Zhi Dai

List of Publications by Year in descending order

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236833 133188 3,924 59 59 25 citations h-index g-index papers 61 61 61 6003 citing authors all docs docs citations times ranked

#	Article	IF	CITATIONS
1	Serial circulating tumor DNA to predict early recurrence in patients with hepatocellular carcinoma: a prospective study. Molecular Oncology, 2022, 16, 549-561.	2.1	21
2	Whole-genome sequencing reveals the evolutionary trajectory of HBV-related hepatocellular carcinoma early recurrence. Signal Transduction and Targeted Therapy, 2022, 7, 24.	7.1	7
3	Validating RRP12 Expression and Its Prognostic Significance in HCC Based on Data Mining and Bioinformatics Methods. Frontiers in Oncology, 2022, 12, 812009.	1.3	5
4	Targeting HNRNPM Inhibits Cancer Stemness and Enhances Antitumor Immunity in Wnt-activated Hepatocellular Carcinoma. Cellular and Molecular Gastroenterology and Hepatology, 2022, 13, 1413-1447.	2.3	15
5	Splicing factors: Insights into their regulatory network in alternative splicing in cancer. Cancer Letters, 2021, 501, 83-104.	3.2	22
6	Structures/cytotoxicity/selectivity relationship of natural steroidal saponins against GSCs and primary mechanism of tribulosaponin A. European Journal of Medicinal Chemistry, 2021, 210, 113068.	2.6	10
7	Identification and validation of a new gene signature predicting prognosis of hepatocellular carcinoma patients by network analysis of stemness indices. Expert Review of Gastroenterology and Hepatology, 2021, 15, 699-709.	1.4	9
8	Bioactivity Ingredients of <i>Chaenomeles speciosa</i> against Microbes: Characterization by LC-MS and Activity Evaluation. Journal of Agricultural and Food Chemistry, 2021, 69, 4686-4696.	2.4	20
9	Splicing factor SRSF1 promotes breast cancer progression via oncogenic splice switching of PTPMT1. Journal of Experimental and Clinical Cancer Research, 2021, 40, 171.	3.5	42
10	The expression profiles and prognostic values of HSP70s in hepatocellular carcinoma. Cancer Cell International, 2021, 21, 286.	1.8	24
11	Profiles of alternative splicing landscape in breast cancer and their clinical significance: an integrative analysis based on large-sequencing data. Annals of Translational Medicine, 2021, 9, 58-58.	0.7	3
12	Furostanol Saponins from Asparagus cochinchinensis and Their Cytotoxicity. Natural Products and Bioprospecting, 2021, 11, 651-658.	2.0	6
13	HNRNPABâ€regulated lncRNAâ€ELF209 inhibits the malignancy of hepatocellular carcinoma. International Journal of Cancer, 2020, 146, 169-180.	2.3	28
14	Plasma hsa_circ_0027089 is a diagnostic biomarker for hepatitis B virus-related hepatocellular carcinoma. Carcinogenesis, 2020, 41, 296-302.	1.3	51
15	Metadherin–PRMT5 complex enhances the metastasis of hepatocellular carcinoma through the WNT–β-catenin signaling pathway. Carcinogenesis, 2020, 41, 130-138.	1.3	27
16	TGM3 promotes epithelial–mesenchymal transition and hepatocellular carcinogenesis and predicts poor prognosis for patients after curative resection. Digestive and Liver Disease, 2020, 52, 668-676.	0.4	15
17	Genetic Alterations and Transcriptional Expression of m6A RNA Methylation Regulators Drive a Malignant Phenotype and Have Clinical Prognostic Impact in Hepatocellular Carcinoma. Frontiers in Oncology, 2020, 10, 900.	1.3	11
18	Age-adjusted Charlson Comorbidity Index predicts survival in intrahepatic cholangiocarcinoma patients after curative resection. Annals of Translational Medicine, 2020, 8, 487-487.	0.7	25

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19	Guidelines for the Diagnosis and Treatment of Hepatocellular Carcinoma (2019 Edition). Liver Cancer, 2020, 9, 682-720.	4.2	427
20	Aspartate aminotransferase-to-platelet ratio index predicts prognosis of hepatocellular carcinoma after postoperative adjuvant transarterial chemoembolization. Cancer Management and Research, 2019, Volume 11, 63-79.	0.9	12
21	The miR-561-5p/CX ₃ CL1 Signaling Axis Regulates Pulmonary Metastasis in Hepatocellular Carcinoma Involving CX ₃ CR1 ⁺ Natural Killer Cells Infiltration. Theranostics, 2019, 9, 4779-4794.	4.6	72
22	Prognostic Value and Predication Model of Microvascular Invasion in Patients with Intrahepatic Cholangiocarcinoma. Journal of Cancer, 2019, 10, 5575-5584.	1.2	28
23	Development and validation of a new tumor-based gene signature predicting prognosis of HBV/HCV-included resected hepatocellular carcinoma patients. Journal of Translational Medicine, 2019, 17, 203.	1.8	18
24	Tissue-infiltrating lymphocytes signature predicts survival in patients with early/intermediate stage hepatocellular carcinoma. BMC Medicine, 2019, 17, 106.	2.3	31
25	Prognostic alternative mRNA splicing signature in hepatocellular carcinoma: a study based on large-scale sequencing data. Carcinogenesis, 2019, 40, 1077-1085.	1.3	34
26	Systemic inflammation score predicts survival in patients with intrahepatic cholangiocarcinoma undergoing curative resection. Journal of Cancer, 2019, 10, 494-503.	1.2	36
27	Daily decrease of post-operative alpha-fetoprotein by 9% discriminates prognosis of HCC: A multicenter retrospective study. Aging, 2019, 11, 11111-11123.	1.4	6
28	A Novel Risk prediction Model for Patients with Combined Hepatocellular-Cholangiocarcinoma. Journal of Cancer, 2018, 9, 1025-1032.	1.2	14
29	Comparative efficacy and safety between ablative therapies or surgery for small hepatocellular carcinoma: a network meta-analysis. Expert Review of Gastroenterology and Hepatology, 2018, 12, 935-945.	1.4	13
30	Surgical Treatment of Combined Hepatocellular-Cholangiocarcinoma is as Effective in Elderly Patients as it is in Younger Patients: A Propensity Score Matching Analysis. Journal of Cancer, 2018, 9, 1106-1112.	1.2	16
31	Guidelines for Diagnosis and Treatment of Primary Liver Cancer in China (2017 Edition). Liver Cancer, 2018, 7, 235-260.	4.2	426
32	Novel role of semaphorin 3A in the growth and progression of hepatocellular carcinoma. Oncology Reports, 2017, 37, 3313-3320.	1.2	17
33	MicroRNA-29a induces loss of 5-hydroxymethylcytosine and promotes metastasis of hepatocellular carcinoma through a TET–SOCS1–MMP9 signaling axis. Cell Death and Disease, 2017, 8, e2906-e2906.	2.7	66
34	Serum exosomal miR-125b is a novel prognostic marker for hepatocellular carcinoma. OncoTargets and Therapy, 2017, Volume 10, 3843-3851.	1.0	117
35	Perioperative blood transfusion does not affect recurrence-free and overall survivals after curative resection for intrahepatic cholangiocarcinoma: a propensity score matching analysis. BMC Cancer, 2017, 17, 762.	1.1	12
36	Coagulopathy associated with poor prognosis in intrahepatic cholangiocarcinoma patients after curative resection. BioScience Trends, 2017, 11, 469-474.	1.1	3

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37	Prognostic value of systemic inflammation score in patients with hepatocellular carcinoma after hepatectomy. Oncotarget, 2017, 8, 79366-79375.	0.8	16
38	CCL24 contributes to HCC malignancy via RhoB- VEGFA-VEGFR2 angiogenesis pathway and indicates poor prognosis. Oncotarget, 2017, 8, 5135-5148.	0.8	35
39	Prostate-derived ETS factor improves prognosis and represses proliferation and invasion in hepatocellular carcinoma. Oncotarget, 2017, 8, 52488-52500.	0.8	10
40	Overexpression of semaphorin 3A promotes tumor progression and predicts poor prognosis in hepatocellular carcinoma after curative resection. Oncotarget, 2016, 7, 51733-51746.	0.8	34
41	miRâ€28â€5pâ€ILâ€34â€macrophage feedback loop modulates hepatocellular carcinoma metastasis. Hepatology 2016, 63, 1560-1575.	'3 . 6	166
42	Tumor-Associated Neutrophils Recruit Macrophages and T-Regulatory Cells to Promote Progression of Hepatocellular Carcinoma and Resistance to Sorafenib. Gastroenterology, 2016, 150, 1646-1658.e17.	0.6	586
43	Caveolin-1 promotes tumor growth and metastasis via autophagy inhibition in hepatocellular carcinoma. Clinics and Research in Hepatology and Gastroenterology, 2016, 40, 169-178.	0.7	32
44	Macrophage-secreted IL-8 induces epithelial-mesenchymal transition in hepatocellular carcinoma cells by activating the JAK2/STAT3/Snail pathway. International Journal of Oncology, 2015, 46, 587-596.	1.4	177
45	CXCR2/CXCL5 axis contributes to epithelial–mesenchymal transition of HCC cells through activating PI3K/Akt/GSK-3β/Snail signaling. Cancer Letters, 2015, 358, 124-135.	3.2	157
46	Endoplasmic reticulum stress induces up-regulation of hepatic \hat{l}^2 -Klotho expression through ATF4 signaling pathway. Biochemical and Biophysical Research Communications, 2015, 459, 300-305.	1.0	17
47	PKM2 promotes metastasis by recruiting myeloid-derived suppressor cells and indicates poor prognosis for hepatocellular carcinoma. Oncotarget, 2015, 6, 846-861.	0.8	84
48	LOXL4 is downregulated in hepatocellular carcinoma with a favorable prognosis. International Journal of Clinical and Experimental Pathology, 2015, 8, 3892-900.	0.5	9
49	Fibroblast Growth Factor 21 Is Regulated by the IRE1α-XBP1 Branch of the Unfolded Protein Response and Counteracts Endoplasmic Reticulum Stress-induced Hepatic Steatosis. Journal of Biological Chemistry, 2014, 289, 29751-29765.	1.6	147
50	Activating Mutations in PTPN3 Promote Cholangiocarcinoma Cell Proliferation and Migration and Are Associated With Tumor Recurrence in Patients. Gastroenterology, 2014, 146, 1397-1407.	0.6	111
51	HNRNPAB Induces Epithelial–Mesenchymal Transition and Promotes Metastasis of Hepatocellular Carcinoma by Transcriptionally Activating <i>SNAIL</i> . Cancer Research, 2014, 74, 2750-2762.	0.4	91
52	Capn4 contributes to tumour growth and metastasis of hepatocellular carcinoma by activation of the FAK-Src signalling pathways. Journal of Pathology, 2014, 234, 316-328.	2.1	48
53	Tacrolimus promotes hepatocellular carcinoma and enhances CXCR4/SDF- $1\hat{l}\pm$ expression in vivo. Molecular Medicine Reports, 2014, 10, 585-592.	1.1	8
54	MiR-302c inhibits tumor growth of hepatocellular carcinoma by suppressing the endothelial-mesenchymal transition of endothelial cells. Scientific Reports, 2014, 4, 5524.	1.6	68

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55	Overexpression of CXCL5 mediates neutrophil infiltration and indicates poor prognosis for hepatocellular carcinoma. Hepatology, 2012, 56, 2242-2254.	3.6	298
56	Lectinâ€based glycoproteomics to explore and analyze hepatocellular carcinomaâ€related glycoprotein markers. Electrophoresis, 2009, 30, 2957-2966.	1.3	69
57	Identification and analysis of $\hat{l}\pm 1,6\hat{a}$ (ucosylated proteins in human normal liver tissues by a target glycoproteomic approach. Electrophoresis, 2007, 28, 4382-4391.	1.3	20
58	Identification and analysis of altered $\hat{l}\pm 1$,6-fucosylated glycoproteins associated with hepatocellular carcinoma metastasis. Proteomics, 2006, 6, 5857-5867.	1.3	50
59	Identification of metastasis candidate proteins among HCC cell lines by comparative proteome and biological function analysis of S100A4 in metastasis in vitro., 2006, 6, 5953.		1