Gang Chen

List of Publications by Year
in descending order

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331 papers	7,756 citations	66315 42 h-index	114418 63 g-index
340 all docs	340 docs citations	340 times ranked	9924 citing authors

#	Article	IF	Citations
1	Clinical Implication of E2F Transcription Factor 1 in Hepatocellular Carcinoma Tissues. Cancer Biotherapy and Radiopharmaceuticals, 2023, 38, 684-707.	0.7	7
2	Clinical Significance of Integrin Subunit Beta 4 in Head and Neck Squamous Cell Carcinoma. Cancer Biotherapy and Radiopharmaceuticals, 2022, 37, 256-275.	0.7	11
3	Downregulation of miR-125b-5p and Its Prospective Molecular Mechanism in Lung Squamous Cell Carcinoma. Cancer Biotherapy and Radiopharmaceuticals, 2022, 37, 125-140.	0.7	4
4	Laryngeal Squamous Cell Carcinoma: Clinical Significance and Potential Mechanism of Cell Division Cycle 45. Cancer Biotherapy and Radiopharmaceuticals, 2022, 37, 300-312.	0.7	4
5	Clinicopathological significance and underlying molecular mechanism of downregulation of basonuclin 1 expression in ovarian carcinoma. Experimental Biology and Medicine, 2022, 247, 106-119.	1.1	7
6	Expression Landscape and Functional Roles of HOXA4 and HOXA5 in Lung Adenocarcinoma. International Journal of Medical Sciences, 2022, 19, 572-587.	1.1	2
7	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.	0.6	1
8	Expression of IER3 in hepatocellular carcinoma: clinicopathology, prognosis, and potential regulatory pathways. PeerJ, 2022, 10, e12944.	0.9	3
9	Decreased expression of transcription factor Homeobox A11 and its potential target genes in bladder cancer. Pathology Research and Practice, 2022, 233, 153847.	1.0	3
10	Ogt Demonstrated Conspicuous Clinical Significance in Cancers, from Pan-Cancer to Small-Cell Lung Cancer. Journal of Oncology, 2022, 2022, 1-16.	0.6	3
11	Upregulation of the transmembrane protease serine 3ÂmRNA level in radioresistant colorectal cancer tissues. Biomarkers in Medicine, 2022, , .	0.6	2
12	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.	0.7	3
13	Clinical significance of cyclin-dependent kinase inhibitor 2C expression in cancers: from small cell lung carcinoma to pan-cancers. BMC Pulmonary Medicine, 2022, 22, .	0.8	8
14	Comprehensive expression analysis reveals upregulated LUZP2 in prostate cancer tissues. Electronic Journal of Biotechnology, 2022, 59, 1-12.	1.2	0
15	Clinical assessment and molecular mechanism of the upregulation of Toll-like receptor 2 (TLR2) in myocardial infarction. BMC Cardiovascular Disorders, 2022, 22, .	0.7	5
16	Upregulation of ATIC in multiple myeloma tissues based on tissue microarray and gene microarrays. International Journal of Laboratory Hematology, 2021, 43, 409-417.	0.7	8
17	Ultrasound Imageâ€“Based Radiomics. Journal of Ultrasound in Medicine, 2021, 40, 1229-1244.	0.8	18
18	Clinical significance and molecular mechanism of angiotensin-converting enzyme 2 in hepatocellular carcinoma tissues. Bioengineered, 2021, 12, 4054-4069.	1.4	11

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19	Clinical significance and potential molecular mechanism of miRNA-222-3p in metastatic prostate cancer. Bioengineered, 2021, 12, 325-340.	1.4	23
20	Expression and Clinical Significance of BCL2 Interacting Protein 3 Like in Multiple Myeloma. Technology in Cancer Research and Treatment, 2021, 20, 153303382110245.	0.8	1
21	MiRNAâ€145â€5p expression and prospective molecular mechanisms in the metastasis of prostate cancer. IET Systems Biology, 2021, 15, 1-13.	0.8	8
22	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.	1.3	26
23	Development and Validation of a Radiomic Nomogram for Predicting the Prognosis of Kidney Renal Clear Cell Carcinoma. Frontiers in Oncology, 2021, 11, 613668.	1.3	7
24	LPCAT1 overexpression promotes the progression of hepatocellular carcinoma. Cancer Cell International, 2021, 21, 442.	1.8	24
25	The Indication of Poor Prognosis by High Expression of ENO1 in Squamous Cell Carcinoma of the Lung. Journal of Oncology, 2021, 2021, 1-11.	0.6	4
26	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, , .	0.7	0
27	Overexpression of cyclinâ€dependent kinase 1 in esophageal squamous cell carcinoma and its clinical significance. FEBS Open Bio, 2021, 11, 3126-3141.	1.0	5
28	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.	1.4	3
29	Upregulation of microRNA miR-141-3p and its prospective targets in endometrial carcinoma: a comprehensive study. Bioengineered, 2021, 12, 2941-2956.	1.4	10
30	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.	1.2	6
31	Down-regulation of microRNA-125b-2-3p is a risk factor for a poor prognosis in hepatocellular carcinoma. Bioengineered, 2021, 12, 1627-1641.	1.4	9
32	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.	0.5	11
33	Clinical Value and Potential Mechanism of miRNA-33a-5p in Lung Squamous Cell Carcinoma. Analytical Cellular Pathology, 2021, 2021, 1-20.	0.7	2
34	Clinical significance and effect of lncRNA BBOX1â€‘AS1 on the proliferation and migration of lung squamous cell carcinoma. Oncology Letters, 2021, 23, 17.	0.8	4
35	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.	1.9	8
36	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.	0.8	3

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37	A novel risk signature that combines 10 long noncoding RNAs to predict neuroblastoma prognosis. Journal of Cellular Physiology, 2020, 235, 3823-3834.	2.0	15
38	A radiogenomics signature for predicting the clinical outcome of bladder urothelial carcinoma. European Radiology, 2020, 30, 547-557.	2.3	39
39	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.	0.7	25
40	The role of upregulated miR-375 expression in breast cancer: An in vitro and in silico study. Pathology Research and Practice, 2020, 216, 152754.	1.0	27
41	The clinical significance and potential molecular mechanism of integrin subunit beta 4 in laryngeal squamous cell carcinoma. Pathology Research and Practice, 2020, 216, 152785.	1.0	7
42	Clinical significance of transcription factor RUNX2 in lung adenocarcinoma and its latent transcriptional regulating mechanism. Computational Biology and Chemistry, 2020, 89, 107383.	1.1	12
43	Clinical significance of CCNE2 protein and mRNA expression in thyroid cancer tissues. Advances in Medical Sciences, 2020, 65, 442-456.	0.9	9
44	Clinical Significance of the Interleukin 24 mRNA Level in Head and Neck Squamous Cell Carcinoma and Its Subgroups: An In Silico Investigation. Journal of Oncology, 2020, 2020, 1-15.	0.6	3
45	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.	1.3	5
46	Downregulation of miRNA-205 Expression and Biological Mechanism in Prostate Cancer Tumorigenesis and Bone Metastasis. BioMed Research International, 2020, 2020, 1-17.	0.9	11
47	Prognostic Values for the mRNA Expression of the ADAMTS Family of Genes in Gastric Cancer. Journal of Oncology, 2020, 2020, 1-24.	0.6	14
48	Clinical value and potential mechanisms of COL8A1 upregulation in breast cancer: a comprehensive analysis. Cancer Cell International, 2020, 20, 392.	1.8	20
49	The clinical value and potential molecular mechanism of the downregulation of MAOA in hepatocellular carcinoma tissues. Cancer Medicine, 2020, 9, 8004-8019.	1.3	24
50	Immunohistochemical basigin expression level in thyroid cancer tissues. World Journal of Surgical Oncology, 2020, 18, 240.	0.8	2
51	Estrogenic activities of compound GL-1, isolated from Ganoderma lucidum. Natural Product Research, 2020, 35, 1-5.	1.0	2
52	The Expression and Potential Role of Tubulin Alpha 1b in Wilmsâ€™ Tumor. BioMed Research International, 2020, 2020, 1-10.	0.9	8
53	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.	0.8	6
54	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.	1.2	4

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55	The clinical significance of interleukin 24 and its potential molecular mechanism in laryngeal squamous cell carcinoma. Cancer Biomarkers, 2020, 29, 111-124.	0.8	5
56	Downregulation of miRNAâ€126â€3p is associated with progression of and poor prognosis for lung squamous cell carcinoma. FEBS Open Bio, 2020, 10, 1624-1641.	1.0	10
57	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.	1.4	10
58	Radiomic profiles in diffuse glioma reveal distinct subtypes with prognostic value. Journal of Cancer Research and Clinical Oncology, 2020, 146, 1253-1262.	1.2	16
59	Development and validation of an immune prognostic classifier for clear cell renal cell carcinoma. Cancer Biomarkers, 2020, 27, 265-275.	0.8	7
60	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.	0.9	0
61	Effect of CELSR3 on the Cell Cycle and Apoptosis of Hepatocellular Carcinoma Cells. Journal of Cancer, 2020, 11, 2830-2844.	1.2	8
62	Clinicopathological value and underlying molecular mechanism of annexin A2 in 992 cases of thyroid carcinoma. Computational Biology and Chemistry, 2020, 86, 107258.	1.1	6
63	Prognostic value of small nucleolar RNAs (snoRNAs) for colon adenocarcinoma based on RNA sequencing data. Pathology Research and Practice, 2020, 216, 152937.	1.0	17
64	The Clinical Significance and Potential Molecular Mechanism of PTTG1 in Esophageal Squamous Cell Carcinoma. Frontiers in Genetics, 2020, 11, 583085.	1.1	11
65	The Latest Overview of circRNA in the Progression, Diagnosis, Prognosis, Treatment, and Drug Resistance of Hepatocellular Carcinoma. Frontiers in Oncology, 2020, 10, 608257.	1.3	16
66	Integrated expression analysis revealed RUNX2 upregulation in lung squamous cell carcinoma tissues. IET Systems Biology, 2020, 14, 252-260.	0.8	7
67	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.	0.5	3
68	Identification of an Immune Score-Based Gene Panel with Prognostic Power for Oral Squamous Cell Carcinoma. Medical Science Monitor, 2020, 26, e922854.	0.5	17
69	Downregulation of CDC14B in 5218 breast cancer patients: A novel prognosticator for triple-negative breast cancer. Mathematical Biosciences and Engineering, 2020, 17, 8152-8181.	1.0	3
70	Nomogram for predicting overall survival in children with neuroblastoma based on SEER database. Annals of Surgical Treatment and Research, 2020, 99, 118.	0.4	10
71	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.	1.2	4
72	Polo like kinase 1 expression in cervical cancer tissues generated from multiple detection methods. PeerJ, 2020, 8, e10458.	0.9	13

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73	Downregulation of miR-193a-3p is involved in the pathogenesis of hepatocellular carcinoma by targeting CCND1. PeerJ, 2020, 8, e8409.	0.9	13
74	Upregulated expression of SAC3D1 is associated with progression in gastric cancer. International Journal of Oncology, 2020, 57, 122-138.	1.4	4
75	The clinical significance of apolipoprotein L1 in head and neck squamous cell carcinoma. Oncology Letters, 2020, 20, 1-1.	0.8	8
76	Small Nucleolar RNAs (snoRNAs)-Based Risk Score Classifier Predicts Overall Survival in Bladder Carcinoma. Medical Science Monitor, 2020, 26, e926273.	0.5	1
77	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.	0.8	9
78	Expression levels and coâ€'targets of miRNAâ€'126â€'3p and miRNAâ€'126â€'5p in lung adenocarcinoma tissues: În exploration with RTâ€'qPCR, microarray and bioinformatic analyses. Oncology Reports, 2019, 41, 939-953.	1.2	13
79	Expression of miRâ€'542â€'3p in osteosarcoma with miRNA microarray data, and its potential signaling pathways. Molecular Medicine Reports, 2019, 19, 974-983.	1.1	3
80	Identification of putative drugs for gastric adenocarcinoma utilizing differentially expressed genes and connectivity map. Molecular Medicine Reports, 2019, 19, 1004-1015.	1.1	3
81	Clinical value of microRNAâ€'198â€'5p downregulation in lung adenocarcinoma and its potential pathways. Oncology Letters, 2019, 18, 2939-2954.	0.8	12
82	Expression significance and potential mechanism of hypoxiaâ€inducible factor 1 alpha in patients with myelodysplastic syndromes. Cancer Medicine, 2019, 8, 6021-6035.	1.3	11
83	Expression and clinical significance of neuropilin-1 in Epstein-Barr virus-associated lymphomas. Cancer Biomarkers, 2019, 25, 259-273.	0.8	5
84	Profiling of prognostic alternative splicing in melanoma. Oncology Letters, 2019, 18, 1081-1088.	0.8	8
85	Development of a prognostic index based on an immunogenomic landscape analysis of papillary thyroid cancer. Aging, 2019, 11, 480-500.	1.4	132
86	Prospective molecular mechanism of COL5A1 in breast cancer based on a microarray, RNA sequencing and immunohistochemistry. Oncology Reports, 2019, 42, 151-175.	1.2	24
87	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.	1.8	12
88	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.	1.4	21
89	Clinical Significance of microRNA-196b-5p in Hepatocellular Carcinoma and its Potential Molecular Mechanism. Journal of Cancer, 2019, 10, 5355-5370.	1.2	12
90	Role of alternative splicing signatures in the prognosis of glioblastoma. Cancer Medicine, 2019, 8, 7623-7636.	1.3	20

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91	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.	1.2	29
92	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.	1.1	8
93	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.	1.8	24
94	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.	1.8	32
95	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.	0.8	2
96	High throughput circRNA sequencing analysis reveals novel insights into the mechanism of nitidine chloride against hepatocellular carcinoma. Cell Death and Disease, 2019, 10, 658.	2.7	50
97	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.	0.1	4
98	Identification and validation of an individualized autophagy-clinical prognostic index in bladder cancer patients . OncoTargets and Therapy, 2019, Volume 12, 3695-3712.	1.0	37
99	MiR-193a-3p inhibits pancreatic ductal adenocarcinoma cell proliferation by targeting CCND1 . Cancer Management and Research, 2019, Volume 11, 4825-4837.	0.9	17
100	Comprehensive evaluation of FKBP10 expression and its prognostic potential in gastric cancer. Oncology Reports, 2019, 42, 615-628.	1.2	19
101	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.	1.0	9
102	Gene profiling of HepG2 cells following nitidine chloride treatment: An investigation with microarray and Connectivity Mapping. Oncology Reports, 2019, 41, 3244-3256.	1.2	9
103	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.	1.4	19
104	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.	1.4	46
105	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.	1.0	9
106	Clinical and genetic characteristics of female dystrophinopathy carriers. Molecular Medicine Reports, 2019, 19, 3035-3044.	1.1	18
107	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.	1.2	25
108	Inï¿½silico analysis identified miRNAâ€‘based therapeutic agents against glioblastoma multiforme. Oncology Reports, 2019, 41, 2194-2208.	1.2	29

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109	miRâ€'146aâ€'5p targets TCSF and influences cell growth and apoptosis to repress NSCLC progression. Oncology Reports, 2019, 41, 2226-2240.	1.2	17
110	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.	1.2	8
111	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.	1.0	12
112	The clinical significance of endothelin receptor type B in hepatocellular carcinoma and its potential molecular mechanism. Experimental and Molecular Pathology, 2019, 107, 141-157.	0.9	21
113	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.	1.0	4
114	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.	1.0	16
115	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.	1.0	13
116	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.	1.0	14
117	EBV as a potential risk factor for hepatobiliary system cancer: A meta-analysis with 918 cases. Pathology Research and Practice, 2019, 215, 278-285.	1.0	8
118	An air freight forwarderâ€™s resource planning and revenue management. Journal of the Operational Research Society, 2019, 70, 294-309.	2.1	5
119	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.	0.3	3
120	A novel alternative splicing-based prediction model for uteri corpus endometrial carcinoma. Aging, 2019, 11, 263-283.	1.4	16
121	Role of global aberrant alternative splicing events in papillary thyroid cancer prognosis. Aging, 2019, 11, 2082-2097.	1.4	37
122	Identification of hub genes in prostate cancer using robust rank aggregation and weighted gene co-expression network analysis. Aging, 2019, 11, 4736-4756.	1.4	82
123	Potential ceRNA networks involved in autophagy suppression of pancreatic cancer caused by chloroquine diphosphate: A study based on differentiallyâ€'expressed circRNAs, lncRNAs, miRNAs and mRNAs. International Journal of Oncology, 2019, 54, 600-626.	3.9	33
124	Comprehensive clinical implications of homeobox A10 in 3,199 cases of non-small cell lung cancer tissue samples combining qRT-PCR, RNA sequencing and microarray data. American Journal of Translational Research (discontinued), 2019, 11, 45-66.	0.0	8
125	Prognostic alternative splicing signatures and underlying regulatory network in esophageal carcinoma. American Journal of Translational Research (discontinued), 2019, 11, 4010-4028.	0.0	7
126	Overexpressed BSG related to the progression of lung adenocarcinoma with high-throughput data-mining, immunohistochemistry, validation and investigation. American Journal of Translational Research (discontinued), 2019, 11, 4835-4850.	0.0	12

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127	Clinical roles of miR-136-5p and its target metadherin in thyroid carcinoma. American Journal of Translational Research (discontinued), 2019, 11, 6754-6774.	0.0	10
128	Integrated assessment of CDK1 upregulation in thyroid cancer. American Journal of Translational Research (discontinued), 2019, 11, 7233-7254.	0.0	10
129	Identifying TF-miRNA-mRNA regulatory modules in nitidine chloride treated HCC xenograft of nude mice. American Journal of Translational Research (discontinued), 2019, 11, 7503-7522.	0.0	7
130	The clinicopathological significance of decreased miR-125b-5p in hepatocellular carcinoma: evidence based on RT-qPCR, microRNA-microarray, and microRNA-sequencing. International Journal of Clinical and Experimental Pathology, 2019, 12, 21-39.	0.5	2
131	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.	1.1	40
132	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.	1.8	36
133	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.	1.0	13
134	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.	1.1	22
135	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.	1.4	6
136	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.	1.0	19
137	Clinical Significance of miR-210 and its Prospective Signaling Pathways in Non-Small Cell Lung Cancer: Evidence from Gene Expression Omnibus and the Cancer Genome Atlas Data Mining with 2763 Samples and Validation via Real-Time Quantitative PCR. Cellular Physiology and Biochemistry, 2018, 46, 925-952.	1.1	26
138	Prognostic Significance of LncRNA PVT1 and Its Potential Target Gene Network in Human Cancers: a Comprehensive Inquiry Based Upon 21 Cancer Types and 9972 Cases. Cellular Physiology and Biochemistry, 2018, 46, 591-608.	1.1	16
139	In silico analysis of the potential mechanism of telocinobufagin on breast cancer MCF-7 cells. Pathology Research and Practice, 2018, 214, 631-643.	1.0	8
140	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.	0.8	17
141	Clinical significance of high expression of miR-452-5p in lung squamous cell carcinoma. Oncology Letters, 2018, 15, 6418-6430.	0.8	37
142	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.	1.0	36
143	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.	1.0	12
144	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.	1.0	4

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145	The clinical value of miRâ€193aâ€3p in nonâ€small cell lung cancer and its potential molecular mechanism explored in silico using <scp>RNA</scp>â€sequencing and microarray data. FEBS Open Bio, 2018, 8, 94-109.	1.0	11
146	Diagnostic value of strandâ€specific mi<scp>RNA</scp>â€101â€3p and mi<scp>RNA</scp>â€101â€5p for hepatocellular carcinoma and a bioinformatic analysis of their possible mechanism of action. FEBS Open Bio, 2018, 8, 64-84.	1.0	22
147	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.	0.8	2
148	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.	0.8	22
149	Oncogenic value of microRNAâ€'15bâ€'5p in hepatocellular carcinoma and a bioinformatics investigation. Oncology Letters, 2018, 17, 1695-1713.	0.8	17
150	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.	0.9	1
151	The Clinicopathological Significance and Correlative Signaling Pathways of an Autophagy-Related Gene, Ambra1, in Breast Cancer: a Study of 25 Microarray RNA-Seq Datasets and in-House Gene Silencing. Cellular Physiology and Biochemistry, 2018, 51, 1027-1040.	1.1	3
152	Expression Signature and Role of miR-30d-5p in Non-Small Cell Lung Cancer: a Comprehensive Study Based on in Silico Analysis of Public Databases and in Vitro Experiments. Cellular Physiology and Biochemistry, 2018, 50, 1964-1987.	1.1	24
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.	1.1	4
154	Genomic analysis of small nucleolar RNAs identifies distinct molecular and prognostic signature in hepatocellular carcinoma. Oncology Reports, 2018, 40, 3346-3358.	1.2	15
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.	0.8	19
156	Upregulation of HOXA11 during the progression of lung adenocarcinoma detected via multiple approaches. International Journal of Molecular Medicine, 2018, 42, 2650-2664.	1.8	29
157	Biological Effect and Mechanism of the miR-23b-3p/ANXA2 Axis in Pancreatic Ductal Adenocarcinoma. Cellular Physiology and Biochemistry, 2018, 50, 823-840.	1.1	20
158	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.	1.1	12
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