

Gang Chen

List of Publications by Year in descending order

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308
papers

8,837
citations

60835

43
h-index

106894

65
g-index

376
all docs

376
docs citations

376
times ranked

13960
citing authors

#	ARTICLE	IF	CITATIONS
1	A circRNA-miRNA-mRNA network identification for exploring underlying pathogenesis and therapy strategy of hepatocellular carcinoma. <i>Journal of Translational Medicine</i> , 2018, 16, 220.	4.5	242
2	miR-146a Inhibits Cell Growth, Cell Migration and Induces Apoptosis in Non-Small Cell Lung Cancer Cells. <i>PLoS ONE</i> , 2013, 8, e60317.	2.5	233
3	Overexpression of MMP Family Members Functions as Prognostic Biomarker for Breast Cancer Patients: A Systematic Review and Meta-Analysis. <i>PLoS ONE</i> , 2015, 10, e0135544.	2.5	158
4	A Pilot Study Examining Activity Participation, Sensory Responsiveness, and Competence in Children with High Functioning Autism Spectrum Disorder. <i>Journal of Autism and Developmental Disorders</i> , 2011, 41, 1496-1506.	3.1	139
5	Expression and Prognostic Significance of lncRNA MALAT1 in Pancreatic Cancer Tissues. <i>Asian Pacific Journal of Cancer Prevention</i> , 2014, 15, 2971-2977.	1.2	138
6	Underexpression of miR-34a in Hepatocellular Carcinoma and Its Contribution towards Enhancement of Proliferating Inhibitory Effects of Agents Targeting c-MET. <i>PLoS ONE</i> , 2013, 8, e61054.	2.5	113
7	Targeting the epidermal growth factor receptor in non-small cell lung cancer cells: the effect of combining RNA interference with tyrosine kinase inhibitors or cetuximab. <i>BMC Medicine</i> , 2012, 10, 28.	5.7	111
8	Increased MiR-221 expression in hepatocellular carcinoma tissues and its role in enhancing cell growth and inhibiting apoptosis in vitro. <i>BMC Cancer</i> , 2013, 13, 21.	2.6	110
9	Ki-67 is a Valuable Prognostic Factor in Gliomas: Evidence from a Systematic Review and Meta-analysis. <i>Asian Pacific Journal of Cancer Prevention</i> , 2015, 16, 411-420.	1.2	110
10	Comprehensive investigation of a novel differentially expressed lncRNA expression profile signature to assess the survival of patients with colorectal adenocarcinoma. <i>Oncotarget</i> , 2017, 8, 16811-16828.	2.1	96
11	A circulating miRNA signature as a diagnostic biomarker for non-invasive early detection of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2015, 154, 423-434.	2.5	95
12	Systematic Analysis of Survival-Associated Alternative Splicing Signatures in Gastrointestinal Pan-Adenocarcinomas. <i>EBioMedicine</i> , 2018, 34, 46-60.	6.0	84
13	Upregulation and Clinicopathological Significance of Long Non-coding NEAT1 RNA in NSCLC Tissues. <i>Asian Pacific Journal of Cancer Prevention</i> , 2015, 16, 2851-2855.	1.2	83
14	Inhibition of β -secretase induces G2/M arrest and triggers apoptosis in breast cancer cells. <i>British Journal of Cancer</i> , 2009, 100, 1879-1888.	6.6	80
15	Time-restricted feeding alters lipid and amino acid metabolite rhythmicity without perturbing clock gene expression. <i>Nature Communications</i> , 2020, 11, 4643.	13.2	80
16	Clinical roles of the aberrantly expressed lncRNAs in lung squamous cell carcinoma: a study based on RNA-sequencing and microarray data mining. <i>Oncotarget</i> , 2017, 8, 61282-61304.	2.1	75
17	Evaluation of the role of radiotherapy in the management of carcinoma of the buccal mucosa. <i>Cancer</i> , 1988, 61, 1326-1331.	4.1	74
18	Identification of a serum microRNA expression signature for detection of lung cancer, involving miR-23b, miR-221, miR-148b and miR-423-3p. <i>Lung Cancer</i> , 2017, 114, 6-11.	2.0	68

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19	Expression and clinicopathological significance of miR-146a in hepatocellular carcinoma tissues. <i>Uppsala Journal of Medical Sciences</i> , 2014, 119, 19-24.	0.9	63
20	Long non-coding RNA TUC338 is functionally involved in sorafenib-sensitized hepatocarcinoma cells by targeting RASAL1. <i>Oncology Reports</i> , 2017, 37, 273-280.	2.6	59
21	Estado del conocimiento de humedales del norte patagónico (Argentina): aspectos relevantes e importancia para la conservación de la biodiversidad regional. <i>Revista Chilena De Historia Natural</i> , 2005, 78, 723.	1.2	58
22	Sp1 cooperates with Sp3 to upregulate MALAT1 expression in human hepatocellular carcinoma. <i>Oncology Reports</i> , 2015, 34, 2403-2412.	2.6	57
23	Comprehensive analysis of the long noncoding RNA HOXA11-AS gene interaction regulatory network in NSCLC cells. <i>Cancer Cell International</i> , 2016, 16, 89.	4.3	56
24	The clinicopathological significance of UBE2C in breast cancer: a study based on immunohistochemistry, microarray and RNA-sequencing data. <i>Cancer Cell International</i> , 2017, 17, 83.	4.3	56
25	Clinicopathological significance of RASSF1A reduced expression and hypermethylation in hepatocellular carcinoma. <i>Hepatology International</i> , 2010, 4, 423-432.	4.4	55
26	DNA topoisomerase 1 and 2A function as oncogenes in liver cancer and may be direct targets of nitidine chloride. <i>International Journal of Oncology</i> , 2018, 53, 1897-1912.	3.2	55
27	High throughput circRNA sequencing analysis reveals novel insights into the mechanism of nitidine chloride against hepatocellular carcinoma. <i>Cell Death and Disease</i> , 2019, 10, 658.	6.4	55
28	Identification of a RNA-Seq based prognostic signature with five lncRNAs for lung squamous cell carcinoma. <i>Oncotarget</i> , 2017, 8, 50761-50773.	2.1	53
29	Effects of miR-152 on Cell Growth Inhibition, Motility Suppression and Apoptosis Induction in Hepatocellular Carcinoma Cells. <i>Asian Pacific Journal of Cancer Prevention</i> , 2014, 15, 4969-4976.	1.2	53
30	Expression of the Long Intergenic Non-Protein Coding RNA 665 (LINC00665) Gene and the Cell Cycle in Hepatocellular Carcinoma Using The Cancer Genome Atlas, the Gene Expression Omnibus, and Quantitative Real-Time Polymerase Chain Reaction. <i>Medical Science Monitor</i> , 2018, 24, 2786-2808.	1.1	52
31	Lithium chloride modulates chondrocyte primary cilia and inhibits Hedgehog signaling. <i>FASEB Journal</i> , 2016, 30, 716-726.	0.5	51
32	A nine-miRNA signature as a potential diagnostic marker for breast carcinoma: An integrated study of 1,110 cases. <i>Oncology Reports</i> , 2017, 37, 3297-3304.	2.6	51
33	Augmented expression of Ki-67 is correlated with clinicopathological characteristics and prognosis for lung cancer patients: an up-dated systematic review and meta-analysis with 108 studies and 14,732 patients. <i>Respiratory Research</i> , 2018, 19, 150.	3.7	50
34	A single-cell polony method reveals low levels of infected <i>Prochlorococcus</i> in oligotrophic waters despite high cyanophage abundances. <i>ISME Journal</i> , 2021, 15, 41-54.	10.0	49
35	Downregulation of MiR-30a is Associated with Poor Prognosis in Lung Cancer. <i>Medical Science Monitor</i> , 2015, 21, 2514-2520.	1.1	49
36	The critical care nurse's role in End-of-Life care: issues and challenges. <i>Nursing in Critical Care</i> , 2011, 16, 116-123.	2.5	48

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37	Decreased expression and clinical significance of miR-148a in hepatocellular carcinoma tissues. <i>European Journal of Medical Research</i> , 2014, 19, 68.	2.2	47
38	Clinical Significance and Effect of lncRNA HOXA11-AS in NSCLC: A Study Based on Bioinformatics, In Vitro and in Vivo Verification. <i>Scientific Reports</i> , 2017, 7, 5567.	3.4	47
39	Down-regulation of microRNA-144-3p and its clinical value in non-small cell lung cancer: a comprehensive analysis based on microarray, miRNA-sequencing, and quantitative real-time PCR data. <i>Respiratory Research</i> , 2019, 20, 48.	3.7	47
40	Prognostic Values of Vimentin Expression and Its Clinicopathological Significance in Non-Small Cell Lung Cancer: A Meta-Analysis of Observational Studies with 4118 Cases. <i>PLoS ONE</i> , 2016, 11, e0163162.	2.5	46
41	Long noncoding RNAs in hepatocellular carcinoma: Novel insights into their mechanism. <i>World Journal of Hepatology</i> , 2015, 7, 2781.	2.0	45
42	Synergistic Effect of Afatinib with Su11274 in Non-Small Cell Lung Cancer Cells Resistant to Gefitinib or Erlotinib. <i>PLoS ONE</i> , 2013, 8, e59708.	2.5	44
43	miR-204 regulates the biological behavior of breast cancer MCF-7 cells by directly targeting FOXA1. <i>Oncology Reports</i> , 2017, 38, 368-376.	2.6	44
44	A radiogenomics signature for predicting the clinical outcome of bladder urothelial carcinoma. <i>European Radiology</i> , 2020, 30, 547-557.	4.6	44
45	Reinvasion by ship rats (<i>Rattus rattus</i>) of forest fragments after eradication. <i>Biological Invasions</i> , 2011, 13, 2391-2408.	2.4	43
46	Clinical Value and Prospective Pathway Signaling of MicroRNA-375 in Lung Adenocarcinoma: A Study Based on the Cancer Genome Atlas (TCGA), Gene Expression Omnibus (GEO) and Bioinformatics Analysis. <i>Medical Science Monitor</i> , 2017, 23, 2453-2464.	1.1	42
47	The suppressive role of miR-542-5p in NSCLC: the evidence from clinical data and in vivo validation using a chick chorioallantoic membrane model. <i>BMC Cancer</i> , 2017, 17, 655.	2.6	42
48	Financial hardship and health risk behavior during COVID-19 in a large US national sample of women. <i>SSM - Population Health</i> , 2021, 13, 100734.	2.9	42
49	Human papillomavirus as a potential risk factor for gastric cancer: a meta-analysis of 1,917 cases. <i>OncoTargets and Therapy</i> , 2016, Volume 9, 7105-7114.	2.1	41
50	Prognostic microRNAs and their potential molecular mechanism in pancreatic cancer: A study based on The Cancer Genome Atlas and bioinformatics investigation. <i>Molecular Medicine Reports</i> , 2018, 17, 939-951.	2.5	40
51	Influence of RT-qPCR primer position on EGFR interference efficacy in lung cancer cells. <i>Biological Procedures Online</i> , 2010, 13, 1.	3.0	39
52	Investigation of miR-136-5p key target genes and pathways in lung squamous cell cancer based on TCGA database and bioinformatics analysis. <i>Pathology Research and Practice</i> , 2018, 214, 644-654.	2.3	39
53	Prospective lncRNA-miRNA-mRNA regulatory network of long non-coding RNA LINC00968 in non-small cell lung cancer A549 cells: A miRNA microarray and bioinformatics investigation. <i>International Journal of Molecular Medicine</i> , 2017, 40, 1895-1906.	4.1	38
54	Downregulated miR-23b-3p expression acts as a predictor of hepatocellular carcinoma progression: A study based on public data and RT-qPCR verification. <i>International Journal of Molecular Medicine</i> , 2018, 41, 2813-2831.	4.1	38

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55	The diagnostic and prognostic values of Ki-67/MIB-1 expression in thyroid cancer: a meta-analysis with 6,051 cases. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 3261-3276.	2.1	37
56	Clinical significance of high expression of miR-452-5p in lung squamous cell carcinoma. <i>Oncology Letters</i> , 2018, 15, 6418-6430.	1.8	37
57	Identification and validation of an individualized autophagy-clinical prognostic index in bladder cancer patients. <i>OncoTargets and Therapy</i> , 2019, Volume 12, 3695-3712.	2.1	37
58	Long non-coding RNA HOTTIP promotes hepatocellular carcinoma tumorigenesis and development: A comprehensive investigation based on bioinformatics, qRT-PCR and meta-analysis of 393 cases. <i>International Journal of Oncology</i> , 2017, 51, 1705-1721.	3.2	36
59	Cervical Cancer Growth Is Regulated by a c-ABL/PLK1 Signaling Axis. <i>Cancer Research</i> , 2017, 77, 1142-1154.	0.9	36
60	Prognosis of clear cell renal cell carcinoma (ccRCC) based on a six-lncRNA-based risk score: an investigation based on RNA-sequencing data. <i>Journal of Translational Medicine</i> , 2019, 17, 281.	4.5	36
61	Effect of siRNAs targeting the EGFR T790M mutation in a non-small cell lung cancer cell line resistant to EGFR tyrosine kinase inhibitors and combination with various agents. <i>Biochemical and Biophysical Research Communications</i> , 2013, 431, 623-629.	2.2	35
62	Down-regulation of ribosomal protein S15A mRNA with a short hairpin RNA inhibits human hepatic cancer cell growth in vitro. <i>Gene</i> , 2014, 536, 84-89.	2.3	35
63	MiR-133a is downregulated in non-small cell lung cancer: a study of clinical significance. <i>European Journal of Medical Research</i> , 2015, 20, 50.	2.2	35
64	Association between underexpression of microRNA-203 and clinicopathological significance in hepatocellular carcinoma tissues. <i>Cancer Cell International</i> , 2015, 15, 62.	4.3	34
65	Neurotensin signaling stimulates glioblastoma cell proliferation by upregulating c-Myc and inhibiting miR-29b-1 and miR-129-3p. <i>Neuro-Oncology</i> , 2016, 18, 216-226.	1.2	34
66	Downregulation of miR-224-5p in prostate cancer and its relevant molecular mechanism via TCGA, GEO database and in silico analyses. <i>Oncology Reports</i> , 2018, 40, 3171-3188.	2.6	34
67	Lower expressed miR-198 and its potential targets in hepatocellular carcinoma: a clinicopathological and in silico study. <i>OncoTargets and Therapy</i> , 2016, Volume 9, 5163-5180.	2.1	33
68	From big data to diagnosis and prognosis: gene expression signatures in liver hepatocellular carcinoma. <i>PeerJ</i> , 2017, 5, e3089.	2.0	33
69	Identification of molecular targets for esophageal carcinoma diagnosis using miRNA-seq and RNA-seq data from The Cancer Genome Atlas: a study of 187 cases. <i>Oncotarget</i> , 2017, 8, 35681-35699.	2.1	32
70	Synergistic Effect of MiR-146a Mimic and Cetuximab on Hepatocellular Carcinoma Cells. <i>BioMed Research International</i> , 2014, 2014, 1-15.	2.0	31
71	Ki-67/MKI67 as a Predictive Biomarker for Clinical Outcome in Gastric Cancer Patients: an Updated Meta-analysis and Systematic Review involving 53 Studies and 7078 Patients. <i>Journal of Cancer</i> , 2019, 10, 5339-5354.	2.6	31
72	Expression and clinicopathological significance of miR-193a-3p and its potential target astrocyte elevated gene-1 in non-small lung cancer tissues. <i>Cancer Cell International</i> , 2015, 15, 80.	4.3	30

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73	Comprehensive investigation of aberrant microRNA profiling in bladder cancer tissues. <i>Tumor Biology</i> , 2016, 37, 12555-12569.	1.7	30
74	Downregulation of miR-136-5p in hepatocellular carcinoma and its clinicopathological significance. <i>Molecular Medicine Reports</i> , 2017, 16, 5393-5405.	2.5	30
75	Role of downregulated miR-133a-3p expression in bladder cancer: a bioinformatics study. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 3667-3683.	2.1	30
76	Upregulation of HOXA11 during the progression of lung adenocarcinoma detected via multiple approaches. <i>International Journal of Molecular Medicine</i> , 2018, 42, 2650-2664.	4.1	30
77	In silico analysis identified miRNA-based therapeutic agents against glioblastoma multiforme. <i>Oncology Reports</i> , 2019, 41, 2194-2208.	2.6	30
78	Caspase-3 over-expression is associated with poor overall survival and clinicopathological parameters in breast cancer: a meta-analysis of 3091 cases. <i>Oncotarget</i> , 2018, 9, 8629-8641.	2.1	30
79	Implication of downregulation and prospective pathway signaling of microRNA-375 in lung squamous cell carcinoma. <i>Pathology Research and Practice</i> , 2017, 213, 364-372.	2.3	29
80	The Prognostic Role of Ki-67/MIB-1 in Cervical Cancer: A Systematic Review with Meta-Analysis. <i>Medical Science Monitor</i> , 2015, 21, 882-889.	1.1	28
81	RNA-sequencing investigation identifies an effective risk score generated by three novel lncRNAs for the survival of papillary thyroid cancer patients. <i>Oncotarget</i> , 2017, 8, 74139-74158.	2.1	28
82	A comprehensive insight into the clinicopathologic significance of miR-144-3p in hepatocellular carcinoma. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 3405-3419.	2.1	28
83	Clinical value of miR-182-5p in lung squamous cell carcinoma: a study combining data from TCGA, GEO, and RT-qPCR validation. <i>World Journal of Surgical Oncology</i> , 2018, 16, 76.	1.9	28
84	The role of upregulated miR-375 expression in breast cancer: An in vitro and in silico study. <i>Pathology Research and Practice</i> , 2020, 216, 152754.	2.3	28
85	Down-Regulation of MiR-193a-3p Dictates Deterioration of HCC: A Clinical Real-Time qRT-PCR Study. <i>Medical Science Monitor</i> , 2015, 21, 2352-2360.	1.1	27
86	Downregulation of microRNA-132 indicates progression in hepatocellular carcinoma. <i>Experimental and Therapeutic Medicine</i> , 2016, 12, 2095-2101.	1.9	27
87	Relationship between TRAF6 and deterioration of HCC: an immunohistochemical and in vitro study. <i>Cancer Cell International</i> , 2016, 16, 76.	4.3	27
88	Transshipment hub selection from a shipper's and freight forwarder's perspective. <i>Expert Systems With Applications</i> , 2017, 83, 396-404.	7.9	27
89	The clinical value of lncRNA NEAT1 in digestive system malignancies: A comprehensive investigation based on 57 microarray and RNA-seq datasets. <i>Oncotarget</i> , 2017, 8, 17665-17683.	2.1	27
90	Osteosarcopenic obesity and its relationship with dyslipidemia in women from different ethnic groups of China. <i>Archives of Osteoporosis</i> , 2018, 13, 65.	2.6	27

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91	MiR-182-5p and its target HOXA9 in non-small cell lung cancer: a clinical and in-silico exploration with the combination of RT-qPCR, miRNA-seq and miRNA-chip. BMC Medical Genomics, 2020, 13, 3.	1.5	27
92	LPCAT1 overexpression promotes the progression of hepatocellular carcinoma. Cancer Cell International, 2021, 21, 442.	4.3	27
93	Clinical significance and effect of AEG-1 on the proliferation, invasion, and migration of NSCLC: a study based on immunohistochemistry, TCGA, bioinformatics, <i>in vitro</i> and <i>in vivo</i> verification. Oncotarget, 2017, 8, 16531-16552.	2.1	27
94	An immunohistochemical study of cyclin-dependent kinase 5 (CDK5) expression in non-small cell lung cancer (NSCLC) and small cell lung cancer (SCLC): a possible prognostic biomarker. World Journal of Surgical Oncology, 2015, 14, 34.	1.9	26
95	An Encapsulation of Gene Signatures for Hepatocellular Carcinoma, MicroRNA-132 Predicted Target Genes and the Corresponding Overlaps. PLoS ONE, 2016, 11, e0159498.	2.5	26
96	The impact of atosiban on pregnancy outcomes in women undergoing in vitro fertilization-embryo transfer: A meta-analysis. PLoS ONE, 2017, 12, e0175501.	2.5	26
97	Diagnostic and prognostic roles of IRAK1 in hepatocellular carcinoma tissues: an analysis of immunohistochemistry and RNA-sequencing data from the cancer genome atlas. OncoTargets and Therapy, 2017, Volume 10, 1711-1723.	2.1	26
98	The clinical value and potential molecular mechanism of the downregulation of MAOA in hepatocellular carcinoma tissues. Cancer Medicine, 2020, 9, 8004-8019.	2.9	26
99	Identification of a Four Hypoxia-Associated Long Non-Coding RNA Signature and Establishment of a Nomogram Predicting Prognosis of Clear Cell Renal Cell Carcinoma. Frontiers in Oncology, 2021, 11, 713346.	2.9	26
100	Clinical Value of miR-101-3p and Biological Analysis of its Prospective Targets in Breast Cancer: A Study Based on The Cancer Genome Atlas (TCGA) and Bioinformatics. Medical Science Monitor, 2017, 23, 1857-1871.	1.1	26
101	Analysis of microarrays of miR-34a and its identification of prospective target gene signature in hepatocellular carcinoma. BMC Cancer, 2018, 18, 12.	2.6	25
102	The underlying molecular mechanism and potential drugs for treatment in papillary renal cell carcinoma: A study based on TCGA and Cmap datasets. Oncology Reports, 2019, 41, 2089-2102.	2.6	25
103	Astrocyte Elevated Gene-1 as a Novel Clinicopathological and Prognostic Biomarker for Gastrointestinal Cancers: A Meta-Analysis with 2999 Patients. PLoS ONE, 2015, 10, e0145659.	2.5	25
104	Utility of miR-133a-3p as a diagnostic indicator for hepatocellular carcinoma: An investigation combined with GEO, TCGA, meta-analysis and bioinformatics. Molecular Medicine Reports, 2018, 17, 1469-1484.	2.5	24
105	miR-1296-5p decreases ERBB2 expression to inhibit the cell proliferation in ERBB2-positive breast cancer. Cancer Cell International, 2017, 17, 95.	4.3	24
106	Expression level and potential target pathways of miR-1-3p in colorectal carcinoma based on 645 cases from 9 microarray datasets. Molecular Medicine Reports, 2018, 17, 5013-5020.	2.5	24
107	Prospective molecular mechanism of COL5A1 in breast cancer based on a microarray, RNA sequencing and immunohistochemistry. Oncology Reports, 2019, 42, 151-175.	2.6	24
108	Determining the prognostic significance of alternative splicing events in soft tissue sarcoma using data from The Cancer Genome Atlas. Journal of Translational Medicine, 2019, 17, 283.	4.5	24

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109	Clinical significance and potential molecular mechanism of miRNA-222-3p in metastatic prostate cancer. <i>Bioengineered</i> , 2021, 12, 325-340.	3.2	24
110	Quantification of epidermal growth factor receptor T790M mutant transcripts in lung cancer cells by real-time reverse transcriptase-quantitative polymerase chain reaction. <i>Analytical Biochemistry</i> , 2010, 398, 266-268.	2.5	23
111	MicroRNA-141 Is a Biomarker for Progression of Squamous Cell Carcinoma and Adenocarcinoma of the Lung: Clinical Analysis of 125 Patients. <i>Tohoku Journal of Experimental Medicine</i> , 2015, 235, 161-169.	1.3	23
112	Longitudinal monitoring of heartbeat dynamics predicts mood changes in bipolar patients: A pilot study. <i>Journal of Affective Disorders</i> , 2017, 209, 30-38.	4.2	23
113	The expression of HOXA13 in lung adenocarcinoma and its clinical significance: A study based on The Cancer Genome Atlas, Oncomine and reverse transcription-quantitative polymerase chain reaction. <i>Oncology Letters</i> , 2018, 15, 8556-8572.	1.8	23
114	The clinical significance of endothelin receptor type B in hepatocellular carcinoma and its potential molecular mechanism. <i>Experimental and Molecular Pathology</i> , 2019, 107, 141-157.	2.3	23
115	Investigation on the structural quality dependent electromagnetic interference shielding performance of few-layer and lamellar Nb ₂ CTx MXene nanostructures. <i>Journal of Alloys and Compounds</i> , 2021, 877, 160235.	5.7	23
116	Survival prediction of kidney renal papillary cell carcinoma by comprehensive LncRNA characterization. <i>Oncotarget</i> , 2017, 8, 110811-110829.	2.1	23
117	Down-regulation of miR-26a-5p in hepatocellular carcinoma: A qRT-PCR and bioinformatics study. <i>Pathology Research and Practice</i> , 2017, 213, 1494-1509.	2.3	22
118	MicroRNA-124-3p expression and its prospective functional pathways in hepatocellular carcinoma: A quantitative polymerase chain reaction, gene expression omnibus and bioinformatics study. <i>Oncology Letters</i> , 2018, 15, 5517-5532.	1.8	22
119	Oncogenic role of miR-183-5p in lung adenocarcinoma: A comprehensive study of qPCR, in-vitro experiments and bioinformatic analysis. <i>Oncology Reports</i> , 2018, 40, 83-100.	2.6	22
120	Clinical value and potential mechanisms of COL8A1 upregulation in breast cancer: a comprehensive analysis. <i>Cancer Cell International</i> , 2020, 20, 392.	4.3	22
121	Expression of Heparanase in Hepatocellular Carcinoma Has Prognostic Significance: A Tissue Microarray Study. <i>Oncology Research</i> , 2008, 17, 183-189.	1.6	21
122	High Ki-67 Immunohistochemical Reactivity Correlates With Poor Prognosis in Bladder Carcinoma. <i>Medicine (United States)</i> , 2016, 95, e3337.	1.1	21
123	Evaluation of the HOXA11 level in patients with lung squamous cancer and insights into potential molecular pathways via bioinformatics analysis. <i>World Journal of Surgical Oncology</i> , 2018, 16, 109.	1.9	21
124	Prognostic index of aberrant mRNA splicing profiling acts as a predictive indicator for hepatocellular carcinoma based on TCGA SpliceSeq data. <i>International Journal of Oncology</i> , 2019, 55, 425-438.	3.2	21
125	Role of alternative splicing signatures in the prognosis of glioblastoma. <i>Cancer Medicine</i> , 2019, 8, 7623-7636.	2.9	21
126	Clinical and genetic characteristics of female dystrophinopathy carriers. <i>Molecular Medicine Reports</i> , 2019, 19, 3035-3044.	2.5	21

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127	Pressure-and temperature induced phase transitions, piezochromism, NLC behaviour and pressure controlled Jahnâ€“Teller switching in a Cu-based framework. <i>Chemical Science</i> , 2020, 11, 8793-8799.	7.8	21
128	Ultrasound Imageâ€“Based Radiomics. <i>Journal of Ultrasound in Medicine</i> , 2021, 40, 1229-1244.	1.8	21
129	TNFRSF6B neutralization antibody inhibits proliferation and induces apoptosis in hepatocellular carcinoma cell. <i>Pathology Research and Practice</i> , 2010, 206, 631-641.	2.3	20
130	High expression of long nonâ€“coding HOTAIR correlated with hepatocarcinogenesis and metastasis. <i>Molecular Medicine Reports</i> , 2018, 17, 1148-1156.	2.5	20
131	Downregulation of HOXA3 in lung adenocarcinoma and its relevant molecular mechanism analysed by RT-qPCR, TCGA and in silico analysis. <i>International Journal of Oncology</i> , 2018, 53, 1557-1579.	3.2	20
132	Comprehensive evaluation of FKBP10 expression and its prognostic potential in gastric cancer. <i>Oncology Reports</i> , 2019, 42, 615-628.	2.6	20
133	The expression, significance and function of cancer susceptibility candidateâ€“19 in lung squamous cell carcinoma: A bioinformatics and inâ€“vitro investigation. <i>International Journal of Oncology</i> , 2019, 54, 1651-1664.	3.2	20
134	The Latest Overview of circRNA in the Progression, Diagnosis, Prognosis, Treatment, and Drug Resistance of Hepatocellular Carcinoma. <i>Frontiers in Oncology</i> , 2020, 10, 608257.	2.9	20
135	An autophagy-related gene expression signature for survival prediction in multiple cohorts of hepatocellular carcinoma patients. <i>Oncotarget</i> , 2018, 9, 17368-17395.	2.1	20
136	Expression of Tumor Necrosis Factor Receptor-associated Factor 6 in Lung Cancer Tissues. <i>Asian Pacific Journal of Cancer Prevention</i> , 2015, 15, 10591-10596.	1.2	20
137	Over-expression of Decoy Receptor 3 in Gastric Precancerous Lesions and Carcinoma. <i>Upsala Journal of Medical Sciences</i> , 2008, 113, 297-304.	0.9	19
138	Significance of decoy receptor 3 in sera of hepatocellular carcinoma patients. <i>Upsala Journal of Medical Sciences</i> , 2010, 115, 232-237.	0.9	19
139	Clinicopathological role of miR-30a-5p in hepatocellular carcinoma tissues and prediction of its function with bioinformatics analysis. <i>OncoTargets and Therapy</i> , 2016, Volume 9, 5061-5071.	2.1	19
140	Biological function of UCA1 in hepatocellular carcinoma and its clinical significance: Investigation with in vitro and meta-analysis. <i>Pathology Research and Practice</i> , 2018, 214, 1260-1272.	2.3	19
141	A metaâ€“analysis and bioinformatics exploration of the diagnostic value and molecular mechanism of miRâ€“193aâ€“5p in lung cancer. <i>Oncology Letters</i> , 2018, 16, 4114-4128.	1.8	19
142	Clinical value of miR-198-5p in lung squamous cell carcinoma assessed using microarray and RT-qPCR. <i>World Journal of Surgical Oncology</i> , 2018, 16, 22.	1.9	19
143	<p>MiR-193a-3p inhibits pancreatic ductal adenocarcinoma cell proliferation by targeting CCND1</p>. <i>Cancer Management and Research</i> , 2019, Volume 11, 4825-4837.	2.0	19
144	Prognostic value of small nucleolar RNAs (snoRNAs) for colon adenocarcinoma based on RNA sequencing data. <i>Pathology Research and Practice</i> , 2020, 216, 152937.	2.3	19

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