Xiao-Dong Zhao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3655331/publications.pdf

Version: 2024-02-01

46 papers

3,290 citations

361296 20 h-index 42 g-index

46 all docs

46 docs citations

46 times ranked

5494 citing authors

#	Article	IF	CITATIONS
1	The Oct4 and Nanog transcription network regulates pluripotency in mouse embryonic stem cells. Nature Genetics, 2006, 38, 431-440.	9.4	2,162
2	MALAT1 long ncRNA promotes gastric cancer metastasis by suppressing <i>PCDH10</i> . Oncotarget, 2016, 7, 12693-12703.	0.8	97
3	NUDT21 negatively regulates PSMB2 and CXXC5 by alternative polyadenylation and contributes to hepatocellular carcinoma suppression. Oncogene, 2018, 37, 4887-4900.	2.6	83
4	A semi-supervised deep learning method based on stacked sparse auto-encoder for cancer prediction using RNA-seq data. Computer Methods and Programs in Biomedicine, 2018, 166, 99-105.	2.6	70
5	Characterization and Potential Antitumor Activity of Polysaccharide from Gracilariopsis lemaneiformis. Marine Drugs, 2017, 15, 100.	2.2	64
6	Integrative epigenomic analysis reveals unique epigenetic signatures involved in unipotency of mouse female germline stem cells. Genome Biology, 2016, 17, 162.	3.8	61
7	Characteristic and human exposure risk assessment of per- and polyfluoroalkyl substances: A study based on indoor dust and drinking water in China. Environmental Pollution, 2019, 254, 112873.	3.7	54
8	<i>Helicobacter pylori</i> CagA induces tumor suppressor gene hypermethylation by upregulating DNMT1 via AKT-NFÎB pathway in gastric cancer development. Oncotarget, 2016, 7, 9788-9800.	0.8	53
9	Genome-Wide Plasma Cell-Free DNA Methylation Profiling Identifies Potential Biomarkers for Lung Cancer. Disease Markers, 2019, 2019, 1-7.	0.6	44
10	Genome-wide profiling of polyadenylation sites reveals a link between selective polyadenylation and cancer metastasis. Human Molecular Genetics, 2015, 24, 3410-3417.	1.4	41
11	Activation of AIFM2 enhances apoptosis of human lung cancer cells undergoing toxicological stress. Toxicology Letters, 2016, 258, 227-236.	0.4	34
12	Post-transcriptional regulation of ERBB2 by miR26a/b and HuR confers resistance to tamoxifen in estrogen receptor-positive breast cancer cells. Journal of Biological Chemistry, 2017, 292, 13551-13564.	1.6	34
13	Identification of Biomarkers for Predicting Lymph Node Metastasis of Stomach Cancer Using Clinical DNA Methylation Data. Disease Markers, 2017, 2017, 1-7.	0.6	31
14	Antibody Binding Epitope Mapping (AbMap) of Hundred Antibodies in a Single Run. Molecular and Cellular Proteomics, 2021, 20, 100059.	2.5	30
15	CPSF6 links alternative polyadenylation to metabolism adaption in hepatocellular carcinoma progression. Journal of Experimental and Clinical Cancer Research, 2021, 40, 85.	3 . 5	29
16	Transcriptome Profiling Reveals the Antitumor Mechanism of Polysaccharide from Marine Algae Gracilariopsis lemaneiformis. PLoS ONE, 2016, 11, e0158279.	1.1	29
17	Organic UV filters exposure induces the production of inflammatory cytokines in human macrophages. Science of the Total Environment, 2018, 635, 926-935.	3.9	27
18	Large scale gene regulatory network inference with a multi-level strategy. Molecular BioSystems, 2016, 12, 588-597.	2.9	26

#	Article	IF	Citations
19	Genomic analyses of unique carbohydrate and phytohormone metabolism in the macroalga Gracilariopsis lemaneiformis (Rhodophyta). BMC Plant Biology, 2018, 18, 94.	1.6	25
20	Temporal requirements for ISL1 in sympathetic neuron proliferation, differentiation, and diversification. Cell Death and Disease, 2018, 9, 247.	2.7	23
21	An array of 60,000 antibodies for proteome-scale antibody generation and target discovery. Science Advances, 2020, 6, eaax2271.	4.7	22
22	The histone chaperone complex FACT promotes proliferative switch of G 0 cancer cells. International Journal of Cancer, 2019, 145, 164-178.	2.3	20
23	Epigenetic Profiling of H3K4Me3 Reveals Herbal Medicine Jinfukang-Induced Epigenetic Alteration Is Involved in Anti-Lung Cancer Activity. Evidence-based Complementary and Alternative Medicine, 2016, 2016, 1-13.	0.5	16
24	Herbal formula YYJD inhibits tumor growth by inducing cell cycle arrest and senescence in lung cancer. Scientific Reports, 2017, 7, 4984.	1.6	16
25	PLZF suppresses differentiation of mouse spermatogonial progenitor cells via binding of differentiation associated genes. Journal of Cellular Physiology, 2020, 235, 3033-3042.	2.0	16
26	Breast Cancer Diagnosis Using an Unsupervised Feature Extraction Algorithm Based on Deep Learning. , 2018, , .		15
27	Jinfukang induces cellular apoptosis through activation of Fas and DR4 in A549 cells. Oncology Letters, 2018, 16, 4343-4352.	0.8	15
28	Characterization of Genome-Wide DNA Methylation and Hydroxymethylation in Mouse Arcuate Nucleus of Hypothalamus During Puberty Process. Frontiers in Genetics, 2020, 11, 626536.	1.1	15
29	Ultra-deep sequencing of ribosome-associated poly-adenylated RNA in early <i>Drosophila</i> embryos reveals hundreds of conserved translated sORFs. DNA Research, 2016, 23, 571-580.	1.5	14
30	An expression based REST signature predicts patient survival and therapeutic response for glioblastoma multiforme. Scientific Reports, 2016, 6, 34556.	1.6	14
31	Busulfan Suppresses Autophagy in Mouse Spermatogonial Progenitor Cells via mTOR of AKT and p53 Signaling Pathways. Stem Cell Reviews and Reports, 2020, 16, 1242-1255.	1.7	14
32	Systematic profiling of SARS-CoV-2-specific IgG epitopes at amino acid resolution. Cellular and Molecular Immunology, 2021, 18, 1067-1069.	4.8	14
33	Identification of Serum Biomarkers for Systemic Lupus Erythematosus Using a Library of Phage Displayed Random Peptides and Deep Sequencing. Molecular and Cellular Proteomics, 2019, 18, 1851-1863.	2.5	13
34	Herbal formula Yangyinjiedu induces lung cancer cell apoptosis via activation of early growth response 1. Journal of Cellular and Molecular Medicine, 2019, 23, 6193-6202.	1.6	12
35	CPF impedes cell cycle reâ€entry of quiescent lung cancer cells through transcriptional suppression of FACT and câ€MYC. Journal of Cellular and Molecular Medicine, 2020, 24, 2229-2239.	1.6	11
36	STAT3 is required for proliferation and exhibits a cell type-specific binding preference in mouse female germline stem cells. Molecular Omics, 2018, 14, 95-102.	1.4	9

3

#	Article	IF	CITATIONS
37	Genome-wide analysis of cell-Free DNA methylation profiling with MeDIP-seq identified potential biomarkers for colorectal cancer. World Journal of Surgical Oncology, 2022, 20, 21.	0.8	9
38	Genome-Wide Profiling Reveals That Herbal Medicine Jinfukang-Induced Polyadenylation Alteration Is Involved in Anti-Lung Cancer Activity. Evidence-based Complementary and Alternative Medicine, 2017, 2017, 1-8.	0.5	8
39	CAsubtype: An R Package to Identify Gene Sets Predictive of Cancer Subtypes and Clinical Outcomes. Interdisciplinary Sciences, Computational Life Sciences, 2018, 10, 169-175.	2.2	5
40	The binding epitope of sintilimab on PD-1 revealed by AbMap. Acta Biochimica Et Biophysica Sinica, 2021, 53, 628-635.	0.9	5
41	Analysis of chromatin accessibility in <i>p53</i> deficient spermatogonial stem cells for high frequency transformation into pluripotent state. Cell Proliferation, 2022, 55, e13195.	2.4	5
42	Integrative RNA-Seq and ATAC-Seq Analysis Reveals the Migration-Associated Genes Involved in Antitumor Effects of Herbal Medicine Feiyanning on Lung Cancer Cells. Frontiers in Genetics, 2021, 12, 799099.	1.1	4
43	Unexpected roles of long non-coding RNAs in cancer biology. Journal of Shanghai Jiaotong University (Science), 2014, 19, 544-549.	0.5	1
44	Enhancing the effectiveness of fungicides by optimizing their combinations. , 2014, , .		0
45	165 ldentification of serum biomarkers for systemic lupus erythematosus using a library of phage displayed random peptides and deep sequencing. , 2019, , .		0
46	Transcriptome-wide alternative polyadenylation profiling reveals the herbal formula Yangyinjiedu-induced preferential poly(A) usage in lung adenocarcinoma cells. Acta Biochimica Et Biophysica Sinica, 2020, 52, 1171-1174.	0.9	0