

Zhao Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

41
papers

2,300
citations

304743

22
h-index

302126

39
g-index

41
all docs

41
docs citations

41
times ranked

3998
citing authors

#	ARTICLE	IF	CITATIONS
1	PD1Hi CD8+ T cells correlate with exhausted signature and poor clinical outcome in hepatocellular carcinoma. , 2019, 7, 331.		213
2	PancanQTL: systematic identification of cis-eQTLs and trans-eQTLs in 33 cancer types. Nucleic Acids Research, 2018, 46, D971-D976.	14.5	191
3	A Pan-cancer Analysis of the Expression and Clinical Relevance of Small Nucleolar RNAs in Human Cancer. Cell Reports, 2017, 21, 1968-1981.	6.4	186
4	Transcriptional landscape and clinical utility of enhancer RNAs for eRNA-targeted therapy in cancer. Nature Communications, 2019, 10, 4562.	12.8	165
5	Characterization of hypoxia-associated molecular features to aid hypoxia-targeted therapy. Nature Metabolism, 2019, 1, 431-444.	11.9	158
6	Comprehensive characterization of circular RNAs in ~1000 human cancer cell lines. Genome Medicine, 2019, 11, 55.	8.2	116
7	Comprehensive Characterization of Alternative Polyadenylation in Human Cancer. Journal of the National Cancer Institute, 2018, 110, 379-389.	6.3	111
8	Activated and Exhausted MAIT Cells Foster Disease Progression and Indicate Poor Outcome in Hepatocellular Carcinoma. Clinical Cancer Research, 2019, 25, 3304-3316.	7.0	109
9	CCL15 Recruits Suppressive Monocytes to Facilitate Immune Escape and Disease Progression in Hepatocellular Carcinoma. Hepatology, 2019, 69, 143-159.	7.3	105
10	Landscape of infiltrating B cells and their clinical significance in human hepatocellular carcinoma. OncoImmunology, 2019, 8, e1571388.	4.6	96
11	Evolutionary Dynamics of the Interferon-Induced Transmembrane Gene Family in Vertebrates. PLoS ONE, 2012, 7, e49265.	2.5	71
12	Spatial and temporal clonal evolution of intrahepatic cholangiocarcinoma. Journal of Hepatology, 2018, 69, 89-98.	3.7	63
13	Global analysis of tRNA and translation factor expression reveals a dynamic landscape of translational regulation in human cancers. Communications Biology, 2018, 1, 234.	4.4	58
14	Association Between Sex and Immune-Related Adverse Events During Immune Checkpoint Inhibitor Therapy. Journal of the National Cancer Institute, 2021, 113, 1396-1404.	6.3	56
15	The lncRNA H19 alleviates muscular dystrophy by stabilizing dystrophin. Nature Cell Biology, 2020, 22, 1332-1345.	10.3	51
16	Clinical significance of PD-1/PD-Ls gene amplification and overexpression in patients with hepatocellular carcinoma. Theranostics, 2018, 8, 5690-5702.	10.0	45
17	Pancan-meQTL: a database to systematically evaluate the effects of genetic variants on methylation in human cancer. Nucleic Acids Research, 2019, 47, D1066-D1072.	14.5	45
18	A noncoding RNA modulator potentiates phenylalanine metabolism in mice. Science, 2021, 373, 662-673.	12.6	42

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19	APAAtlas: decoding alternative polyadenylation across human tissues. <i>Nucleic Acids Research</i> , 2020, 48, D34-D39.	14.5	41
20	A Multi-Omics Perspective of Quantitative Trait Loci in Precision Medicine. <i>Trends in Genetics</i> , 2020, 36, 318-336.	6.7	41
21	Association of antibiotic treatment with immune-related adverse events in patients with cancer receiving immunotherapy. , 2022, 10, e003779.		34
22	Small non-coding RNAs in human cancer: function, clinical utility, and characterization. <i>Oncogene</i> , 2021, 40, 1570-1577.	5.9	33
23	Maximizing the Utility of Cancer Transcriptomic Data. <i>Trends in Cancer</i> , 2018, 4, 823-837.	7.4	32
24	Characterization of the dual functional effects of heat shock proteins (HSPs) in cancer hallmarks to aid development of HSP inhibitors. <i>Genome Medicine</i> , 2020, 12, 101.	8.2	31
25	Isoflurane-Induced Burst Suppression Increases Intrinsic Functional Connectivity of the Monkey Brain. <i>Frontiers in Neuroscience</i> , 2019, 13, 296.	2.8	29
26	HeRA: an atlas of enhancer RNAs across human tissues. <i>Nucleic Acids Research</i> , 2021, 49, D932-D938.	14.5	27
27	Efficacy and Safety of Transcatheter Arterial Chemoembolization and Transcatheter Arterial Chemotherapy Infusion in Hepatocellular Carcinoma: A Systematic Review and Meta-Analysis. <i>Oncology Research</i> , 2018, 26, 231-239.	1.5	20
28	tRic: a user-friendly data portal to explore the expression landscape of tRNAs in human cancers. <i>RNA Biology</i> , 2020, 17, 1674-1679.	3.1	18
29	The genetic and pharmacogenomic landscape of snoRNAs in human cancer. <i>Molecular Cancer</i> , 2020, 19, 108.	19.2	17
30	Brain Map of Intrinsic Functional Flexibility in Anesthetized Monkeys and Awake Humans. <i>Frontiers in Neuroscience</i> , 2019, 13, 174.	2.8	15
31	Animal-eRNAdb: a comprehensive animal enhancer RNA database. <i>Nucleic Acids Research</i> , 2022, 50, D46-D53.	14.5	14
32	Profiling of immune features to predict immunotherapy efficacy. <i>Innovation(China)</i> , 2021, 3, 100194.	9.1	13
33	Genetic, Pharmacogenomic, and Immune Landscapes of Enhancer RNAs Across Human Cancers. <i>Cancer Research</i> , 2022, 82, 785-790.	0.9	11
34	Molecular Treasures of Cancer Cell Lines. <i>Trends in Molecular Medicine</i> , 2019, 25, 657-659.	6.7	9
35	Functional significance of gain-of-function H19 lncRNA in skeletal muscle differentiation and anti-obesity effects. <i>Genome Medicine</i> , 2021, 13, 137.	8.2	8
36	3D Spheroids Propel Tumor Characterization. <i>Trends in Cancer</i> , 2020, 6, 622-624.	7.4	6

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37	Circular RNAs sequenced at last. Nature Biotechnology, 2021, 39, 811-812.	17.5	5
38	PWWP2B Fine-tunes Adipose Thermogenesis by Stabilizing HDACs in a NuRD Subcomplex. Advanced Science, 2021, 8, e2102060.	11.2	5
39	Identification of a STIM1 Splicing Variant that Promotes Glioblastoma Growth. Advanced Science, 2022, 9, e2103940.	11.2	5
40	Impact of Enhanced Recovery After Surgery on Long-Term Outcomes and Postoperative Recovery in Patients Undergoing Hepatectomy: A Retrospective Cohort Study. Cancer Management and Research, 2021, Volume 13, 2681-2690.	1.9	4
41	Resolving Spliceosomal Malfunctions Advances RNA-Based Therapeutics. Trends in Molecular Medicine, 2020, 26, 135-137.	6.7	1