Gang Chen

List of Publications by Year
in descending order

Source: //exaly.com/author-pdf/4724337/publications.pdf

Version: 2024-02-01

308	8,837	60835	106894
papers	citations	43 h-index	65 g-index

376	376	376	13960
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Global nitrous oxide budget (1980â€“2020). Earth System Science Data, 2024, 16, 2543-2604.	8.9	0
2	Global bibliometric mapping of the research trends in artificial intelligence-based digital pathology for lung cancer over the past two decades. Digital Health, 2024, 10, .	1.8	0
3	Clinical Implication of E2F Transcription Factor 1 in Hepatocellular Carcinoma Tissues. Cancer Biotherapy and Radiopharmaceuticals, 2023, 38, 684-707.	1.1	7
4	Clinical Significance of Integrin Subunit Beta 4 in Head and Neck Squamous Cell Carcinoma. Cancer Biotherapy and Radiopharmaceuticals, 2022, 37, 256-275.	1.1	12
5	Downregulation of miR-125b-5p and Its Prospective Molecular Mechanism in Lung Squamous Cell Carcinoma. Cancer Biotherapy and Radiopharmaceuticals, 2022, 37, 125-140.	1.1	4
6	Laryngeal Squamous Cell Carcinoma: Clinical Significance and Potential Mechanism of Cell Division Cycle 45. Cancer Biotherapy and Radiopharmaceuticals, 2022, 37, 300-312.	1.1	5
7	Expression Landscape and Functional Roles of HOXA4 and HOXA5 in Lung Adenocarcinoma. International Journal of Medical Sciences, 2022, 19, 572-587.	2.6	3
8	Effect of CELSR3 on the Cell Cycle and Apoptosis of Hepatocellular Carcinoma Cells: Erratum. Journal of Cancer, 2022, 13, 1386-1387.	2.6	0
9	Downregulation of MicroRNA-1 and Its Potential Molecular Mechanism in Nasopharyngeal Cancer: An Investigation Combined with In Silico and In-House Immunohistochemistry Validation. Disease Markers, 2022, 2022, 1-13.	1.4	1
10	Expression of IER3 in hepatocellular carcinoma: clinicopathology, prognosis, and potential regulatory pathways. PeerJ, 2022, 10, e12944.	2.0	4
11	Decreased expression of transcription factor Homeobox A11 and its potential target genes in bladder cancer. Pathology Research and Practice, 2022, 233, 153847.	2.3	3
12	Ogt Demonstrated Conspicuous Clinical Significance in Cancers, from Pan-Cancer to Small-Cell Lung Cancer. Journal of Oncology, 2022, 2022, 1-16.	1.4	3
13	SYNJ2 is a novel and potential biomarker for the prediction and treatment of cancers: from lung squamous cell carcinoma to pan-cancer. BMC Medical Genomics, 2022, 15, 114.	1.5	4
14	Clinical significance of cyclin-dependent kinase inhibitor 2C expression in cancers: from small cell lung carcinoma to pan-cancers. BMC Pulmonary Medicine, 2022, 22, .	2.0	11
15	Comprehensive expression analysis reveals upregulated LUZP2 in prostate cancer tissues. Electronic Journal of Biotechnology, 2022, 59, 1-12.	2.3	0
16	Clinical assessment and molecular mechanism of the upregulation of Toll-like receptor 2 (TLR2) in myocardial infarction. BMC Cardiovascular Disorders, 2022, 22, .	1.7	6
17	Estrogenic activities of compound GL-1, isolated from Ganoderma lucidum. Natural Product Research, 2021, 35, 6062-6066.	1.8	2
18	Apparent Intracellular Helicobacter pylori Detected by Immunohistochemistry: The Missing Link in Eradication Failure. Clinical Infectious Diseases, 2021, 73, e1719-e1726.	5.7	8

#	Article	IF	Citations
19	Upregulation of ATIC in multiple myeloma tissues based on tissue microarray and gene microarrays. International Journal of Laboratory Hematology, 2021, 43, 409-417.	1.3	10
20	Ultrasound Imageâ€“Based Radiomics. Journal of Ultrasound in Medicine, 2021, 40, 1229-1244.	1.8	21
21	A single-cell polony method reveals low levels of infected Prochlorococcus in oligotrophic waters despite high cyanophage abundances. ISME Journal, 2021, 15, 41-54.	10.0	49
22	Measurement of hadronic event shapes in high-pT multijet final states at $$ sqrt{s} $$ = 13 TeV with the ATLAS detector. Journal of High Energy Physics, 2021, 2021, 1.	4.8	6
23	Clinical significance and molecular mechanism of angiotensin-converting enzyme 2 in hepatocellular carcinoma tissues. Bioengineered, 2021, 12, 4054-4069.	3.2	12
24	Clinical significance and potential molecular mechanism of miRNA-222-3p in metastatic prostate cancer. Bioengineered, 2021, 12, 325-340.	3.2	24
25	Expression and Clinical Significance of BCL2 Interacting Protein 3 Like in Multiple Myeloma. Technology in Cancer Research and Treatment, 2021, 20, 153303382110245.	1.9	1
26	Index-stable compact $${varvec{p}}$$-adic analytic groups. Archiv Der Mathematik, 2021, 116, 153-160.	0.5	1
27	MiRNAâ€145â€5p expression and prospective molecular mechanisms in the metastasis of prostate cancer. IET Systems Biology, 2021, 15, 1-13.	1.5	8
28	Financial hardship and health risk behavior during COVID-19 in a large US national sample of women. SSM - Population Health, 2021, 13, 100734.	2.9	42
29	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
30	Development and Validation of a Radiomic Nomogram for Predicting the Prognosis of Kidney Renal Clear Cell Carcinoma. Frontiers in Oncology, 2021, 11, 613668.	2.9	8
31	LPCAT1 overexpression promotes the progression of hepatocellular carcinoma. Cancer Cell International, 2021, 21, 442.	4.3	27
32	The Indication of Poor Prognosis by High Expression of ENO1 in Squamous Cell Carcinoma of the Lung. Journal of Oncology, 2021, 2021, 1-11.	1.4	4
33	Clinical Significance and Underlying Mechanisms of CELSR3 in Metastatic Prostate Cancer Based on Immunohistochemistry, Data Mining, and In Silico Analysis. Cancer Biotherapy and Radiopharmaceuticals, 2021, , .	1.1	0
34	Investigation on the structural quality dependent electromagnetic interference shielding performance of few-layer and lamellar Nb2CTx MXene nanostructures. Journal of Alloys and Compounds, 2021, 877, 160235.	5.7	23
35	Incomplete thermal ablation-induced up-regulation of transcription factor nuclear receptor subfamily 2, group F, member 6 (NR2F6) contributes to the rapid progression of residual liver tumor in hepatoblastoma. Bioengineered, 2021, 12, 4289-4303.	3.2	4
36	Upregulation of microRNA miR-141-3p and its prospective targets in endometrial carcinoma: a comprehensive study. Bioengineered, 2021, 12, 2941-2956.	3.2	12

#	Article	IF	Citations
37	Identification of a novel therapeutic candidate, NRK, in primary cancer-associated fibroblasts of lung adenocarcinoma microenvironment. Journal of Cancer Research and Clinical Oncology, 2021, 147, 1049-1064.	2.6	6
38	Down-regulation of microRNA-125b-2-3p is a risk factor for a poor prognosis in hepatocellular carcinoma. Bioengineered, 2021, 12, 1627-1641.	3.2	11
39	The Use of Cangrelor Infusions After Endovascular Aortic Repair With Prophylactic Lumbar Drain Placement. Journal of Cardiothoracic and Vascular Anesthesia, 2021, 35, 3723-3726.	1.3	3
40	Expression of Cell Division Cycle Protein 45 in Tissue Microarrays and the CDC45 Gene by Bioinformatics Analysis in Human Hepatocellular Carcinoma and Patient Outcomes. Medical Science Monitor, 2021, 27, e928800.	1.1	11
41	Clinical significance and effect of lncRNA BBOX1â€‘AS1 on the proliferation and migration of lung squamous cell carcinoma. Oncology Letters, 2021, 23, 17.	1.8	6
42	Identification of the susceptibility genes for COVID-19 in lung adenocarcinoma with global data and biological computation methods. Computational and Structural Biotechnology Journal, 2021, 19, 6229-6239.	4.2	12
43	Down-Regulation of Activating Transcription Factor 3 (ATF3) in Hepatoblastoma and Its Relationship with Ferroptosis. International Journal of General Medicine, 2021, Volume 14, 9401-9418.	1.8	3
44	A novel risk signature that combines 10 long noncoding RNAs to predict neuroblastoma prognosis. Journal of Cellular Physiology, 2020, 235, 3823-3834.	4.2	15
45	A radiogenomics signature for predicting the clinical outcome of bladder urothelial carcinoma. European Radiology, 2020, 30, 547-557.	4.6	44
46	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
47	The role of upregulated miR-375 expression in breast cancer: An in vitro and in silico study. Pathology Research and Practice, 2020, 216, 152754.	2.3	28
48	The clinical significance and potential molecular mechanism of integrin subunit beta 4 in laryngeal squamous cell carcinoma. Pathology Research and Practice, 2020, 216, 152785.	2.3	8
49	Clinical significance of transcription factor RUNX2 in lung adenocarcinoma and its latent transcriptional regulating mechanism. Computational Biology and Chemistry, 2020, 89, 107383.	2.4	13
50	Clinical significance of CCNE2 protein and mRNA expression in thyroid cancer tissues. Advances in Medical Sciences, 2020, 65, 442-456.	2.2	9
51	Clinical significance and biological function of transcriptional repressor GATA binding 1 in gastric cancer: a study based on data mining, RT-qPCR, immunochemistry, and vitro experiment. Cell Cycle, 2020, 19, 2866-2885.	2.8	6
52	Downregulation of miRNA-205 Expression and Biological Mechanism in Prostate Cancer Tumorigenesis and Bone Metastasis. BioMed Research International, 2020, 2020, 1-17.	2.0	12
53	Pressure-and temperature induced phase transitions, piezochromism, NLC behaviour and pressure controlled Jahnâ€“Teller switching in a Cu-based framework. Chemical Science, 2020, 11, 8793-8799.	7.8	21
54	Prognostic Values for the mRNA Expression of the ADAMTS Family of Genes in Gastric Cancer. Journal of Oncology, 2020, 2020, 1-24.	1.4	15

#	Article	IF	Citations
55	Clinical value and potential mechanisms of COL8A1 upregulation in breast cancer: a comprehensive analysis. Cancer Cell International, 2020, 20, 392.	4.3	22
56	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
57	Time-restricted feeding alters lipid and amino acid metabolite rhythmicity without perturbing clock gene expression. Nature Communications, 2020, 11, 4643.	13.2	80
58	Immunohistochemical basigin expression level in thyroid cancer tissues. World Journal of Surgical Oncology, 2020, 18, 240.	1.9	2
59	The Expression and Potential Role of Tubulin Alpha 1b in Wilmsâ€™ Tumor. BioMed Research International, 2020, 2020, 1-10.	2.0	10
60	Downregulation of miR-199a-3p in Hepatocellular Carcinoma and Its Relevant Molecular Mechanism via GEO, TCGA Database and In Silico Analyses. Technology in Cancer Research and Treatment, 2020, 19, 153303382097967.	1.9	6
61	Predictive value of hypoxia, metabolism and immune factors for prognosis in hepatocellular carcinoma: a retrospective analysis and multicenter validation study. Journal of Cancer, 2020, 11, 4145-4156.	2.6	5
62	Calcaneus traction compression with orthopaedic reduction forceps combined with percutaneous minimally invasive treatment of intra-articular calcaneal fractures: An analysis of efficacy. Biomedicine and Pharmacotherapy, 2020, 128, 110295.	5.8	5
63	The clinical significance of interleukin 24 and its potential molecular mechanism in laryngeal squamous cell carcinoma. Cancer Biomarkers, 2020, 29, 111-124.	1.7	6
64	Erratum to â€œThe role of upregulated miR-375 expression in breast cancer: An in vitro and in silico studyâ€•[Pathol. Res. Pract. 216 (January (1)) (2020) 152754]. Pathology Research and Practice, 2020, 216, 152929.	2.3	0
65	Downregulation of hsa-microRNA-204-5p and identification of its potential regulatory network in non-small cell lung cancer: RT-qPCR, bioinformatic- and meta-analyses. Respiratory Research, 2020, 21, 60.	3.7	11
66	Radiomic profiles in diffuse glioma reveal distinct subtypes with prognostic value. Journal of Cancer Research and Clinical Oncology, 2020, 146, 1253-1262.	2.6	17
67	Development and validation of an immune prognostic classifier for clear cell renal cell carcinoma. Cancer Biomarkers, 2020, 27, 265-275.	1.7	7
68	Clinical Significance and Effect of MTDH/AEG-I in Bladder Urothelial Cancer: A Study Based on Immunohistochemistry, RNA-Seq, and in vitro Verification [Retraction] . Cancer Management and Research, 2020, Volume 12, 461-462.	2.0	0
69	Effect of CELSR3 on the Cell Cycle and Apoptosis of Hepatocellular Carcinoma Cells. Journal of Cancer, 2020, 11, 2830-2844.	2.6	9
70	Clinicopathological value and underlying molecular mechanism of annexin A2 in 992 cases of thyroid carcinoma. Computational Biology and Chemistry, 2020, 86, 107258.	2.4	6
71	Prognostic value of small nucleolar RNAs (snoRNAs) for colon adenocarcinoma based on RNA sequencing data. Pathology Research and Practice, 2020, 216, 152937.	2.3	19
72	The Clinical Significance and Potential Molecular Mechanism of PTTG1 in Esophageal Squamous Cell Carcinoma. Frontiers in Genetics, 2020, 11, 583085.	2.3	11

#	Article	IF	Citations
73	The Latest Overview of circRNA in the Progression, Diagnosis, Prognosis, Treatment, and Drug Resistance of Hepatocellular Carcinoma. Frontiers in Oncology, 2020, 10, 608257.	2.9	20
74	Integrated expression analysis revealed RUNX2 upregulation in lung squamous cell carcinoma tissues. IET Systems Biology, 2020, 14, 252-260.	1.5	8
75	RNA-Sequencing, Connectivity Mapping, and Molecular Docking to Investigate Ligand-Protein Binding for Potential Drug Candidates for the Treatment of Wilms Tumor. Medical Science Monitor, 2020, 26, e920725.	1.1	3
76	Identification of an Immune Score-Based Gene Panel with Prognostic Power for Oral Squamous Cell Carcinoma. Medical Science Monitor, 2020, 26, e922854.	1.1	18
77	Downregulation of CDC14B in 5218 breast cancer patients: A novel prognosticator for triple-negative breast cancer. Mathematical Biosciences and Engineering, 2020, 17, 8152-8181.	2.0	5
78	Genome-wide Analysis of the Alternative Splicing Profiles Revealed Novel Prognostic Index for Kidney Renal Cell Clear Cell Carcinoma. Journal of Cancer, 2020, 11, 1542-1554.	2.6	4
79	Downregulation of miR-193a-3p is involved in the pathogenesis of hepatocellular carcinoma by targeting CCND1. PeerJ, 2020, 8, e8409.	2.0	14
80	Upregulated expression of SAC3D1 is associated with progression in gastric cancer. International Journal of Oncology, 2020, 57, 122-138.	3.2	5
81	The clinical significance of apolipoprotein L1 in head and neck squamous cell carcinoma. Oncology Letters, 2020, 20, 1-1.	1.8	9
82	Small Nucleolar RNAs (snoRNAs)-Based Risk Score Classifier Predicts Overall Survival in Bladder Carcinoma. Medical Science Monitor, 2020, 26, e926273.	1.1	2
83	Investigation of the clinical significance and molecular mechanism of miRâ€'21â€'5p in hepatocellular carcinoma: A systematic review based on 24 studies and bioinformatics investigation. Oncology Letters, 2019, 17, 230-246.	1.8	9
84	Oncogenic value of microRNAâ€'15bâ€'5p in hepatocellular carcinoma and a bioinformatics investigation. Oncology Letters, 2019, 17, 1695-1713.	1.8	17
85	Expression levels and coâ€'targets of miRNAâ€'126â€'3p and miRNAâ€'126â€'5p in lung adenocarcinoma tissues: Îˆn exploration with RTâ€'qPCR, microarray and bioinformatic analyses. Oncology Reports, 2019, 41, 939-953.	2.6	15
86	Expression of miRâ€'542â€'3p in osteosarcoma with miRNA microarray data, and its potential signaling pathways. Molecular Medicine Reports, 2019, 19, 974-983.	2.5	3
87	Identification of putative drugs for gastric adenocarcinoma utilizing differentially expressed genes and connectivity map. Molecular Medicine Reports, 2019, 19, 1004-1015.	2.5	3
88	Clinical value of microRNAâ€'198â€'5p downregulation in lung adenocarcinoma and its potential pathways. Oncology Letters, 2019, 18, 2939-2954.	1.8	12
89	Expression significance and potential mechanism of hypoxiaâ€inducible factor 1 alpha in patients with myelodysplastic syndromes. Cancer Medicine, 2019, 8, 6021-6035.	2.9	11
90	Expression and clinical significance of neuropilin-1 in Epstein-Barr virus-associated lymphomas. Cancer Biomarkers, 2019, 25, 259-273.	1.7	5

#	Article	IF	Citations
91	Profiling of prognostic alternative splicing in melanoma. Oncology Letters, 2019, 18, 1081-1088.	1.8	8
92	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
93	Protective potential of miR-146a-5p and its underlying molecular mechanism in diverse cancers: a comprehensive meta-analysis and bioinformatics analysis. Cancer Cell International, 2019, 19, 167.	4.3	12
94	Prognostic index of aberrant mRNA splicing profiling acts as a predictive indicator for hepatocellular carcinoma based on TCGA SpliceSeq data. International Journal of Oncology, 2019, 55, 425-438.	3.2	21
95	Clinical Significance of microRNA-196b-5p in Hepatocellular Carcinoma and its Potential Molecular Mechanism. Journal of Cancer, 2019, 10, 5355-5370.	2.6	12
96	Role of alternative splicing signatures in the prognosis of glioblastoma. Cancer Medicine, 2019, 8, 7623-7636.	2.9	21
97	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. Journal of Cancer, 2019, 10, 5339-5354.	2.6	31
98	Differentially expressed gene profile and relevant pathways of the traditional Chinese medicine cinobufotalin on MCFâ€7 breast cancer cells. Molecular Medicine Reports, 2019, 19, 4256-4270.	2.5	8
99	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
100	Prognosis of clear cell renal cell carcinoma (ccRCC) based on a six-lncRNA-based risk score: an investigation based on RNA-sequencing data. Journal of Translational Medicine, 2019, 17, 281.	4.5	36
101	Identification of potential agents for thymoma by integrated analyses of differentially expressed tumourâ€associated genes and molecular docking experiments. Experimental and Therapeutic Medicine, 2019, 18, 2001-2014.	1.9	2
102	High throughput circRNA sequencing analysis reveals novel insights into the mechanism of nitidine chloride against hepatocellular carcinoma. Cell Death and Disease, 2019, 10, 658.	6.4	55
103	CD117 expression is correlated with poor survival of patients and progression of lung carcinoma:a meta-analysis with a panel of 2645 patients. Polish Journal of Pathology, 2019, 70, 63-78.	0.4	4
104	Identification and validation of an individualized autophagy-clinical prognostic index in bladder cancer patients . OncoTargets and Therapy, 2019, Volume 12, 3695-3712.	2.1	37
105	MiR-193a-3p inhibits pancreatic ductal adenocarcinoma cell proliferation by targeting CCND1 . Cancer Management and Research, 2019, Volume 11, 4825-4837.	2.0	19
106	Comprehensive evaluation of FKBP10 expression and its prognostic potential in gastric cancer. Oncology Reports, 2019, 42, 615-628.	2.6	20
107	Evaluation of miR-302b-5p expression and molecular mechanism in hepatocellular carcinoma: Findings based on RT-qPCR and in silico analysis. Pathology Research and Practice, 2019, 215, 152424.	2.3	9
108	Gene profiling of HepG2 cells following nitidine chloride treatment: An investigation with microarray and Connectivity Mapping. Oncology Reports, 2019, 41, 3244-3256.	2.6	9

#	Article	IF	Citations
109	The expression, significance and function of cancer susceptibility candidateï¿½9 in lung squamous cell carcinoma: A bioinformatics and inï¿½vitro investigation. International Journal of Oncology, 2019, 54, 1651-1664.	3.2	20
110	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. Respiratory Research, 2019, 20, 48.	3.7	47
111	Expression of vimentin in nasopharyngeal carcinoma and its possible molecular mechanism: A study based on immunohistochemistry and bioinformatics analysis. Pathology Research and Practice, 2019, 215, 1020-1032.	2.3	9
112	Clinical and genetic characteristics of female dystrophinopathy carriers. Molecular Medicine Reports, 2019, 19, 3035-3044.	2.5	21
113	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
114	Inï¿½silico analysis identified miRNAâ€˜based therapeutic agents against glioblastoma multiforme. Oncology Reports, 2019, 41, 2194-2208.	2.6	30
115	miRâ€˜146aâ€˜5p targets TCSF and influences cell growth and apoptosis to repress NSCLC progression. Oncology Reports, 2019, 41, 2226-2240.	2.6	18
116	Novel drug candidate for the treatment of several softâ€˜tissue sarcoma histologic subtypes: A computational method using survivalâ€˜associated gene signatures for drug repurposing. Oncology Reports, 2019, 41, 2241-2253.	2.6	9
117	Drug repositioning in head and neck squamous cell carcinoma: An integrated pathway analysis based on connectivity map and differential gene expression. Pathology Research and Practice, 2019, 215, 152378.	2.3	13
118	The clinical significance of endothelin receptor type B in hepatocellular carcinoma and its potential molecular mechanism. Experimental and Molecular Pathology, 2019, 107, 141-157.	2.3	23
119	The coexistence of a Wilms’ tumor and renal cell carcinoma in children: a case report and review of the literature . OncoTargets and Therapy, 2019, Volume 12, 953-958.	2.1	5
120	Research about the Integration of Urban and Rural Public Transportation and Rizhaoâ€™s Reform. , 2019, , .		1
121	Influence of sodium diffusion from substrates on performance of SnS/CdS thin-film solar cells. Journal of Materials Chemistry A, 2019, 7, 24186-24190.	10.5	11
122	MIR22HG As A Tumor Suppressive lncRNA In HCC: A Comprehensive Analysis Integrating RT-qPCR, mRNA-Seq, And Microarrays . OncoTargets and Therapy, 2019, Volume 12, 9827-9848.	2.1	16
123	Upregulated LINC00565 Accelerates Ovarian Cancer Progression By Targeting GAS6 . OncoTargets and Therapy, 2019, Volume 12, 10011-10022.	2.1	18
124	Effective Thermoelasticity of Polymer-Bonded Particle Composites with Imperfect Interfaces and Thermally Expansive Interphases. Journal of Elasticity, 2019, 136, 55-85.	2.0	9
125	Clinical and prognostic value of chaperonin containing T-complex 1 subunit 3 in hepatocellular carcinoma: A Study based on microarray and RNA-sequencing with 4272 cases. Pathology Research and Practice, 2019, 215, 177-194.	2.3	15
126	Prognostic value of small nuclear RNAs (snRNAs) for digestive tract pan- adenocarcinomas identified by RNA sequencing data. Pathology Research and Practice, 2019, 215, 414-426.	2.3	14

#	Article	IF	Citations
127	EBV as a potential risk factor for hepatobiliary system cancer: A meta-analysis with 918 cases. Pathology Research and Practice, 2019, 215, 278-285.	2.3	9
128	An air freight forwarderâ€™s resource planning and revenue management. Journal of the Operational Research Society, 2019, 70, 294-309.	3.4	6
129	Improvements in the process efficiency and bond strength when friction surfacing stainless steel onto aluminium substrates. Proceedings of the Institution of Mechanical Engineers, Part L: Journal of Materials: Design and Applications, 2019, 233, 687-698.	1.0	6
130	Primitive neuroectodermal tumors of the abdominal wall and vulva in children: Report of two cases and review of the literature. World Journal of Clinical Cases, 2019, 7, 3671-3682.	0.8	4
131	A novel alternative splicing-based prediction model for uteri corpus endometrial carcinoma. Aging, 2019, 11, 263-283.	2.9	16
132	Diagnostic significance and potential function of miR-338-5p in hepatocellular carcinoma: A bioinformatics study with microarray and RNA sequencing data. Molecular Medicine Reports, 2018, 17, 2297-2312.	2.5	11
133	Utility of miRâ€‘133aâ€‘3p as a diagnostic indicator for hepatocellular carcinoma: An investigation combined with GEO, TCGA, metaâ€‘analysis and bioinformatics. Molecular Medicine Reports, 2018, 17, 1469-1484.	2.5	24
134	Prognostic microRNAs and their potential molecular mechanism in pancreatic cancer: A study based on The Cancer Genome Atlas and bioinformatics investigation. Molecular Medicine Reports, 2018, 17, 939-951.	2.5	40
135	Potential role of microRNAâ€‘223â€‘3p in the tumorigenesis of hepatocellular carcinoma: A comprehensive study based on data mining and bioinformatics. Molecular Medicine Reports, 2018, 17, 2211-2228.	2.5	10
136	High expression of long nonâ€‘coding HOTAIR correlated with hepatocarcinogenesis and metastasis. Molecular Medicine Reports, 2018, 17, 1148-1156.	2.5	20
137	Downregulated miR-23b-3p expression acts as a predictor of hepatocellular carcinoma progression: A study based on public data and RT-qPCR verification. International Journal of Molecular Medicine, 2018, 41, 2813-2831.	4.1	38
138	Role of upregulated miR-136-5p in lung adenocarcinoma: A study of 1242 samples utilizing bioinformatics analysis. Pathology Research and Practice, 2018, 214, 750-766.	2.3	13
139	Expression level and potential target pathways of miR-1-3p in colorectal carcinoma based on 645Ã¯Â¿Â½cases from 9 microarray datasets. Molecular Medicine Reports, 2018, 17, 5013-5020.	2.5	24
140	MicroRNA-671-3p inhibits the development of breast cancer: A study based on in vitro experiments, in-house quantitative polymerase chain reaction and bioinformatics analysis. International Journal of Oncology, 2018, 52, 1801-1814.	3.2	6
141	Biological function of UCA1 in hepatocellular carcinoma and its clinical significance: Investigation with in vitro and meta-analysis. Pathology Research and Practice, 2018, 214, 1260-1272.	2.3	19
142	In silico analysis of the potential mechanism of telocinobufagin on breast cancer MCF-7 cells. Pathology Research and Practice, 2018, 214, 631-643.	2.3	8
143	A comprehensive analysis of the predicted targets of miR-642b-3p associated with the long non-coding RNA HOXA11-AS in NSCLC cells. Oncology Letters, 2018, 15, 6147-6160.	1.8	17
144	Clinical significance of high expression of miR-452-5p in lung squamous cell carcinoma. Oncology Letters, 2018, 15, 6418-6430.	1.8	37

#	Article	IF	Citations
145	Investigation of miR-136-5p key target genes and pathways in lung squamous cell cancer based on TCGA database and bioinformatics analysis. Pathology Research and Practice, 2018, 214, 644-654.	2.3	39
146	Clinical value of survivin and its underlying mechanism in ovarian cancer: A bioinformatics study based on GEO and TCGA data mining. Pathology Research and Practice, 2018, 214, 385-401.	2.3	12
147	Role of miR-1 expression in clear cell renal cell carcinoma (ccRCC): A bioinformatics study based on GEO, ArrayExpress microarrays and TCGA database. Pathology Research and Practice, 2018, 214, 195-206.	2.3	4
148	Clinical significance of microRNA-449a in hepatocellular carcinoma with microarray data mining together with initial bioinformatics analysis. Experimental and Therapeutic Medicine, 2018, 15, 3247-3258.	1.9	2
149	MicroRNAâ€'124â€'3p expression and its prospective functional pathways in hepatocellular carcinoma: A quantitative polymerase chain reaction, gene expression omnibus and bioinformatics study. Oncology Letters, 2018, 15, 5517-5532.	1.8	22
150	Asynchronous Multi-sensor Data Fusion with Decentralized IMM-PDAF. , 2018, , .		2
151	Clinical significance and effect of MTDH/AEG-1 in bladder urothelial cancer: a study based on immunohistochemistry, RNA-seq, and in vitro verification. Cancer Management and Research, 2018, Volume 10, 6921-6936.	2.0	1
152	Health Status Benefits of SuccessfulÂChronic Total Occlusion Revascularization Across the SpectrumÂofÂLeft Ventricular Function. JACC: Cardiovascular Interventions, 2018, 11, 2276-2283.	3.6	11
153	Expression and potential molecular mechanisms of miRâ€'204â€'5p in breast cancer, based on bioinformatics and a metaâ€'analysis of 2,306ï¿½cases. Molecular Medicine Reports, 2018, 19, 1168-1184.	2.5	4
154	Genomic analysis of small nucleolar RNAs identifies distinct molecular and prognostic signature in hepatocellular carcinoma. Oncology Reports, 2018, 40, 3346-3358.	2.6	16
155	A metaâ€'analysis and bioinformatics exploration of the diagnostic value and molecular mechanism of miRâ€'193aâ€'5p in lung cancer. Oncology Letters, 2018, 16, 4114-4128.	1.8	19
156	Upregulation of HOXA11 during the progression of lung adenocarcinoma detected via multiple approaches. International Journal of Molecular Medicine, 2018, 42, 2650-2664.	4.1	30
157	Exploration of the diagnostic value and molecular mechanism of miRâ€'1 in prostate cancer: A study based on metaâ€'analyses and bioinformatics. Molecular Medicine Reports, 2018, 18, 5630-5646.	2.5	14
158	Survival analysis of genome-wide profiles coupled with Connectivity Map database mining to identify potential therapeutic targets for cholangiocarcinoma. Oncology Reports, 2018, 40, 3189-3198.	2.6	6
159	Downregulation of miRâ€'224â€'5p in prostate cancer and its relevant molecular mechanism via TCGA, GEO database and inï¿½silico analyses. Oncology Reports, 2018, 40, 3171-3188.	2.6	34
160	DNA topoisomerase 1 and 2A function as oncogenes in liver cancer and may be direct targets of nitidine chloride. International Journal of Oncology, 2018, 53, 1897-1912.	3.2	55
161	Identification of potential drugs for diffuse large b-cell lymphoma based on bioinformatics and Connectivity Map database. Pathology Research and Practice, 2018, 214, 1854-1867.	2.3	17
162	Upregulation of HOXA1 promotes tumorigenesis and development of nonâ€'small cell lung cancer: A comprehensive investigation based on reverse transcription-quantitative polymerase chain reaction and bioinformatics analysis. International Journal of Oncology, 2018, 53, 73-86.	3.2	18

#	Article	IF	Citations
163	Microarrayâ€‘based bioinformatics analysis of the prospective target gene network of key miRNAs influenced by long nonâ€‘coding RNA PVT1 in HCC. Oncology Reports, 2018, 40, 226-240.	2.6	11
164	Regulatory interactions between long noncoding RNA LINC00968 and miRâ€‘9â€‘3p in nonâ€‘small cell lung cancer: A bioinformatic analysis based on miRNA microarray, GEO and TCGA. Oncology Letters, 2018, 15, 9487-9497.	1.8	9
165	Downregulation of miRâ€‘486â€‘5p in papillary thyroid carcinoma tissue: A study based on microarray and miRNA sequencing. Molecular Medicine Reports, 2018, 18, 2631-2642.	2.5	14
166	Comprehensive analysis of the clinical significance and prospective molecular mechanisms of differentially expressed autophagy-related genes in thyroid cancer. International Journal of Oncology, 2018, 53, 603-619.	3.2	14
167	Evaluation of the HOXA11 level in patients with lung squamous cancer and insights into potential molecular pathways via bioinformatics analysis. World Journal of Surgical Oncology, 2018, 16, 109.	1.9	21
168	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
169	Clinical value of miR-182-5p in lung squamous cell carcinoma: a study combining data from TCGA, GEO, and RT-qPCR validation. World Journal of Surgical Oncology, 2018, 16, 76.	1.9	28
170	Clinical value of miR-198-5p in lung squamous cell carcinoma assessed using microarray and RT-qPCR. World Journal of Surgical Oncology, 2018, 16, 22.	1.9	19
171	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. Medical Science Monitor, 2018, 24, 2786-2808.	1.1	52
172	Expression of microRNA-99a-3p in Prostate Cancer Based on Bioinformatics Data and Meta-Analysis of a Literature Review of 965 Cases. Medical Science Monitor, 2018, 24, 4807-4822.	1.1	5
173	A Network Pharmacology-Based Analysis of Multi-Target, Multi-Pathway, Multi-Compound Treatment for Ovarian Serous Cystadenocarcinoma. Clinical Drug Investigation, 2018, 38, 909-925.	2.2	12
174	A circRNAâ€“miRNAâ€“mRNA network identification for exploring underlying pathogenesis and therapy strategy of hepatocellular carcinoma. Journal of Translational Medicine, 2018, 16, 220.	4.5	242
175	Long nonâ€‘coding RNAs in small cell lung cancer: A potential opening to combat the disease (Review). Oncology Reports, 2018, 40, 1831-1842.	2.6	10
176	Systematic Analysis of Survival-Associated Alternative Splicing Signatures in Gastrointestinal Pan-Adenocarcinomas. EBioMedicine, 2018, 34, 46-60.	6.0	84
177	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. Respiratory Research, 2018, 19, 150.	3.7	50
178	Downregulation of HOXA3 in lung adenocarcinoma and its relevant molecular mechanism analysed by RT-qPCR, TCGA and in silico analysis. International Journal of Oncology, 2018, 53, 1557-1579.	3.2	20
179	The expression of HOXA13 in lung adenocarcinoma and its clinical significance: A study based on The Cancer Genome Atlas, Oncomine and reverse transcriptionâ€‘quantitative polymerase chain reaction. Oncology Letters, 2018, 15, 8556-8572.	1.8	23
180	Oncogenic role of miRâ€‘183â€‘5p in lung adenocarcinoma: A comprehensive study of qPCR, inÃ¯Â¿Â½vitro experiments and bioinformatic analysis. Oncology Reports, 2018, 40, 83-100.	2.6	22

#	Article	IF	Citations
181	Osteosarcopenic obesity and its relationship with dyslipidemia in women from different ethnic groups of China. Archives of Osteoporosis, 2018, 13, 65.	2.6	27
182	Investigation of miR-490-3p Expression in Hepatocellular Carcinoma Based on Reverse Transcription-Polymerase Chain Reaction (RT-qPCR) and a Meta-Analysis of 749 Cases. Medical Science Monitor, 2018, 24, 4914-4925.	1.1	8
183	A meta-analysis of the lymphatic microvessel density and survival in gastric cancer with 1809 cases. Oncotarget, 2018, 9, 5406-5415.	2.1	4
184	Caspase-3 over-expression is associated with poor overall survival and clinicopathological parameters in breast cancer: a meta-analysis of 3091 cases. Oncotarget, 2018, 9, 8629-8641.	2.1	30
185	An autophagy-related gene expression signature for survival prediction in multiple cohorts of hepatocellular carcinoma patients. Oncotarget, 2018, 9, 17368-17395.	2.1	20
186	Clinical significances of p27 in digestive tract cancers: a comprehensive analysis on immunohistochemistry staining, published literatures, microarray and RNA-seq data. Oncotarget, 2018, 9, 12284-12303.	2.1	3
187	Expression and clinical significance of ubiquitinâ€'specificâ€'processing protease 34 in diffuse large Bâ€'cell lymphoma. Molecular Medicine Reports, 2018, 18, 4543-4554.	2.5	8
188	Implication of downregulation and prospective pathway signaling of microRNA-375 in lung squamous cell carcinoma. Pathology Research and Practice, 2017, 213, 364-372.	2.3	29
189	Long non-coding RNA TUC338 is functionally involved in sorafenib-sensitized hepatocarcinoma cells by targeting RASAL1. Oncology Reports, 2017, 37, 273-280.	2.6	59
190	A qRT-PCR and Gene Functional Enrichment Study Focused on Downregulation of miR-141-3p in Hepatocellular Carcinoma and Its Clinicopathological Significance. Technology in Cancer Research and Treatment, 2017, 16, 835-849.	1.9	12
191	Clinical value of miR-452-5p expression in lung adenocarcinoma: A retrospective quantitative real-time polymerase chain reaction study and verification based on The Cancer Genome Atlas and Gene Expression Omnibus databases. Tumor Biology, 2017, 39, 101042831770575.	1.7	5
192	Transshipment hub selection from a shipper's and freight forwarder's perspective. Expert Systems With Applications, 2017, 83, 396-404.	7.9	27
193	A nine-miRNA signature as a potential diagnostic marker for breast carcinoma: An integrated study of 1,110 cases. Oncology Reports, 2017, 37, 3297-3304.	2.6	51
194	miR-204 regulates the biological behavior of breast cancer MCF-7 cells by directly targeting FOXA1. Oncology Reports, 2017, 38, 368-376.	2.6	44
195	Clinical value of miR-145-5p in NSCLC and potential molecular mechanism exploration: A retrospective study based on GEO, qRT-PCR, and TCGA data. Tumor Biology, 2017, 39, 101042831769168.	1.7	18
196	Identification of a serum microRNA expression signature for detection of lung cancer, involving miR-23b, miR-221, miR-148b and miR-423-3p. Lung Cancer, 2017, 114, 6-11.	2.0	68
197	Downregulation of miR-136-5p in hepatocellular carcinoma and its clinicopathological significance. Molecular Medicine Reports, 2017, 16, 5393-5405.	2.5	30
198	Morphological characteristics of fatal pediatric hand, foot and mouth disease: A clinicopathological study with related receptors of EV71. Pathology Research and Practice, 2017, 213, 1144-1151.	2.3	6

#	Article	IF	Citations
199	Clinical Significance and Effect of lncRNA HOXA11-AS in NSCLC: A Study Based on Bioinformatics, In Vitro and in Vivo Verification. Scientific Reports, 2017, 7, 5567.	3.4	47
200	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. International Journal of Oncology, 2017, 51, 1705-1721.	3.2	36
201	Down-regulation of miR-26a-5p in hepatocellular carcinoma: A qRT-PCR and bioinformatics study. Pathology Research and Practice, 2017, 213, 1494-1509.	2.3	22
202	Quantitative Analysis of Hepatic Microcirculation inÂRabbitsÂAfter Liver Ischemia-Reperfusion InjuryÂUsingÂContrast-Enhanced Ultrasound. Ultrasound in Medicine and Biology, 2017, 43, 2469-2476.	1.6	8
203	Longitudinal monitoring of heartbeat dynamics predicts mood changes in bipolar patients: A pilot study. Journal of Affective Disorders, 2017, 209, 30-38.	4.2	23
204	Cervical Cancer Growth Is Regulated by a c-ABLâ€“PLK1 Signaling Axis. Cancer Research, 2017, 77, 1142-1154.	0.9	36
205	Genetic analysis of the dystrophin gene in children with Duchenne and Becker muscular dystrophies. Muscle and Nerve, 2017, 56, 117-121.	2.3	8
206	Effect of miR-146a-5p on tumor growth in NSCLC using chick chorioallantoic membrane assay and bioinformatics investigation. Molecular Medicine Reports, 2017, 16, 8781-8792.	2.5	12
207	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
208	The diagnostic and prognostic values of Ki-67/MIB-1 expression in thyroid cancer: a meta-analysis with 6,051 cases. OncoTargets and Therapy, 2017, Volume 10, 3261-3276.	2.1	37
209	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. Medical Science Monitor, 2017, 23, 2453-2464.	1.1	42
210	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
211	The suppressive role of miR-542-5p in NSCLC: the evidence from clinical data and in vivo validation using a chick chorioallantoic membrane model. BMC Cancer, 2017, 17, 655.	2.6	42
212	The clinicopathological significance of UBE2C in breast cancer: a study based on immunohistochemistry, microarray and RNA-sequencing data. Cancer Cell International, 2017, 17, 83.	4.3	56
213	Comprehensive investigation of a novel differentially expressed lncRNA expression profile signature to assess the survival of patients with colorectal adenocarcinoma. Oncotarget, 2017, 8, 16811-16828.	2.1	96
214	RNA-sequencing investigation identifies an effective risk score generated by three novel lncRNAs for the survival of papillary thyroid cancer patients. Oncotarget, 2017, 8, 74139-74158.	2.1	28
215	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. International Journal of Molecular Medicine, 2017, 40, 1895-1906.	4.1	38
216	A comprehensive insight into the clinicopathologic significance of miR-144-3p in hepatocellular carcinoma. OncoTargets and Therapy, 2017, Volume 10, 3405-3419.	2.1	28

#	Article	IF	Citations
217	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
218	Role of downregulated miR-133a-3p expression in bladder cancer: a bioinformatics study. OncoTargets and Therapy, 2017, Volume 10, 3667-3683.	2.1	30
219	Progression-free survival of up to 8 months of an advanced intrahepatic cholangiocarcinoma patient treated with apatinib: a case report. OncoTargets and Therapy, 2017, Volume 10, 5237-5242.	2.1	5
220	Clinical roles of the aberrantly expressed lncRNAs in lung squamous cell carcinoma: a study based on RNA-sequencing and microarray data mining. Oncotarget, 2017, 8, 61282-61304.	2.1	75
221	Identification of a RNA-Seq based prognostic signature with five lncRNAs for lung squamous cell carcinoma. Oncotarget, 2017, 8, 50761-50773.	2.1	53
222	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. Oncotarget, 2017, 8, 35681-35699.	2.1	32
223	The anticipating value of PLK1 for diagnosis, progress and prognosis and its prospective mechanism in gastric cancer: a comprehensive investigation based on high-throughput data and immunohistochemical validation. Oncotarget, 2017, 8, 92497-92521.	2.1	18
224	The clinical value of lncRNA NEAT1 in digestive system malignancies: A comprehensive investigation based on 57 microarray and RNA-seq datasets. Oncotarget, 2017, 8, 17665-17683.	2.1	27
225	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
226	Clinical significance and effect of AEG-1 on the proliferation, invasion, and migration of NSCLC: a study based on immunohistochemistry, TCGA, bioinformatics, in vitro and in vivo verification. Oncotarget, 2017, 8, 16531-16552.	2.1	27
227	Prediction of clinical outcome and survival in soft-tissue sarcoma using a ten-lncRNA signature. Oncotarget, 2017, 8, 80336-80347.	2.1	18
228	The protective value of miR-204-5p for prognosis and its potential gene network in various malignancies: a comprehensive exploration based on RNA-seq high-throughput data and bioinformatics. Oncotarget, 2017, 8, 104960-104980.	2.1	10
229	Clinical role and biological function of CDK5 in hepatocellular carcinoma: A study based on immunohistochemistry, RNA-seq and in vitro investigation. Oncotarget, 2017, 8, 108333-108354.	2.1	14
230	Survival prediction of kidney renal papillary cell carcinoma by comprehensive LncRNA characterization. Oncotarget, 2017, 8, 110811-110829.	2.1	23
231	From big data to diagnosis and prognosis: gene expression signatures in liver hepatocellular carcinoma. PeerJ, 2017, 5, e3089.	2.0	33
232	Expression and clinicopathological implication of DcR3 in lung cancer tissues: a tissue microarray study with 365 cases. OncoTargets and Therapy, 2016, Volume 9, 4959-4968.	2.1	16
233	Effect of DcR3-specific siRNA on cell growth suppression and apoptosis induction in glioma cells via affecting ERK and AKT. OncoTargets and Therapy, 2016, Volume 9, 5195-5202.	2.1	10
234	Clinicopathological significance of STAT4 in hepatocellular carcinoma and its effect on cell growth and apoptosis. OncoTargets and Therapy, 2016, 9, 1721.	2.1	9

#	Article	IF	Citations
235	Clinicopathological role of miR-30a-5p in hepatocellular carcinoma tissues and prediction of its function with bioinformatics analysis. OncoTargets and Therapy, 2016, Volume 9, 5061-5071.	2.1	19
236	Lower expressed miR-198 and its potential targets in hepatocellular carcinoma: a clinicopathological and in silico study. OncoTargets and Therapy, 2016, Volume 9, 5163-5180.	2.1	33
237	Human papillomavirus as a potential risk factor for gastric cancer: a meta-analysis of 1,917 cases. OncoTargets and Therapy, 2016, Volume 9, 7105-7114.	2.1	41
238	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
239	Prognostic Values of Vimentin Expression and Its Clinicopathological Significance in Non-Small Cell Lung Cancer: A Meta-Analysis of Observational Studies with 4118 Cases. PLoS ONE, 2016, 11, e0163162.	2.5	46
240	High Ki-67 Immunohistochemical Reactivity Correlates With Poor Prognosis in Bladder Carcinoma. Medicine (United States), 2016, 95, e3337.	1.1	21
241	Comprehensive analysis of the long noncoding RNA HOXA11-AS gene interaction regulatory network in NSCLC cells. Cancer Cell International, 2016, 16, 89.	4.3	56
242	Downregulation of microRNA-132 indicates progression in hepatocellular carcinoma. Experimental and Therapeutic Medicine, 2016, 12, 2095-2101.	1.9	27
243	Evaluation and clinical significance of cyclin-dependent kinase5 expression in cervical lesions: a clinical research study in Guangxi, China. European Journal of Medical Research, 2016, 21, 28.	2.2	3
244	Relationship between TRAF6 and deterioration of HCC: an immunohistochemical and in vitro study. Cancer Cell International, 2016, 16, 76.	4.3	27
245	Comprehensive investigation of aberrant microRNA profiling in bladder cancer tissues. Tumor Biology, 2016, 37, 12555-12569.	1.7	30
246	Prognostic Value of Expression of Cyclin E in Gastrointestinal Cancer. Technology in Cancer Research and Treatment, 2016, 15, 12-19.	1.9	14
247	Lithium chloride modulates chondrocyte primary cilia and inhibits Hedgehog signaling. FASEB Journal, 2016, 30, 716-726.	0.5	51
248	Importance of the intraoperative appearance of preserved parathyroid glands after total thyroidectomy. Surgery Today, 2016, 46, 356-362.	1.5	11
249	Neurotensin signaling stimulates glioblastoma cell proliferation by upregulating c-Myc and inhibiting miR-29b-1 and miR-129-3p. Neuro-Oncology, 2016, 18, 216-226.	1.2	34
250	Decoy Receptor 3 (DcR3) as a Biomarker of Tumor Deterioration in Female Reproductive Cancers: A Meta-Analysis. Medical Science Monitor, 2016, 22, 1850-1857.	1.1	11
251	Expression and clinicopathological significance of miR-193a-3p and its potential target astrocyte elevated gene-1 in non-small lung cancer tissues. Cancer Cell International, 2015, 15, 80.	4.3	30
252	Association between underexpression of microrna-203 and clinicopathological significance in hepatocellular carcinoma tissues. Cancer Cell International, 2015, 15, 62.	4.3	34

#	Article	IF	Citations
253	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
254	Overexpression of MMP Family Members Functions as Prognostic Biomarker for Breast Cancer Patients: A Systematic Review and Meta-Analysis. PLoS ONE, 2015, 10, e0135544.	2.5	158
255	Down-Regulation of MiR-193a-3p Dictates Deterioration of HCC: A Clinical Real-Time qRT-PCR Study. Medical Science Monitor, 2015, 21, 2352-2360.	1.1	27
256	The Prognostic Role of Ki-67/MIB-1 in Cervical Cancer: A Systematic Review with Meta-Analysis. Medical Science Monitor, 2015, 21, 882-889.	1.1	28
257	MiR-133a is downregulated in non-small cell lung cancer: a study of clinical significance. European Journal of Medical Research, 2015, 20, 50.	2.2	35
258	Upregulation and Clinicopathological Significance of Long Non-coding NEAT1 RNA in NSCLC Tissues. Asian Pacific Journal of Cancer Prevention, 2015, 16, 2851-2855.	1.2	83
259	Sp1 cooperates with Sp3 to upregulate MALAT1 expression in human hepatocellular carcinoma. Oncology Reports, 2015, 34, 2403-2412.	2.6	57
260	A Galerkin method with spherical splines for the shallow water equations on a sphere: error analysis. Numerische Mathematik, 2015, 129, 783-814.	1.8	1
261	MicroRNA-141 Is a Biomarker for Progression of Squamous Cell Carcinoma and Adenocarcinoma of the Lung: Clinical Analysis of 125 Patients. Tohoku Journal of Experimental Medicine, 2015, 235, 161-169.	1.3	23
262	A circulating miRNA signature as a diagnostic biomarker for non-invasive early detection of breast cancer. Breast Cancer Research and Treatment, 2015, 154, 423-434.	2.5	95
263	Downregulation of MiR-30a is Associated with Poor Prognosis in Lung Cancer. Medical Science Monitor, 2015, 21, 2514-2520.	1.1	49
264	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
265	Long noncoding RNAs in hepatocellular carcinoma: Novel insights into their mechanism. World Journal of Hepatology, 2015, 7, 2781.	2.0	45
266	Expression of Tumor Necrosis Factor Receptor-associated Factor 6 in Lung Cancer Tissues. Asian Pacific Journal of Cancer Prevention, 2015, 15, 10591-10596.	1.2	20
267	Ki-67 is a Valuable Prognostic Factor in Gliomas: Evidence from a Systematic Review and Meta-analysis. Asian Pacific Journal of Cancer Prevention, 2015, 16, 411-420.	1.2	110
268	Overexpression of DcR3 and Its Significance on Tumor Cell Differentiation and Proliferation in Glioma. Scientific World Journal, The, 2014, 2014, 1-7.	2.2	16
269	Decreased expression and clinical significance of miR-148a in hepatocellular carcinoma tissues. European Journal of Medical Research, 2014, 19, 68.	2.2	47
270	Synergistic Effect of MiR-146a Mimic and Cetuximab on Hepatocellular Carcinoma Cells. BioMed Research International, 2014, 2014, 1-15.	2.0	31

#	Article	IF	Citations
271	Expression and clinicopathological significance of miR-146a in hepatocellular carcinoma tissues. Upsala Journal of Medical Sciences, 2014, 119, 19-24.	0.9	63
272	Down-regulation of ribosomal protein S15A mRNA with a short hairpin RNA inhibits human hepatic cancer cell growth in vitro. Gene, 2014, 536, 84-89.	2.3	35
273	SLIP FACTOR OF HIGH STRENGTH BOLTED FRICTIONAL JOINTS WITH ROUGHENED STEEL SURFACE AND INORGANIC ZINC RICH PAINTED SURFACE. Journal of Japan Society of Civil Engineers Ser A1 (Structural) Tj ETQq1 1 0.784314 rgBT /Ove		
274	Effects of miR-152 on Cell Growth Inhibition, Motility Suppression and Apoptosis Induction in Hepatocellular Carcinoma Cells. Asian Pacific Journal of Cancer Prevention, 2014, 15, 4969-4976.	1.2	53
275	Overexpression and Clinicopathological Contribution of DcR3 in Bladder Urothelial Carcinoma Tissues. Asian Pacific Journal of Cancer Prevention, 2014, 15, 9137-9142.	1.2	9
276	Expression and Prognostic Significance of lncRNA MALAT1 in Pancreatic Cancer Tissues. Asian Pacific Journal of Cancer Prevention, 2014, 15, 2971-2977.	1.2	138
277	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. Biochemical and Biophysical Research Communications, 2013, 431, 623-629.	2.2	35
278	Increased MiR-221 expression in hepatocellular carcinoma tissues and its role in enhancing cell growth and inhibiting apoptosis in vitro. BMC Cancer, 2013, 13, 21.	2.6	110
279	Synergistic Effect of Afatinib with Su11274 in Non-Small Cell Lung Cancer Cells Resistant to Gefitinib or Erlotinib. PLoS ONE, 2013, 8, e59708.	2.5	44
280	miR-146a Inhibits Cell Growth, Cell Migration and Induces Apoptosis in Non-Small Cell Lung Cancer Cells. PLoS ONE, 2013, 8, e60317.	2.5	233
281	Underexpression of miR-34a in Hepatocellular Carcinoma and Its Contribution towards Enhancement of Proliferating Inhibitory Effects of Agents Targeting c-MET. PLoS ONE, 2013, 8, e61054.	2.5	113
282	Application of Recrystallization Texture Evolution Model to Type 430 Stainless-Steel Strip Production. Materials Transactions, 2012, 53, 1837-1846.	1.3	2
283	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. BMC Medicine, 2012, 10, 28.	5.7	111
284	The critical care nurse's role in Endâ€ofâ€Life care: issues and challenges. Nursing in Critical Care, 2011, 16, 116-123.	2.5	48
285	A Pilot Study Examining Activity Participation, Sensory Responsiveness, and Competence in Children with High Functioning Autism Spectrum Disorder. Journal of Autism and Developmental Disorders, 2011, 41, 1496-1506.	3.1	139
286	Reinvasion by ship rats (Rattus rattus) of forest fragments after eradication. Biological Invasions, 2011, 13, 2391-2408.	2.4	43
287	Clinicopathological significance of RASSF1A reduced expression and hypermethylation in hepatocellular carcinoma. Hepatology International, 2010, 4, 423-432.	4.4	55
288	TNFRSF6B neutralization antibody inhibits proliferation and induces apoptosis in hepatocellular carcinoma cell. Pathology Research and Practice, 2010, 206, 631-641.	2.3	20

#	Article	IF	Citations
289	Quantification of epidermal growth factor receptor T790M mutant transcripts in lung cancer cells by real-time reverse transcriptaseâ€“quantitative polymerase chain reaction. Analytical Biochemistry, 2010, 398, 266-268.	2.5	23
290	Significance of decoy receptor 3 in sera of hepatocellular carcinoma patients. Upsala Journal of Medical Sciences, 2010, 115, 232-237.	0.9	19
291	Influence of RT-qPCR primer position on EGFR interference efficacy in lung cancer cells. Biological Procedures Online, 2010, 13, 1.	3.0	39
292	Inhibition of Î³-secretase induces G2/M arrest and triggers apoptosis in breast cancer cells. British Journal of Cancer, 2009, 100, 1879-1888.	6.6	80
293	Over-expression of Decoy Receptor 3 in Gastric Precancerous Lesions and Carcinoma. Upsala Journal of Medical Sciences, 2008, 113, 297-304.	0.9	19
294	Expression of Heparanase in Hepatocellular Carcinoma Has Prognostic Significance: A Tissue Microarray Study. Oncology Research, 2008, 17, 183-189.	1.6	21
295	Estado del conocimiento de humedales del norte patagÃ³nico (Argentina): aspectos relevantes e importancia para la conservaciÃ³n de la biodiversidad regional. Revista Chilena De Historia Natural, 2005, 78, 723.	1.2	58
296	Bootstrap Current Increment After Siliconization on the HT-7 Tokamak. Chinese Physics Letters, 2002, 19, 1141-1143.	3.4	5
297	Evaluation of the role of radiotherapy in the management of carcinoma of the buccal mucosa. Cancer, 1988, 61, 1326-1331.	4.1	74
298	Retention of mercury in sediments of the Baltic Sea near Oulu, Finland. Journal of Paleolimnology, 1988, 1, 133.	1.5	1
299	Species differences in pharmacokinetics. Veterinary Research Communications, 1983, 7, 285-300.	1.6	6
300	137. Indikationen und Ergebnisse bei Totalkorrektur der Transposition der groÃŸen Arterien nach Mustard. Langenbeck's Archives of Surgery, 1967, 319, 811-816.	1.9	0
301	Validation of BET proteins as therapeutic targets in Neurofibromatosis type 2. Neuro-Oncology Advances, 0, , .	0.7	0
302	Expression Profile and Molecular Basis of Cyclin-Dependent Kinases Regulatory Subunit 2 in Endometrial Carcinoma Detected by Diversified Methods. Pathology and Oncology Research, 0, 28, .	1.9	0
303	Deep Learning-Based Multi-Omics Integration Robustly Predicts Relapse in Prostate Cancer. Frontiers in Oncology, 0, 12, .	2.9	12
304	Metadherin Promotes the Development of Bladder Cancer by Enhancing Cell Division. Cancer Biotherapy and Radiopharmaceuticals, 0, , .	1.1	1
305	Upregulation of CCNB2 and Its Perspective Mechanisms in Cerebral Ischemic Stroke and All Subtypes of Lung Cancer: A Comprehensive Study. Frontiers in Integrative Neuroscience, 0, 16, .	2.1	8
306	Downregulated Dual-Specificity Protein Phosphatase 1 in Ovarian Carcinoma: A Comprehensive Study With Multiple Methods. Pathology and Oncology Research, 0, 28, .	1.9	2

#	Article	IF	Citations
307	What enlightenment has the development of lung cancer bone metastasis brought in the last 22 years. World Journal of Clinical Oncology, 0, 15, 765-782.	2.3	0
308	A Comprehensive Analysis of LYAR in Colorectal Cancer: Prognostic Marker and Therapeutic Target. Cancer Biotherapy and Radiopharmaceuticals, 0, , .	1.1	0