Induced Autophagy-Related Cell Death. PLoS ONE, 2015, 10, e0116782.	2.5	29
34	Genome-Wide Mapping of Nucleosome Occupancy, Histone Modifications, and Gene Expression Using Next-Generation Sequencing Technology. Methods in Enzymology, 2012, 513, 297-313.	1.0	24
35	Retinoid signaling controls spermatogonial differentiation by regulating expression of replication-dependent core histone genes. Development (Cambridge), 2016, 143, 1502-11.	2.5	23
36	A positive role for polycomb in transcriptional regulation via H4K20me1. Cell Research, 2016, 26, 529-542.	12.0	18

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37	Baf60b-mediated ATM-p53 activation blocks cell identity conversion by sensing chromatin opening. Cell Research, 2017, 27, 642-656.	12.0	18
38	Changes in white and brown adipose tissue microRNA expression in cold-induced mice. Biochemical and Biophysical Research Communications, 2015, 463, 193-199.	2.1	17
39	Integrated Chromatin Accessibility and Transcriptome Landscapes of Doxorubicin-Resistant Breast Cancer Cells. Frontiers in Cell and Developmental Biology, 2021, 9, 708066.	3.7	17
40	Rrp6 Regulates Heterochromatic Gene Silencing via ncRNA RUF6 Decay in Malaria Parasites. MBio, 2020, 11, .	4.1	15
41	Indirubin, a small molecular deriving from connectivity map (CMAP) screening, ameliorates obesity-induced metabolic dysfunction by enhancing brown adipose thermogenesis and white adipose browning. Nutrition and Metabolism, 2020, 17, 21.	3.0	15
42	SMYD2 Drives Mesendodermal Differentiation of Human Embryonic Stem Cells Through Mediating the Transcriptional Activation of Key Mesendodermal Genes. Stem Cells, 2019, 37, 1401-1415.	3.2	14
43	Accelerated evolution of an Lhx2 enhancer shapes mammalian social hierarchies. Cell Research, 2020, 30, 408-420.	12.0	14
44	The CTCF/LncRNAâ€PACERR complex recruits E1A binding protein p300 to induce proâ€ŧumour macrophages in pancreatic ductal adenocarcinoma via directly regulating PTGS2 expression. Clinical and Translational Medicine, 2022, 12, e654.	4.0	14
45	Reconstruction of the Gene Regulatory Network Involved in the Sonic Hedgehog Pathway with a Potential Role in Early Development of the Mouse Brain. PLoS Computational Biology, 2014, 10, e1003884.	3.2	13
46	Inhibition of XBP1s ubiquitination enhances its protein stability and improves glucose homeostasis. Metabolism: Clinical and Experimental, 2020, 105, 154046.	3.4	12
47	Triterpenoids from functional mushroom Ganoderma resinaceum and the novel role of Resinacein S in enhancing the activity of brown/beige adipocytes. Food Research International, 2020, 136, 109303.	6.2	12
48	Polycomb Group Gene E(z) Is Required for Spermatogonial Dedifferentiation in Drosophila Adult Testis. Journal of Molecular Biology, 2017, 429, 2030-2041.	4.2	11
49	Epigenomic analysis in a cell-based model reveals the roles of H3K9me3 in breast cancer transformation. Epigenomics, 2017, 9, 1077-1092.	2.1	11
50	Membrane-bound TNF mediates microtubule-targeting chemotherapeutics-induced cancer cytolysis via juxtacrine inter-cancer-cell death signaling. Cell Death and Differentiation, 2020, 27, 1569-1587.	11.2	11
51	The Architectural Factor HMGB1 Is Involved in Genome Organization in the Human Malaria Parasite Plasmodium falciparum. MBio, 2021, 12, .	4.1	11
52	Independent manipulation of histone H3 modifications in individual nucleosomes reveals the contributions of sister histones to transcription. ELife, 2017, 6, .	6.0	8
53	Spautin-A41 Attenuates Cerulein-Induced Acute Pancreatitis through Inhibition of Dysregulated Autophagy. Biological and Pharmaceutical Bulletin, 2019, 42, 1789-1798.	1.4	8
54	Comparative Transcriptional Profiling of Three Super-Hybrid Rice Combinations. International Journal of Molecular Sciences, 2014, 15, 3799-3815.	4.1	7

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55	Autonomous Bionanorobots <i>via</i> a Cage-Shaped Silsesquioxane Vehicle for <i>In Vivo</i> Heavy Metal Detoxification. ACS Applied Materials & Interfaces, 2022, 14, 29238-29249.	8.0	7
56	Perineural invasion is related to p38 mitogenâ€activated protein kinase pathway activation and promotes tumor growth and chemoresistance in pancreatic cancer. Journal of Cellular Biochemistry, 2019, 120, 11775-11783.	2.6	5
57	3C-based methods to detect long-range chromatin interactions. Frontiers in Biology, 2011, 6, 76-81.	0.7	4
58	Pre-existing chromatin accessibility of switchable repressive compartment delineates cell plasticity. National Science Review, 2022, 9, .	9.5	4
59	Understanding human diseases with high-throughput quantitative measurement and analysis of molecular signatures. Science China Life Sciences, 2013, 56, 213-219.	4.9	3
60	Establishment and Investigation of a Multiple Gene Expression Signature to Predict Long-Term Survival in Pancreatic Cancer. BioMed Research International, 2020, 2020, 1-20.	1.9	1