

Qian Yang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3547155/publications.pdf>

Version: 2024-02-01

17
papers

508
citations

759233

12
h-index

839539

18
g-index

20
all docs

20
docs citations

20
times ranked

577
citing authors

#	ARTICLE	IF	CITATIONS
1	Fatty Acid Signaling Impacts Prostate Cancer Lineage Plasticity in an Autocrine and Paracrine Manner. <i>Cancers</i> , 2022, 14, 3449.	3.7	2
2	A Transcriptional Regulatory Loop of Master Regulator Transcription Factors, PPARC, and Fatty Acid Synthesis Promotes Esophageal Adenocarcinoma. <i>Cancer Research</i> , 2021, 81, 1216-1229.	0.9	41
3	Potential Mechanism of Immune Evasion Associated with the Master Regulator ASCL2 in Microsatellite Stability in Colorectal Cancer. <i>Journal of Immunology Research</i> , 2021, 2021, 1-12.	2.2	9
4	A pan-cancer analysis of CpG Island gene regulation reveals extensive plasticity within Polycomb target genes. <i>Nature Communications</i> , 2021, 12, 2485.	12.8	21
5	Interplay and cooperation between SREBF1 and master transcription factors regulate lipid metabolism and tumor-promoting pathways in squamous cancer. <i>Nature Communications</i> , 2021, 12, 4362.	12.8	50
6	Activation of bivalent factor DLX5 cooperates with master regulator TP63 to promote squamous cell carcinoma. <i>Nucleic Acids Research</i> , 2021, 49, 9246-9263.	14.5	13
7	Integrated single-cell transcriptome analysis reveals heterogeneity of esophageal squamous cell carcinoma microenvironment. <i>Nature Communications</i> , 2021, 12, 7335.	12.8	69
8	Master transcription factors form interconnected circuitry and orchestrate transcriptional networks in oesophageal adenocarcinoma. <i>Gut</i> , 2020, 69, 630-640.	12.1	68
9	Lineage-Specific Epigenomic and Genomic Activation of Oncogene HNF4A Promotes Gastrointestinal Adenocarcinomas. <i>Cancer Research</i> , 2020, 80, 2722-2736.	0.9	37
10	Identification of key genes by integrating DNA methylation and next-generation transcriptome sequencing for esophageal squamous cell carcinoma. <i>Aging</i> , 2020, 12, 1332-1365.	3.1	17
11	D-Inc: a comprehensive database and analytical platform to dissect the modification of drugs on lncRNA expression. <i>RNA Biology</i> , 2019, 16, 1586-1591.	3.1	25
12	Long Read Single-Molecule Real-Time Sequencing Elucidates Transcriptome-Wide Heterogeneity and Complexity in Esophageal Squamous Cells. <i>Frontiers in Genetics</i> , 2019, 10, 915.	2.3	12
13	Pathway enrichment analysis approach based on topological structure and updated annotation of pathway. <i>Briefings in Bioinformatics</i> , 2019, 20, 168-177.	6.5	32
14	Systematical analysis of lncRNA-mRNA competing endogenous RNA network in breast cancer subtypes. <i>Breast Cancer Research and Treatment</i> , 2018, 169, 267-275.	2.5	47
15	Accurate prediction and elucidation of drug resistance based on the robust and reproducible chemoresponse communities. <i>International Journal of Cancer</i> , 2018, 142, 1427-1439.	5.1	3
16	The histone deacetylase inhibitor panobinostat exerts anticancer effects on esophageal squamous cell carcinoma cells by inducing cell cycle arrest. <i>Cell Biochemistry and Function</i> , 2018, 36, 398-407.	2.9	16
17	The gain and loss of long noncoding RNA associated-competing endogenous RNAs in prostate cancer. <i>Oncotarget</i> , 2016, 7, 57228-57238.	1.8	43