Xiang-Huo He

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Circular RNA is enriched and stable in exosomes: a promising biomarker for cancer diagnosis. Cell Research, 2015, 25, 981-984.	5.7	1,777
2	Circular RNA profiling reveals an abundant circHIPK3 that regulates cell growth by sponging multiple miRNAs. Nature Communications, 2016, 7, 11215.	5.8	1,729
3	Circular RNA profile identifies circPVT1 as a proliferative factor and prognostic marker in gastric cancer. Cancer Letters, 2017, 388, 208-219.	3.2	603
4	IKKβ Suppression of TSC1 Links Inflammation and Tumor Angiogenesis via the mTOR Pathway. Cell, 2007, 130, 440-455.	13.5	585
5	exoRBase: a database of circRNA, IncRNA and mRNA in human blood exosomes. Nucleic Acids Research, 2018, 46, D106-D112.	6.5	415
6	Multiple microRNAs modulate p21Cip1/Waf1 expression by directly targeting its 3′ untranslated region. Oncogene, 2010, 29, 2302-2308.	2.6	351
7	Degradation of Mcl-1 by β-TrCP Mediates Glycogen Synthase Kinase 3-Induced Tumor Suppression and Chemosensitization. Molecular and Cellular Biology, 2007, 27, 4006-4017.	1.1	348
8	Gain of miR-151 on chromosome 8q24.3 facilitates tumour cell migration and spreading through downregulating RhoGDIA. Nature Cell Biology, 2010, 12, 390-399.	4.6	290
9	Diagnostic and prognostic implications of microRNAs in human hepatocellular carcinoma. International Journal of Cancer, 2008, 123, 1616-1622.	2.3	287
10	<i>MicroRNA-148a</i> Suppresses Tumor Cell Invasion and Metastasis by Downregulating <i>ROCK1</i> in Gastric Cancer. Clinical Cancer Research, 2011, 17, 7574-7583.	3.2	258
11	MicroRNA-193a-3p and -5p suppress the metastasis of human non-small-cell lung cancer by downregulating the ERBB4/PIK3R3/mTOR/S6K2 signaling pathway. Oncogene, 2015, 34, 413-423.	2.6	238
12	MicroRNA-125b suppressesed human liver cancer cell proliferation and metastasis by directly targeting oncogene LIN28B2. Hepatology, 2010, 52, 1731-1740.	3.6	225
13	The role of microRNAs in liver cancer progression. British Journal of Cancer, 2011, 104, 235-240.	2.9	208
14	Upregulation of miRâ€23aâ^1⁄427aâ^1⁄424 decreases transforming growth factorâ€betaâ€induced tumorâ€suppre activities in human hepatocellular carcinoma cells. International Journal of Cancer, 2008, 123, 972-978.	ssive 2.3	198
15	MiRâ€199aâ€5p is negatively associated with malignancies and regulates glycolysis and lactate production by targeting hexokinase 2 in liver cancer. Hepatology, 2015, 62, 1132-1144.	3.6	196
16	MicroRNA-30d promotes tumor invasion and metastasis by targeting Galphai2 in hepatocellular carcinoma. Hepatology, 2010, 51, NA-NA.	3.6	195
17	Long noncoding RNA TSLNC8 is a tumor suppressor that inactivates the interleukinâ€6/STAT3 signaling pathway. Hepatology, 2018, 67, 171-187.	3.6	183
18	Comprehensive transcriptome analysis identifies novel molecular subtypes and subtype-specific RNAs of triple-negative breast cancer. Breast Cancer Research, 2016, 18, 33.	2.2	176

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19	Extracellular Vesicles Long RNA Sequencing Reveals Abundant mRNA, circRNA, and IncRNA in Human Blood as Potential Biomarkers for Cancer Diagnosis. Clinical Chemistry, 2019, 65, 798-808.	1.5	174
20	Myeloid Cell Leukemia-1 Inversely Correlates with Glycogen Synthase Kinase-3Î ² Activity and Associates with Poor Prognosis in Human Breast Cancer. Cancer Research, 2007, 67, 4564-4571.	0.4	171
21	Hypoxia-inducible MicroRNA-210 augments the metastatic potential of tumor cells by targeting vacuole membrane protein 1 in hepatocellular carcinoma. Hepatology, 2011, 54, 2064-2075.	3.6	162
22	The LINC01138 drives malignancies via activating arginine methyltransferase 5 in hepatocellular carcinoma. Nature Communications, 2018, 9, 1572.	5.8	157
23	Long noncoding RNA miR503HG, a prognostic indicator, inhibits tumor metastasis by regulating the HNRNPA2B1/NF-κB pathway in hepatocellular carcinoma. Theranostics, 2018, 8, 2814-2829.	4.6	151
24	CXCR6 Upregulation Contributes to a Proinflammatory Tumor Microenvironment That Drives Metastasis and Poor Patient Outcomes in Hepatocellular Carcinoma. Cancer Research, 2012, 72, 3546-3556.	0.4	150
25	Hypoxia-inducible factor 1 alpha-activated angiopoietin-like protein 4 contributes to tumor metastasis via vascular cell adhesion molecule-1/integrin l²1 signaling in human hepatocellular carcinoma. Hepatology, 2011, 54, 910-919.	3.6	144
26	Plasma extracellular vesicle long RNA profiling identifies a diagnostic signature for the detection of pancreatic ductal adenocarcinoma. Gut, 2020, 69, 540-550.	6.1	142
27	Increased expression of long noncoding RNA TUG1 predicts a poor prognosis of gastric cancer and regulates cell proliferation by epigenetically silencing of p57. Cell Death and Disease, 2016, 7, e2109-e2109.	2.7	140
28	MetaLnc9 Facilitates Lung Cancer Metastasis via a PGK1-Activated AKT/mTOR Pathway. Cancer Research, 2017, 77, 5782-5794.	0.4	139
29	Disruption of xCT inhibits cancer cell metastasis via the caveolin-1/β-catenin pathway. Oncogene, 2009, 28, 599-609.	2.6	131
30	MicroRNA-95 Promotes Cell Proliferation and Targets Sorting Nexin 1 in Human Colorectal Carcinoma. Cancer Research, 2011, 71, 2582-2589.	0.4	129
31	Genome-wide screening reveals that miR-195 targets the TNF-α/NF-κB pathway by down-regulating lκB kinase alpha and TAB3 in hepatocellular carcinoma. Hepatology, 2013, 58, 654-666.	3.6	118
32	MicroRNA-409 suppresses tumour cell invasion and metastasis by directly targeting radixin in gastric cancers. Oncogene, 2012, 31, 4509-4516.	2.6	116
33	Comprehensive characterization of circular RNAs in ~ 1000 human cancer cell lines. Genome Medicine, 2019, 11, 55.	3.6	116
34	Exome sequencing of hepatoblastoma reveals novel mutations and cancer genes in the Wnt pathway and ubiquitin ligase complex. Hepatology, 2014, 60, 1686-1696.	3.6	115
35	MiR-181a confers resistance of cervical cancer to radiation therapy through targeting the pro-apoptotic PRKCD gene. Oncogene, 2013, 32, 3019-3027.	2.6	113
36	Genome-wide copy number analyses identified novel cancer genes in hepatocellular carcinoma. Hepatology, 2011, 54, 1227-1236.	3.6	112

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37	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
38	miRNA-200c inhibits invasion and metastasis of human non-small cell lung cancer by directly targeting ubiquitin specific peptidase 25. Molecular Cancer, 2014, 13, 166.	7.9	110
39	MicroRNA-30d-5p inhibits tumour cell proliferation and motility by directly targeting CCNE2 in non-small cell lung cancer. Cancer Letters, 2015, 362, 208-217.	3.2	110
40	MicroRNA-423 promotes cell growth and regulates G 1 /S transition by targeting p21Cip1/Waf1 in hepatocellular carcinoma. Carcinogenesis, 2011, 32, 1641-1647.	1.3	107
41	Disruption of xCT inhibits cell growth via the ROS/autophagy pathway in hepatocellular carcinoma. Cancer Letters, 2011, 312, 55-61.	3.2	95
42	The endogenous retrovirus-derived long noncoding RNA TROJAN promotes triple-negative breast cancer progression via ZMYND8 degradation. Science Advances, 2019, 5, eaat9820.	4.7	95
43	Hypoxia induced LUCAT1/PTBP1 axis modulates cancer cell viability and chemotherapy response. Molecular Cancer, 2020, 19, 11.	7.9	92
44	Knockdown of splicing factor SRp20 causes apoptosis in ovarian cancer cells and its expression is associated with malignancy of epithelial ovarian cancer. Oncogene, 2011, 30, 356-365.	2.6	91
45	Programmed death ligand 1 promotes lymph node metastasis and glucose metabolism in cervical cancer by activating integrin β4/SNAI1/SIRT3 signaling pathway. Oncogene, 2018, 37, 4164-4180.	2.6	91
46	Histone lysine methyltransferase, suppressor of variegation 3-9 homolog 1, promotes hepatocellular carcinoma progression and is negatively regulated by microRNA-125b. Hepatology, 2013, 57, 637-647.	3.6	90
47	Involvement of polypyrimidine tract-binding protein (PTBP1) in maintaining breast cancer cell growth and malignant properties. Oncogenesis, 2014, 3, e84-e84.	2.1	90
48	Sphingosine kinase 1 promotes tumour cell migration and invasion via the <scp>S1P</scp> / <scp>EDG</scp> 1 axis in hepatocellular carcinoma. Liver International, 2012, 32, 331-338.	1.9	89
49	Transcriptomeâ€Wide Analysis Reveals the Landscape of Aberrant Alternative Splicing Events in Liver Cancer. Hepatology, 2019, 69, 359-375.	3.6	86
50	Integrative Analyses Identify Osteopontin, LAMB3 and ITGB1 as Critical Pro-Metastatic Genes for Lung Cancer. PLoS ONE, 2013, 8, e55714.	1.1	81
51	MicroRNA-124 Reduces the Pentose Phosphate Pathway and Proliferation by Targeting PRPS1 and RPIA mRNAs in Human Colorectal Cancer Cells. Gastroenterology, 2015, 149, 1587-1598.e11.	0.6	80
52	MicroRNA-181a modulates gene expression of zinc finger family members by directly targeting their coding regions. Nucleic Acids Research, 2010, 38, 7211-7218.	6.5	79
53	Genome-Wide Analysis of Long Noncoding RNA (IncRNA) Expression in Hepatoblastoma Tissues. PLoS ONE, 2014, 9, e85599.	1.1	78
54	Acetylcholinesterase, a key prognostic predictor for hepatocellular carcinoma, suppresses cell growth and induces chemosensitization. Hepatology, 2011, 53, 493-503.	3.6	75

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55	miR-27b synergizes with anticancer drugs via p53 activation and CYP1B1 suppression. Cell Research, 2015, 25, 477-495.	5.7	75
56	MicroRNA-1285 inhibits the expression of p53 by directly targeting its 3′ untranslated region. Biochemical and Biophysical Research Communications, 2010, 396, 435-439.	1.0	73
57	MicroRNA-202-3p Inhibits Cell Proliferation by Targeting ADP-Ribosylation Factor-like 5A in Human Colorectal Carcinoma. Clinical Cancer Research, 2014, 20, 1146-1157.	3.2	72
58	MiR-200b/200c/429 subfamily negatively regulates Rho/ROCK signaling pathway to suppress hepatocellular carcinoma metastasis. Oncotarget, 2015, 6, 13658-13670.	0.8	70
59	Genome-Wide Screening Identified That miR-134 Acts as a Metastasis Suppressor by Targeting Integrin β1 in Hepatocellular Carcinoma. PLoS ONE, 2014, 9, e87665.	1.1	69
60	miR-192, a prognostic indicator, targets the SLC39A6/SNAIL pathway to reduce tumor metastasis in human hepatocellular carcinoma. Oncotarget, 2016, 7, 2672-2683.	0.8	68
61	Development of a highly metastatic model that reveals a crucial role of fibronectin in lung cancer cell migration and invasion. BMC Cancer, 2010, 10, 364.	1.1	65
62	TRIM35 Interacts with pyruvate kinase isoform M2 to suppress the Warburg effect and tumorigenicity in hepatocellular carcinoma. Oncogene, 2015, 34, 3946-3956.	2.6	65
63	Arginine methylation of <scp>SIRT</scp> 7 couples glucose sensing with mitochondria biogenesis. EMBO Reports, 2018, 19, .	2.0	64
64	MicroRNA-135b, a HSF1 target, promotes tumor invasion and metastasis by regulating RECK and EVI5 in hepatocellular carcinoma. Oncotarget, 2015, 6, 2421-2433.	0.8	64
65	Amplification of MPZL1/PZR promotes tumor cell migration through Src-mediated phosphorylation of cortactin in hepatocellular carcinoma. Cell Research, 2014, 24, 204-217.	5.7	61
66	LncRNA SNHG11 facilitates tumor metastasis by interacting with and stabilizing HIF-11±. Oncogene, 2020, 39, 7005-7018.	2.6	60
67	NF-κB signaling relieves negative regulation by miR-194 in hepatocellular carcinoma by suppressing the transcription factor HNF-1α. Science Signaling, 2015, 8, ra75.	1.6	59
68	MicroRNA-129-5p Regulates Glycolysis and Cell Proliferation by Targeting the Glucose Transporter SLC2A3 in Gastric Cancer Cells. Frontiers in Pharmacology, 2018, 9, 502.	1.6	59
69	Choline Transporters in Human Lung Adenocarcinoma: Expression and Functional Implications. Acta Biochimica Et Biophysica Sinica, 2007, 39, 668-674.	0.9	55
70	MicroRNAâ€127â€5p targets the biliverdin reductase B/nuclear factorâ€₽̂B pathway to suppress cell growth in hepatocellular carcinoma cells. Cancer Science, 2016, 107, 258-266.	1.7	55
71	HNRNPL Circularizes ARHGAP35 to Produce an Oncogenic Protein. Advanced Science, 2021, 8, 2001701.	5.6	55
72	MicroRNA-550a Acts as a Pro-Metastatic Gene and Directly Targets Cytoplasmic Polyadenylation Element-Binding Protein 4 in Hepatocellular Carcinoma. PLoS ONE, 2012, 7, e48958.	1.1	54

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73	Application of microRNA and mRNA expression profiling on prognostic biomarker discovery for hepatocellular carcinoma. BMC Genomics, 2014, 15, S13.	1.2	54
74	LncRNA ID2-AS1 suppresses tumor metastasis by activating the HDAC8/ID2 pathway in hepatocellular carcinoma. Cancer Letters, 2020, 469, 399-409.	3.2	54
75	Genome-wide analysis reveals that exon methylation facilitates its selective usage in the human transcriptome. Briefings in Bioinformatics, 2018, 19, 754-764.	3.2	52
76	Inflammationâ€Induced Long Intergenic Noncoding RNA (LINC00665) Increases Malignancy Through Activating the Doubleâ€Stranded RNA–Activated Protein Kinase/Nuclear Factor Kappa B Pathway in Hepatocellular Carcinoma. Hepatology, 2020, 72, 1666-1681.	3.6	52
77	Quantitative Proteomic Analysis Identifies CPNE3 as a Novel Metastasis-promoting Gene in NSCLC. Journal of Proteome Research, 2013, 12, 3423-3433.	1.8	50
78	Co-expression of PKM2 and TRIM35 predicts survival and recurrence in hepatocellular carcinoma. Oncotarget, 2015, 6, 2539-2548.	0.8	50
79	GNAI1 Suppresses Tumor Cell Migration and Invasion and is Post-Transcriptionally Regulated by Mir-320a/c/d in Hepatocellular Carcinoma. Cancer Biology and Medicine, 2012, 9, 234-41.	1.4	48
80	Analysis of acetylcholine, choline and butyrobetaine in human liver tissues by hydrophilic interaction liquid chromatography-tandem mass spectrometry. Journal of Pharmaceutical and Biomedical Analysis, 2008, 47, 870-875.	1.4	46
81	Quantitative Proteomic Analysis of the Metastasis-Inhibitory Mechanism of miR-193a-3p in Non-Small Cell Lung Cancer. Cellular Physiology and Biochemistry, 2015, 35, 1677-1688.	1.1	44
82	Gain of LINC00624 Enhances Liver Cancer Progression by Disrupting the Histone Deacetylase 6/Tripartite Motif Containing 28/Zinc Finger Protein 354C Corepressor Complex. Hepatology, 2021, 73, 1764-1782.	3.6	42
83	SERPINA5 inhibits tumor cell migration by modulating the fibronectinâ€integrin β1 signaling pathway in hepatocellular carcinoma. Molecular Oncology, 2014, 8, 366-377.	2.1	41
84	GNAI3 inhibits tumor cell migration and invasion and is post-transcriptionally regulated by miR-222 in hepatocellular carcinoma. Cancer Letters, 2015, 356, 978-984.	3.2	40
85	Splicing Regulator p54nrb/Non–POU Domain–Containing Octamerâ€Binding Protein Enhances Carcinogenesis Through Oncogenic Isoform Switch of MYC Box–Dependent Interacting Protein 1 in Hepatocellular Carcinoma. Hepatology, 2020, 72, 548-568.	3.6	40
86	An LTR Retrotransposon-Derived Long Noncoding RNA IncMER52A Promotes Hepatocellular Carcinoma Progression by Binding p120-Catenin. Cancer Research, 2020, 80, 976-987.	0.4	39
87	Transcriptomic analyses of <scp>RNA</scp> â€binding proteins reveal <i><scp>elF</scp>3c</i> promotes cell proliferation in hepatocellular carcinoma. Cancer Science, 2017, 108, 877-885.	1.7	38
88	MiR-525-3p Enhances the Migration and Invasion of Liver Cancer Cells by Downregulating ZNF395. PLoS ONE, 2014, 9, e90867.	1.1	33
89	Inactivation of the tumor suppressor p53 by long noncoding RNA RMRP. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	33
90	Integrated Analysis of Mutation Data from Various Sources Identifies Key Genes and Signaling Pathways in Hepatocellular Carcinoma. PLoS ONE, 2014, 9, e100854.	1.1	32

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91	Over-Expression of c-FLIP Confers the Resistance to TRAIL-Induced Apoptosis on Gallbladder Carcinoma. Tohoku Journal of Experimental Medicine, 2009, 217, 203-208.	0.5	29
92	Molecular cloning and characterization of the human ASB-8 gene encoding a novel member of ankyrin repeat and SOCS box containing protein family. Biochemical and Biophysical Research Communications, 2003, 300, 972-979.	1.0	28
93	microRNAs: tiny RNA molecules, huge driving forces to move the cell. Protein and Cell, 2010, 1, 916-926.	4.8	27
94	Cloning and characterization of a novel gene which encodes a protein interacting with the mitosis-associated kinase-like protein NTKL. Journal of Human Genetics, 2003, 48, 315-321.	1.1	26
95	Hepatic SMARCA4 predicts HCC recurrence and promotes tumour cell proliferation by regulating SMAD6 expression. Cell Death and Disease, 2018, 9, 59.	2.7	26
96	CRISPR/Cas9 Screens Reveal that Hexokinase 2 Enhances Cancer Stemness and Tumorigenicity by Activating the ACSL4â€Fatty Acid <i>β</i> â€Oxidation Pathway. Advanced Science, 2022, 9, .	5.6	26
97	Macro-management of microRNAs in cell cycle progression of tumor cells and its implications in anti-cancer therapy. Acta Pharmacologica Sinica, 2011, 32, 1311-1320.	2.8	24
98	Resection of liver metastases from breast cancer: a multicentre analysis. Clinical and Translational Oncology, 2020, 22, 512-521.	1.2	24
99	Speckle-type POZ protein is negatively associated with malignancies and inhibits cell proliferation and migration in liver cancer. Tumor Biology, 2015, 36, 9753-9761.	0.8	22
100	A LIN28B Tumor-Specific Transcript in Cancer. Cell Reports, 2018, 22, 2016-2025.	2.9	22
101	Molecular cloning and characterization of CT120, a novel membrane-associated gene involved in amino acid transport and glutathione metabolism. Biochemical and Biophysical Research Communications, 2002, 297, 528-536.	1.0	20
102	Role of microRNAs in inflammation-associated liver cancer. Cancer Biology and Medicine, 2016, 13, 407.	1.4	20
103	Transcriptome analysis of Luminal Breast Cancer Reveals a Role for LOL in Tumor Progression and Tamoxifen Resistance. International Journal of Cancer, 2019, 145, 842-856.	2.3	20
104	Ciliary neurotrophic factor receptor α subunit-modulated multiple downstream signaling pathways in hepatic cancer cell lines and their biological implications. Hepatology, 2008, 47, 1298-1308.	3.6	19
105	Predicting Value of ALCAM as a Target Gene of microRNA-483-5p in Patients with Early Recurrence in Hepatocellular Carcinoma. Frontiers in Pharmacology, 2017, 8, 973.	1.6	19
106	Tumorâ€ S pecific Transcripts Are Frequently Expressed in Hepatocellular Carcinoma With Clinical Implication and Potential Function. Hepatology, 2020, 71, 259-274.	3.6	16
107	Altered gene expression profiles of NIH3T3 cells regulated by human lung cancer associated gene CT120. Cell Research, 2004, 14, 487-496.	5.7	14
108	Real-time imaging nuclear translocation of Akt1 in HCC cells. Biochemical and Biophysical Research Communications, 2007, 356, 1038-1043.	1.0	14

Xiang-Huo He

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109	TC21 promotes cell motility and metastasis by regulating the expression of E-cadherin and N-cadherin in hepatocellular carcinoma. International Journal of Oncology, 2010, 37, 853-9.	1.4	13
110	The Mutational and Transcriptional Landscapes of Hepatocarcinogenesis in a Rat Model. IScience, 2020, 23, 101690.	1.9	12
111	Suppression of Human Liver Cancer Cell Migration and Invasion via the GABAA Receptor. Cancer Biology and Medicine, 2012, 9, 90-8.	1.4	12
112	ASJA: A Program for Assembling Splice Junctions Analysis. Computational and Structural Biotechnology Journal, 2019, 17, 1143-1150.	1.9	11
113	Differential gene expression in human hepatocellular carcinoma Hep3B cells induced by apoptosis-related gene <i>BNIPL-2</i> . World Journal of Gastroenterology, 2004, 10, 1286.	1.4	11
114	LncRNA RP11-295G20.2 regulates hepatocellular carcinoma cell growth and autophagy by targeting PTEN to lysosomal degradation. Cell Discovery, 2021, 7, 118.	3.1	11
115	RNA binding protein RALY activates the cholesterol synthesis pathway through an MTA1 splicing switch in hepatocellular carcinoma. Cancer Letters, 2022, 538, 215711.	3.2	11
116	Cloning and characterization of human IC53-2, a novel CDK5 activator binding protein. Cell Research, 2003, 13, 83-91.	5.7	10
117	Molecular cloning and characterization of human Aph2 gene, involved in AP-1 regulation by interaction with JAB1. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2006, 1759, 514-525.	2.4	10
118	B7-H4 enhances the differentiation of murine leukemia-initiating cells via the PTEN/AKT/RCOR2/RUNX1 pathways. Leukemia, 2017, 31, 2260-2264.	3.3	10
119	HNRNPH1-stabilized LINC00662 promotes ovarian cancer progression by activating the GRP78/p38 pathway. Oncogene, 2021, 40, 4770-4782.	2.6	10
120	HCC-Associated Protein HCAP1, a Variant of GEMIN4, Interacts with Zinc-Finger Proteins. Journal of Biochemistry, 2003, 133, 713-718.	0.9	9
121	RNA Helicase DHX37 Facilitates Liver Cancer Progression by Cooperating with PLRG1 to Drive Superenhancer-Mediated Transcription of Cyclin D1. Cancer Research, 2022, 82, 1937-1952.	0.4	9
122	A decrease in serum 1,5â€anhydroglucitol levels is associated with the presence of a firstâ€degree family history of diabetes in a Chinese population with normal glucose tolerance. Diabetic Medicine, 2018, 35, 131-136.	1.2	8
123	Integrative Analysis of Transcriptional Regulatory Network and Copy Number Variation in Intrahepatic Cholangiocarcinoma. PLoS ONE, 2014, 9, e98653.	1.1	6
124	Application of third-generation sequencing in cancer research. Medical Review, 2021, 1, 150-171.	0.3	6
125	cDNA Expression Array Analysis of Gene Expression in Human Hepatocarcinoma Hep3B Cells Induced By BNIPL-1. Acta Biochimica Et Biophysica Sinica, 2005, 37, 618-624.	0.9	5
126	BNIPL-2 promotes the invasion and metastasis of human hepatocellular carcinoma cells. Oncology Reports, 0, , .	1.2	5

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127	Psychometric assessment and application of the Chinese version of the Compliance with Annual Diabetic Eye Exams Survey in people with diabetic retinopathy. Diabetic Medicine, 2020, 37, 84-94.	1.2	5
128	A pathway-guided strategy identifies a metabolic signature for prognosis prediction and precision therapy for hepatocellular carcinoma. Computers in Biology and Medicine, 2022, 144, 105376.	3.9	5
129	SRTdb: an omnibus for human tissue and cancer-specific RNA transcripts. Biomarker Research, 2022, 10, 27.	2.8	5
130	Dysfunction of Murine Dendritic Cells Induced by Incubation with Tumor Cells. Cellular and Molecular Immunology, 2008, 5, 133-140.	4.8	4
131	Optimized protocol for an inducible rat model of liver tumor with chronic hepatocellular injury, inflammation, fibrosis, and cirrhosis. STAR Protocols, 2021, 2, 100353.	0.5	4
132	CD26/dipeptidyl peptidase IV contributes to tumor metastasis in human lung adenocarcinoma. Bangladesh Journal of Pharmacology, 2013, 8, .	0.1	1
133	FAM57A (family with sequence similarity 57, member A). Atlas of Genetics and Cytogenetics in Oncology and Haematology, 2011, , .	0.1	0
134	Construction and selection of human anti-idiotypic antibody single chain variable fragments or CDR3 fragments of nasopharyngeal carcinoma. Journal of Experimental and Clinical Cancer Research, 2004,	0.4	0

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