Haiming Liu

List of Publications by Year
in descending order

20 papers	694 citations	623734 14 h-index	794594 19 g-index
20 all docs	20 docs citations	20 times ranked	886 citing authors

#	Article	IF	Citations
1	LINC00941 promotes CRC metastasis through preventing SMAD4 protein degradation and activating the TGF-β/SMAD2/3 signaling pathway. Cell Death and Differentiation, 2021, 28, 219-232.	11.2	99
2	DMfold: A Novel Method to Predict RNA Secondary Structure With Pseudoknots Based on Deep Learning and Improved Base Pair Maximization Principle. Frontiers in Genetics, 2019, 10, 143.	2.3	63
3	Integrative Analysis of Dysregulated lncRNA-Associated ceRNA Network Reveals Functional lncRNAs in Gastric Cancer. Genes, 2018, 9, 303.	2.4	60
4	Role of plant MicroRNA in cross-species regulatory networks of humans. BMC Systems Biology, 2016, 10, 60.	3.0	53
5	miR-218 inhibited tumor angiogenesis by targeting ROBO1 in gastric cancer. Gene, 2017, 615, 42-49.	2.2	52
6	Long non‐coding RNA MYOSLID functions as a competing endogenous RNA to regulate MCL‐1 expression by sponging miR‐29c‐3p in gastric cancer. Cell Proliferation, 2019, 52, e12678.	5.3	51
7	Identification of Potential Prognostic Genes for Neuroblastoma. Frontiers in Genetics, 2018, 9, 589.	2.3	47
8	Long Non-coding RNA LINC00941 as a Potential Biomarker Promotes the Proliferation and Metastasis of Gastric Cancer. Frontiers in Genetics, 2019, 10, 5.	2.3	47
9	miR‐204‐5p suppresses hepatocellular cancer proliferation by regulating homeoprotein SIX1 expression. FEBS Open Bio, 2018, 8, 189-200.	2.3	40
10	O-GlcNAcylation of SIX1 enhances its stability and promotes Hepatocellular Carcinoma Proliferation. Theranostics, 2020, 10, 9830-9842.	10.0	33
11	miR-5590-3p inhibited tumor growth in gastric cancer by targeting DDX5/AKT/m-TOR pathway. Biochemical and Biophysical Research Communications, 2018, 503, 1491-1497.	2.1	32
12	O‐GlcNAcylation promotes colorectal cancer progression by regulating protein stability and potential catcinogenic function of DDX5. Journal of Cellular and Molecular Medicine, 2019, 23, 1354-1362.	3.6	31
13	Aberrantly high activation of a FoxM1–STMN1 axis contributes to progression and tumorigenesis in FoxM1-driven cancers. Signal Transduction and Targeted Therapy, 2021, 6, 42.	17.1	28
14	Cellular components in tumor microenvironment of neuroblastoma and the prognostic value. PeerJ, 2019, 7, e8017.	2.0	18
15	Comparative pharmacoproteomics reveals potential targets for berberine, a promising therapy for colorectal cancer. Biochemical and Biophysical Research Communications, 2020, 525, 244-250.	2.1	13
16	The FENDRR/FOXC2 Axis Contributes to Multidrug Resistance in Gastric Cancer and Correlates With Poor Prognosis. Frontiers in Oncology, 2021, 11, 634579.	2.8	11
17	ncRFP: A Novel end-to-end Method for Non-Coding RNAs Family Prediction Based on Deep Learning. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2021, 18, 784-789.	3.0	9
18	A Dynamic Service Placement Based on Deep Reinforcement Learning in Mobile Edge Computing. Network, 2022, 2, 106-122.	2.4	6

#	Article	IF	Citations
19	Identification of novel Phytophthora infestans small RNAs involved in potato late blight reveals potential cross-kingdom regulation to facilitate oomycete infection. International Journal of Data Mining and Bioinformatics, 2020, 23, 119.	0.1	1
20	Resource provisioning in collaborative fog computing for multiple delay€sensitive users. Software - Practice and Experience, 2023, 53, 243-262.	3.6	0