## Hui-Yun Wang

List of Publications by Year in descending order

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63 papers 2,849 citations

30 h-index 51 g-index

65 all docs

65
docs citations

65 times ranked 5222 citing authors

#	Article	IF	CITATIONS
1	Prognostic value of a microRNA signature in nasopharyngeal carcinoma: a microRNA expression analysis. Lancet Oncology, The, 2012, 13, 633-641.	5.1	274
2	Rab1A Is an mTORC1 Activator and a Colorectal Oncogene. Cancer Cell, 2014, 26, 754-769.	7.7	218
3	PKD1 Phosphorylation-Dependent Degradation of SNAIL by SCF-FBXO11 Regulates Epithelial-Mesenchymal Transition and Metastasis. Cancer Cell, 2014, 26, 358-373.	7.7	196
4	Expanding roles of superoxide dismutases in cell regulation and cancer. Drug Discovery Today, 2016, 21, 143-149.	3.2	180
5	Downregulation of Six MicroRNAs Is Associated with Advanced Stage, Lymph Node Metastasis and Poor Prognosis in Small Cell Carcinoma of the Cervix. PLoS ONE, 2012, 7, e33762.	1.1	108
6	Clinical Significance and Prognostic Value of microRNA Expression Signatures in Hepatocellular Carcinoma. Clinical Cancer Research, 2013, 19, 4780-4791.	3.2	95
7	SOD1 Phosphorylation by mTORC1 Couples Nutrient Sensing and Redox Regulation. Molecular Cell, 2018, 70, 502-515.e8.	4.5	94
8	miR-665 expression predicts poor survival and promotes tumor metastasis by targeting NR4A3 in breast cancer. Cell Death and Disease, 2019, 10, 479.	2.7	81
9	Significance and mechanism of androgen receptor overexpression and androgen receptor/mechanistic target of rapamycin crossâ€talk in hepatocellular carcinoma. Hepatology, 2018, 67, 2271-2286.	3.6	78
10	Identification of a geneâ€expression signature for predicting lymph node metastasis in patients with early stage cervical carcinoma. Cancer, 2011, 117, 3363-3373.	2.0	77
11	Emerging Role of MicroRNAs in mTOR Signaling. Cellular and Molecular Life Sciences, 2017, 74, 2613-2625.	2.4	74
12	Checkpoint kinase 1 is negatively regulated by miR-497 in hepatocellular carcinoma. Medical Oncology, 2014, 31, 844.	1.2	66
13	Overexpressed HDAC4 is associated with poor survival and promotes tumor progression in esophageal carcinoma. Aging, 2016, 8, 1236-1248.	1.4	66
14	A genotyping system capable of simultaneously analyzing > 1000 single nucleotide polymorphisms in a haploid genome. Genome Research, 2005, 15, 276-283.	2.4	63
15	Identification of ADH4 as a novel and potential prognostic marker in hepatocellular carcinoma. Medical Oncology, 2012, 29, 2737-2743.	1.2	61
16	MAF1 suppresses AKTâ€mTOR signaling and liver cancer through activation of PTEN transcription. Hepatology, 2016, 63, 1928-1942.	3.6	61
17	Aberrant amino acid signaling promotes growth and metastasis of hepatocellular carcinomas through Rab1A-dependent activation of mTORC1 by Rab1A. Oncotarget, 2015, 6, 20813-20828.	0.8	61
18	RBM24 suppresses cancer progression by upregulating miR-25 to target MALAT1 in nasopharyngeal carcinoma. Cell Death and Disease, 2016, 7, e2352-e2352.	2.7	58

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19	Upregulated TRIM29 promotes proliferation and metastasis of nasopharyngeal carcinoma via PTEN/AKT/mTOR signal pathway. Oncotarget, 2016, 7, 13634-13650.	0.8	57
20	TERT Polymorphism rs2736100-C Is Associated with EGFR Mutation–Positive Non–Small Cell Lung Cancer. Clinical Cancer Research, 2015, 21, 5173-5180.	3.2	47
21	Serine and one-carbon metabolism, a bridge that links mTOR signaling and DNA methylation in cancer. Pharmacological Research, 2019, 149, 104352.	3.1	45
22	Identification of a 7-gene signature that predicts relapse and survival for early stage patients with cervical carcinoma. Medical Oncology, 2012, 29, 2911-2918.	1.2	42
23	Low serum level of miR-485-3p predicts poor survival in patients with glioblastoma. PLoS ONE, 2017, 12, e0184969.	1.1	40
24	TRIM29 overexpression is associated with poor prognosis and promotes tumor progression by activating Wnt $\hat{l}^2$ -catenin pathway in cervical cancer. Oncotarget, 2016, 7, 28579-28591.	0.8	39
25	Prognostic and predictive value of a microRNA signature in adults with T-cell lymphoblastic lymphoma. Leukemia, 2019, 33, 2454-2465.	3.3	38
26	Reduced SOD2 expression is associated with mortality of hepatocellular carcinoma patients in a mutant p53-dependent manner. Aging, 2016, 8, 1184-1200.	1.4	34
27	Decreased expression of ALDH1L1 is associated with a poor prognosis in hepatocellular carcinoma. Medical Oncology, 2012, 29, 1843-1849.	1.2	32
28	Identification of a novel microRNA signature associated with intrahepatic cholangiocarcinoma (ICC) patient prognosis. BMC Cancer, 2015, 15, 64.	1.1	32
29	Overexpression of Rab1B and MMP9 predicts poor survival and good response to chemotherapy in patients with colorectal cancer. Aging, 2017, 9, 914-931.	1.4	32
30	Beyond regulation of pol III: Role of MAF1 in growth, metabolism, aging and cancer. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2018, 1861, 338-343.	0.9	31
31	Generation of Urine Cell-Derived Non-integrative Human iPSCs and iNSCs: A Step-by-Step Optimized Protocol. Frontiers in Molecular Neuroscience, 2017, 10, 348.	1.4	28
32	Efficacy and safety of nimotuzumab in addition to radiotherapy and temozolomide for cerebral glioblastoma: a phase II multicenter clinical trial. Journal of Cancer, 2019, 10, 3214-3223.	1.2	28
33	Identification of two microRNA signatures in whole blood as novel biomarkers for diagnosis of nasopharyngeal carcinoma. Journal of Translational Medicine, 2019, 17, 186.	1.8	27
34	EIF4EBP1 Overexpression Is Associated with Poor Survival and Disease Progression in Patients with Hepatocellular Carcinoma. PLoS ONE, 2015, 10, e0117493.	1.1	27
35	LncRNA CSMD1-1 promotes the progression of Hepatocellular Carcinoma by activating MYC signaling. Theranostics, 2020, 10, 7527-7544.	4.6	26
36	Identification of microRNA-615-3p as a novel tumor suppressor in non-small cell lung cancer. Oncology Letters, 2017, 13, 2403-2410.	0.8	23

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37	MicroRNA-34c-3p promotes cell proliferation and invasion in hepatocellular carcinoma by regulation of NCKAP1 expression. Journal of Cancer Research and Clinical Oncology, 2017, 143, 263-273.	1.2	21
38	Synergistic Inhibitory Effect of Hyperbaric Oxygen Combined with Sorafenib on Hepatoma Cells. PLoS ONE, 2014, 9, e100814.	1.1	19
39	MicroRNA-711 is a prognostic factor for poor overall survival and has an oncogenic role in breast cancer. Oncology Letters, 2016, 11, 2155-2163.	0.8	18
40	Sorafenib and Carfilzomib Synergistically Inhibit the Proliferation, Survival, and Metastasis of Hepatocellular Carcinoma. Molecular Cancer Therapeutics, 2018, 17, 2610-2621.	1.9	18
41	p53R2 overexpression in cervical cancer promotes AKT signaling and EMT, and is correlated with tumor progression, metastasis and poor prognosis. Cell Cycle, 2017, 16, 1673-1682.	1.3	17
42	Identification of an 88-microRNA signature in whole blood for diagnosis of hepatocellular carcinoma and other chronic liver diseases. Aging, 2017, 9, 1565-1584.	1.4	17
43	Expression of Rac-1 related to tumor depth, lymph node metastasis and patient prognosis in esophageal squamous cell carcinoma. Medical Oncology, 2013, 30, 689.	1.2	15
44	MT1G is Silenced by DNA Methylation and Contributes to the Pathogenesis of Hepatocellular Carcinoma. Journal of Cancer, 2018, 9, 2807-2816.	1.2	15
45	Lnc-GAN1 expression is associated with good survival and suppresses tumor progression by sponging mir-26a-5p to activate PTEN signaling in non-small cell lung cancer. Journal of Experimental and Clinical Cancer Research, 2021, 40, 9.	3.5	15
46	Allele Loss and Down-Regulation of Heparanase Gene Are Associated with the Progression and Poor Prognosis of Hepatocellular Carcinoma. PLoS ONE, 2012, 7, e44061.	1.1	13
47	Identification of immunological subtypes of hepatocellular carcinoma with expression profiling of immune-modulating genes. Aging, 2020, 12, 12187-12205.	1.4	13
48	CircSCAP interacts with SF3A3 to inhibit the malignance of non-small cell lung cancer by activating p53 signaling. Journal of Experimental and Clinical Cancer Research, 2022, 41, 120.	3.5	13
49	mTORC1 Promotes ARID1A Degradation and Oncogenic Chromatin Remodeling in Hepatocellular Carcinoma. Cancer Research, 2021, 81, 5652-5665.	0.4	12
50	Clinical significance and prognostic value of TRIM24 expression in esophageal squamous cell carcinoma. Aging, 2016, 8, 2204-2221.	1.4	11
51	Reduced expression of Dicer11 is associated with poor prognosis in patients with nasopharyngeal carcinoma. Medical Oncology, 2013, 30, 360.	1.2	10
52	Rapamycin and trametinib: a rational combination for treatment of NSCLC. International Journal of Biological Sciences, 2021, 17, 3211-3223.	2.6	10
53	Methyl-methanesulfonate sensitivity 19 expression is associated with metastasis and chemoradiotherapy response in esophageal cancer. World Journal of Gastroenterology, 2015, 21, 4240.	1.4	10
54	Phosphorylation of androgen receptor by mTORC1 promotes liver steatosis and tumorigenesis. Hepatology, 2022, 75, 1123-1138.	3 <b>.</b> 6	9

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55	Rab1 GTPases as oncogenes. Aging, 2015, 7, 897-898.	1.4	9
56	Rab1A promotes cancer metastasis and radioresistance through activating GSK-3β/Wnt/β-catenin signaling in nasopharyngeal carcinoma. Aging, 2020, 12, 20380-20395.	1.4	9
57	The prognostic value of a seven-lncRNA signature in patients with esophageal squamous cell carcinoma: a lncRNA expression analysis. Journal of Translational Medicine, 2020, 18, 47.	1.8	8
58	Androgen Receptor Promotes Gastric Carcinogenesis via Upregulating Cell Cycle-Related Kinase Expression. Journal of Cancer, 2019, 10, 4178-4188.	1.2	7
59	Identification of a 4-IncRNA signature predicting prognosis of patients with non-small cell lung cancer: a multicenter study in China. Journal of Translational Medicine, 2020, 18, 320.	1.8	7
60	Prognostic values of apoptosis-stimulating P53-binding protein 1 and 2 and their relationships with clinical characteristics of esophageal squamous cell carcinoma patients: a retrospective study. Chinese Journal of Cancer, 2017, 36, 15.	4.9	6
61	Combination Treatment With Inhibitors of ERK and Autophagy Enhances Antitumor Activity of Betulinic Acid in Non–small-Cell Lung Cancer In Vivo and In Vitro. Frontiers in Pharmacology, 2021, 12, 684243.	1.6	6
62	IDDF2018-ABS-0140â€The crosstalk of MTORC1 and DNA methylation in hepatocellular carcinoma. , 2018, , .		2
63	IDDF2018-ABS-0139 $\hat{a} \in \mathbb{R}$ Metallothionein $1G$ is silenced by DNA methylation and contributes to the pathogenesis of hepatocellular carcinoma. , 2018, , .		O