Fengbiao Mao

List of Publications by Year in descending order

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43 papers

1,119 citations

393982 19 h-index 454577 30 g-index

46 all docs

46 docs citations

46 times ranked

1927 citing authors

#	Article	IF	Citations
1	HOXA9 Reprograms the Enhancer Landscape to Promote Leukemogenesis. Cancer Cell, 2018, 34, 643-658.e5.	7.7	94
2	Genetic landscape of papillary thyroid carcinoma in the Chinese population. Journal of Pathology, 2018, 244, 215-226.	2.1	90
3	Antiâ€Friction MSCs Delivery System Improves the Therapy for Severe Osteoarthritis. Advanced Materials, 2021, 33, e2104758.	11.1	66
4	OncoVar: an integrated database and analysis platform for oncogenic driver variants in cancers. Nucleic Acids Research, 2021, 49, D1289-D1301.	6.5	64
5	Transcriptomic signature associated with carcinogenesis and aggressiveness of papillary thyroid carcinoma. Theranostics, 2018, 8, 4345-4358.	4.6	63
6	TET1 modulates H4K16 acetylation by controlling auto-acetylation of hMOF to affect gene regulation and DNA repair function. Nucleic Acids Research, 2017, 45, 672-684.	6.5	56
7	RBP-Var: a database of functional variants involved in regulation mediated by RNA-binding proteins. Nucleic Acids Research, 2016, 44, D154-D163.	6.5	52
8	Population Genomics Reveals Speciation and Introgression between Brown Norway Rats and Their Sibling Species. Molecular Biology and Evolution, 2017, 34, 2214-2228.	3.5	47
9	RNAi-dependent <i>Polycomb</i> repression controls transposable elements in <i>Tetrahymena</i> Genes and Development, 2019, 33, 348-364.	2.7	42
10	ER-associated degradation preserves hematopoietic stem cell quiescence and self-renewal by restricting mTOR activity. Blood, 2020, 136, 2975-2986.	0.6	40
11	Ras-induced Epigenetic Inactivation of the RRAD (Ras-related Associated with Diabetes) Gene Promotes Glucose Uptake in a Human Ovarian Cancer Model. Journal of Biological Chemistry, 2014, 289, 14225-14238.	1.6	37
12	Anxiety-Related Behaviours Associated with microRNA-206-3p and BDNF Expression in Pregnant Female Mice Following Psychological Social Stress. Molecular Neurobiology, 2018, 55, 1097-1111.	1.9	37
13	Histone Acetyltransferase MOF Blocks Acquisition of Quiescence in Ground-State ESCs through Activating Fatty Acid Oxidation. Cell Stem Cell, 2020, 27, 441-458.e10.	5.2	37
14	OncoBase: a platform for decoding regulatory somatic mutations in human cancers. Nucleic Acids Research, 2019, 47, D1044-D1055.	6.5	33
15	Prevalence and architecture of posttranscriptionally impaired synonymous mutations in 8,320 genomes across 22 cancer types. Nucleic Acids Research, 2020, 48, 1192-1205.	6.5	31
16	CirGRDB: a database for the genome-wide deciphering circadian genes and regulators. Nucleic Acids Research, 2018, 46, D64-D70.	6.5	29
17	A Polycomb repressive complex is required for RNAi-mediated heterochromatin formation and dynamic distribution of nuclear bodies. Nucleic Acids Research, 2021, 49, 5407-5425.	6.5	27
18	Q-RRBS: a quantitative reduced representation bisulfite sequencing method for single-cell methylome analyses. Epigenetics, 2015, 10, 775-783.	1.3	23

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19	Functional analysis of the methyltransferase SMYD in the single-cell model organism Tetrahymena thermophila. Marine Life Science and Technology, 2020, 2, 109-122.	1.8	22
20	Mechanism for DPY30 and ASH2L intrinsically disordered regions to modulate the MLL/SET1 activity on chromatin. Nature Communications, 2021, 12, 2953.	5.8	21
21	CircleBase: an integrated resource and analysis platform for human eccDNAs. Nucleic Acids Research, 2022, 50, D72-D82.	6.5	20
22	Al-Driver: an ensemble method for identifying driver mutations in personal cancer genomes. NAR Genomics and Bioinformatics, 2020, 2, Iqaa084.	1.5	19
23	Comprehensive evaluation of computational methods for predicting cancer driver genes. Briefings in Bioinformatics, 2022, 23, .	3.2	19
24	Transcriptomic signatures and repurposing drugs for COVID-19 patients: findings of bioinformatics analyses. Computational and Structural Biotechnology Journal, 2021, 19, 1-15.	1.9	18
25	EpiDenovo: a platform for linking regulatory de novo mutations to developmental epigenetics and diseases. Nucleic Acids Research, 2018, 46, D92-D99.	6.5	17
26	p53 Integrates Temporal WDR5 Inputs during Neuroectoderm and Mesoderm Differentiation of Mouse Embryonic Stem Cells. Cell Reports, 2020, 30, 465-480.e6.	2.9	17
27	Discovery of first-in-class inhibitors of ASH1L histone methyltransferase with anti-leukemic activity. Nature Communications, 2021, 12, 2792.	5.8	17
28	Comparative RNA-seq analysis reveals potential mechanisms mediating the conversion to androgen independence in an LNCaP progression cell model. Cancer Letters, 2014, 342, 130-138.	3.2	16
29	Whole-Genome Sequencing Reveals Genetic Variation in the Asian House Rat. G3: Genes, Genomes, Genetics, 2016, 6, 1969-1977.	0.8	8
30	Whole-exome sequencing identifies a somatic missense mutation of NBN in clear cell sarcoma of the salivary gland. Oncology Reports, 2016, 35, 3349-3356.	1,2	8
31	MLL1 Inhibition and Vitamin D Signaling Cooperate to Facilitate the Expanded Pluripotency State. Cell Reports, 2019, 29, 2659-2671.e6.	2.9	8
32	Identification of a novel missense (C7W) mutation of SOD1 in a large familial amyotrophic lateral sclerosis pedigree. Neurobiology of Aging, 2014, 35, 725.e11-725.e15.	1.5	6
33	MBRidge: an accurate and cost-effective method for profiling DNA methylome at single-base resolution. Journal of Molecular Cell Biology, 2015, 7, 299-313.	1.5	5
34	Genome-wide identification and divergent transcriptional expression of StAR-related lipid transfer (START) genes in teleosts. Gene, 2013, 519, 18-25.	1.0	4
35	A Bayesian Framework to Identify Methylcytosines from High-Throughput Bisulfite Sequencing Data. PLoS Computational Biology, 2014, 10, e1003853.	1.5	4
36	p53 inactivation unmasks histone methylation-independent WDR5 functions that drive self-renewal and differentiation of pluripotent stem cells. Stem Cell Reports, 2021, 16, 2642-2658.	2.3	4

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37	A novel 10-base pair insertion mutation in exon 5 of the SOD1 gene in a Chinese family with amyotrophic lateral sclerosis. Neurobiology of Aging, 2016, 45, 212.e1-212.e4.	1.5	3
38	Prenatal witness stress induces intergenerational anxiety-like behaviors and altered gene expression profiles in male mice. Neuropharmacology, 2022, 202, 108857.	2.0	3
39	Natural Products as a Source for New Leads in Depression Treatment. Evidence-based Complementary and Alternative Medicine, 2022, 2022, 1-2.	0.5	3
40	Genetic association and single-cell transcriptome analyses reveal distinct features connecting autoimmunity with cancers. IScience, 2022, 25, 104631.	1.9	3
41	p53 Coordinates Temporal WDR5 Inputs During Neuroectoderm and Mesoderm Differentiation of Mouse Embryonic Stem Cells. SSRN Electronic Journal, 0, , .	0.4	1
42	Identification and analysis of intermediate-size noncoding RNAs in the rhesus macaque fetal brain. Journal of Genetics and Genomics, 2017, 44, 171-174.	1.7	0
43	Transcriptomic Signatures and Immunomodulatory Treatments for COVID-19 Patients. SSRN Electronic Journal, 0, , .	0.4	O