

Jian Zheng

List of Publications by Year in descending order

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Version: 2024-02-01

36
papers

3,016
citations

279487

23
h-index

344852

36
g-index

37
all docs

37
docs citations

37
times ranked

4299
citing authors

#	ARTICLE	IF	CITATIONS
1	Circular RNA ITCH has inhibitory effect on ESCC by suppressing the Wnt/ β -catenin pathway. <i>Oncotarget</i> , 2015, 6, 6001-6013.	0.8	626
2	METTL3 facilitates tumor progression via an m6A-IGF2BP2-dependent mechanism in colorectal carcinoma. <i>Molecular Cancer</i> , 2019, 18, 112.	7.9	515
3	Excessive miR-25-3p maturation via N6-methyladenosine stimulated by cigarette smoke promotes pancreatic cancer progression. <i>Nature Communications</i> , 2019, 10, 1858.	5.8	242
4	Pancreatic cancer risk variant in LINC00673 creates a miR-1231 binding site and interferes with PTPN11 degradation. <i>Nature Genetics</i> , 2016, 48, 747-757.	9.4	237
5	Increased Levels of the Long Intergenic Non-coding Protein Coding RNA POU3F3 Promote DNA Methylation in Esophageal Squamous Cell Carcinoma Cells. <i>Gastroenterology</i> , 2014, 146, 1714-1726.e5.	0.6	169
6	PIWI-interacting RNA-36712 restrains breast cancer progression and chemoresistance by interaction with SEPW1 pseudogene SEPW1P RNA. <i>Molecular Cancer</i> , 2019, 18, 9.	7.9	139
7	PIWI-interacting RNA-54265 is oncogenic and a potential therapeutic target in colorectal adenocarcinoma. <i>Theranostics</i> , 2018, 8, 5213-5230.	4.6	115
8	Long Noncoding RNA p53 β -Stabilizing and Activating RNA Promotes p53 Signaling by Inhibiting Heterogeneous Nuclear Ribonucleoprotein K deSUMOylation and Suppresses Hepatocellular Carcinoma. <i>Hepatology</i> , 2020, 71, 112-129.	3.6	104
9	A Sequence Polymorphism in <i>miR-608</i> Predicts Recurrence after Radiotherapy for Nasopharyngeal Carcinoma. <i>Cancer Research</i> , 2013, 73, 5151-5162.	0.4	64
10	N(6)-methyladenosine-binding protein YTHDF1 suppresses EBV replication and promotes EBV RNA decay. <i>EMBO Reports</i> , 2021, 22, e50128.	2.0	59
11	NSUN2-mediated RNA 5-methylcytosine promotes esophageal squamous cell carcinoma progression via LIN28B-dependent GRB2 mRNA stabilization. <i>Oncogene</i> , 2021, 40, 5814-5828.	2.6	59
12	Serum piRNA-54265 is a New Biomarker for early detection and clinical surveillance of Human Colorectal Cancer. <i>Theranostics</i> , 2020, 10, 8468-8478.	4.6	58
13	Functional genetic variations in the IL-23 receptor gene are associated with risk of breast, lung and nasopharyngeal cancer in Chinese populations. <i>Carcinogenesis</i> , 2012, 33, 2409-2416.	1.3	55
14	Functional <i>NBS1</i> polymorphism is associated with occurrence and advanced disease status of nasopharyngeal carcinoma. <i>Molecular Carcinogenesis</i> , 2011, 50, 689-696.	1.3	48
15	<i>N6</i> -methyladenosine-Mediated Upregulation of WTAPP1 Promotes WTAP Translation and Wnt Signaling to Facilitate Pancreatic Cancer Progression. <i>Cancer Research</i> , 2021, 81, 5268-5283.	0.4	46
16	Solute Carrier Family 39 Member 6 Gene Promotes Aggressiveness of Esophageal Carcinoma Cells by Increasing Intracellular Levels of Zinc, Activating Phosphatidylinositol 3-Kinase Signaling, and Up-regulating Genes That Regulate Metastasis. <i>Gastroenterology</i> , 2017, 152, 1985-1997.e12.	0.6	40
17	Identification of chimeric TSNAX-DISC1 resulting from intergenic splicing in endometrial carcinoma through high-throughput RNA sequencing. <i>Carcinogenesis</i> , 2014, 35, 2687-2697.	1.3	36
18	A Polymorphism rs12325489C>T in the lincRNA-ENST00000515084 Exon Was Found to Modulate Breast Cancer Risk via GWAS-Based Association Analyses. <i>PLoS ONE</i> , 2014, 9, e98251.	1.1	36

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19	Inflammatory cytokine-regulated tRNA-derived fragment tRF-21 suppresses pancreatic ductal adenocarcinoma progression. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	36
20	LINC00842 inactivates transcription co-regulator PGC-1 β to promote pancreatic cancer malignancy through metabolic remodelling. <i>Nature Communications</i> , 2021, 12, 3830.	5.8	34
21	The protective role of polymorphism <i>MKK4</i> 1304 T>G in nasopharyngeal carcinoma is modulated by Epstein-Barr virus' infection status. <i>International Journal of Cancer</i> , 2012, 130, 1981-1990.	2.3	32
22	Association between the Cytotoxic T-Lymphocyte Antigen 4 +49G > A polymorphism and cancer risk: a meta-analysis. <i>BMC Cancer</i> , 2010, 10, 522.	1.1	30
23	Genome landscapes of rectal cancer before and after preoperative chemoradiotherapy. <i>Theranostics</i> , 2019, 9, 6856-6866.	4.6	27
24	Heterozygous Genetic Variations of <i>FOXP3</i> in Xp11.23 Elevate Breast Cancer Risk in Chinese Population via Skewed X-Chromosome Inactivation. <i>Human Mutation</i> , 2013, 34, n/a-n/a.	1.1	26
25	BRCA1-Associated Protein Increases Invasiveness of Esophageal Squamous Cell Carcinoma. <i>Gastroenterology</i> , 2017, 153, 1304-1319.e5.	0.6	23
26	Dysregulation, functional implications, and prognostic ability of the circadian clock across cancers. <i>Cancer Medicine</i> , 2019, 8, 1710-1720.	1.3	23
27	pCysMod: Prediction of Multiple Cysteine Modifications Based on Deep Learning Framework. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 617366.	1.8	21
28	Functional role of BTB and CNC Homology 1 gene in pancreatic cancer and its association with survival in patients treated with gemcitabine. <i>Theranostics</i> , 2018, 8, 3366-3379.	4.6	19
29	ATXN2-mediated translation of TNFR1 promotes esophageal squamous cell carcinoma via m6A-dependent manner. <i>Molecular Therapy</i> , 2022, 30, 1089-1103.	3.7	17
30	LncPipe: A Nextflow-based pipeline for identification and analysis of long non-coding RNAs from RNA-Seq data. <i>Journal of Genetics and Genomics</i> , 2018, 45, 399-401.	1.7	15
31	A micropeptide XBP1SBM encoded by lncRNA promotes angiogenesis and metastasis of TNBC via XBP1s pathway. <i>Oncogene</i> , 2022, 41, 2163-2172.	2.6	15
32	Integrative analysis of gene expression profiles reveals specific signaling pathways associated with pancreatic duct adenocarcinoma. <i>Cancer Communications</i> , 2018, 38, 1-12.	3.7	14
33	PIWI-interacting RNAs in human cancer. <i>Seminars in Cancer Biology</i> , 2021, 75, 15-28.	4.3	12
34	CircVPS13C promotes pituitary adenoma growth by decreasing the stability of IFITM1 mRNA via interacting with RRBP1. <i>Oncogene</i> , 2022, 41, 1550-1562.	2.6	12
35	Genome-wide identification and characterization of circular RNA m6A modification in pancreatic cancer. <i>Genome Medicine</i> , 2021, 13, 183.	3.6	10
36	Clinical and genomic characterization of neutral tumor evolution in Head and Neck Squamous Cell Carcinoma. <i>Genomics</i> , 2020, 112, 3448-3454.	1.3	2