Bangshun He

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Novel insights into the interaction between N6-methyladenosine modification and circular RNA. Molecular Therapy - Nucleic Acids, 2022, 27, 824-837.	5.1	19
2	Association Between SNPs in the One-Carbon Metabolism Pathway and the Risk of Female Breast Cancer in a Chinese Population. Pharmacogenomics and Personalized Medicine, 2022, Volume 15, 9-16.	0.7	1
3	Identification of autophagy related genes in predicting the prognosis and aiding 5- fluorouracil therapy of colorectal cancer. Heliyon, 2022, 8, e09033.	3.2	3
4	Susceptibility of Genetic Variations in Methylation Pathway to Gastric Cancer. Pharmacogenomics and Personalized Medicine, 2022, Volume 15, 441-448.	0.7	1
5	LncRNA SPINT1-AS1 promotes breast cancer proliferation and metastasis by sponging let-7 a/b/i-5p. Pathology Research and Practice, 2021, 217, 153268.	2.3	26
6	The diagnostic and prognostic values of microRNA-196a in cancer. Bioscience Reports, 2021, 41, .	2.4	17
7	3044 Cases reveal important prognosis signatures of COVID-19 patients. Computational and Structural Biotechnology Journal, 2021, 19, 1163-1175.	4.1	11
8	Long intergenic non-coding RNA LINC00485 exerts tumor-suppressive activity by regulating miR-581/EDEM1 axis in colorectal cancer. Aging, 2021, 13, 3866-3885.	3.1	5
9	Magnetic Colloid Antibodies Accelerate Small Extracellular Vesicles Isolation for Point-of-Care Diagnostics. Nano Letters, 2021, 21, 2001-2009.	9.1	26
10	Upregulated IL-6 Indicates a Poor COVID-19 Prognosis: A Call for Tocilizumab and Convalescent Plasma Treatment. Frontiers in Immunology, 2021, 12, 598799.	4.8	24
11	Genetic Variation of Inflammatory Genes to Ischemic Stroke Risk in a Chinese Han Population. Pharmacogenomics and Personalized Medicine, 2021, Volume 14, 977-986.	0.7	2
12	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. Gut, 2020, 69, 641-651.	12.1	36
13	FoxO3 reverses 5-fluorouracil resistance in human colorectal cancer cells by inhibiting the Nrf2/TR1 signaling pathway. Cancer Letters, 2020, 470, 29-42.	7.2	48
14	Macrophage-derived CCL5 facilitates immune escape of colorectal cancer cells via the p65/STAT3-CSN5-PD-L1 pathway. Cell Death and Differentiation, 2020, 27, 1765-1781.	11.2	115
15	Susceptibility of PON1 / PON2 Genetic Variations to Ischemic Stroke Risk in a Chinese Han Population. Pharmacogenomics and Personalized Medicine, 2020, Volume 13, 563-570.	0.7	7
16	Tumor biomarkers predict clinical outcome of COVID-19 patients. Journal of Infection, 2020, 81, 452-482.	3.3	6
17	METTL14-mediated N6-methyladenosine modification of SOX4 mRNA inhibits tumor metastasis in colorectal cancer. Molecular Cancer, 2020, 19, 106.	19.2	188
18	Cerium metal organic framework mediated molecular threading for point-of-care colorimetric assays. Biosensors and Bioelectronics, 2020, 165, 112406.	10.1	24

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19	MiR-485-5p as a potential biomarker and tumor suppressor in human colorectal cancer. Biomarkers in Medicine, 2020, 14, 239-248.	1.4	20
20	LRIG3 represses cell motility by inhibiting slug via inactivating ERK signaling in human colorectal cancer. IUBMB Life, 2020, 72, 1393-1403.	3.4	7
21	Potential False-Positive and False-Negative Results for COVID-19 IgG/IgM Antibody Testing After Heat-Inactivation. Frontiers in Medicine, 2020, 7, 589080.	2.6	6
22	Clinical efficacy of convalescent plasma therapy on treating COVIDâ€19 patients: Evidence from matched study and a metaâ€analysis. Clinical and Translational Medicine, 2020, 10, e259.	4.0	18
23	IGF2 loss of imprinting enhances colorectal cancer stem cells pluripotency by promoting tumor autophagy. Aging, 2020, 12, 21236-21252.	3.1	11
24	Analysis of METTL3 and METTL14 in hepatocellular carcinoma. Aging, 2020, 12, 21638-21659.	3.1	44
25	<p>MicroRNA-371-3 cluster as biomarkers for the diagnosis and prognosis of cancers</p> . Cancer Management and Research, 2019, Volume 11, 5437-5457.	1.9	8
26	Identification of Serum Exosomal hsa-circ-0004771 as a Novel Diagnostic Biomarker of Colorectal Cancer. Frontiers in Genetics, 2019, 10, 1096.	2.3	157
27	IncRNA SNHG6 regulates EZH2 expression by sponging miR-26a/b and miR-214 in colorectal cancer. Journal of Hematology and Oncology, 2019, 12, 3.	17.0	175
28	LncRNA SATB2-AS1 inhibits tumor metastasis and affects the tumor immune cell microenvironment in colorectal cancer by regulating SATB2. Molecular Cancer, 2019, 18, 135.	19.2	205
29	Circulating miR-1290 and miR-320d as Novel Diagnostic Biomarkers of Human Colorectal Cancer. Journal of Cancer, 2019, 10, 43-50.	2.5	53
30	An electrochemiluminescent aptasensor for amplified detection of exosomes from breast tumor cells (MCF-7 cells) based on G-quadruplex/hemin DNAzymes. Analyst, The, 2019, 144, 3668-3675.	3.5	54
31	A self-powered microfluidic chip integrated with fluorescent microscopic counting for biomarkers assay. Sensors and Actuators B: Chemical, 2019, 291, 192-199.	7.8	14
32	Polymorphisms of IL-23R predict survival of gastric cancer patients in a Chinese population. Cytokine, 2019, 117, 79-83.	3.2	6
33	P53-induced miR-1249 inhibits tumor growth, metastasis, and angiogenesis by targeting VEGFA and HMGA2. Cell Death and Disease, 2019, 10, 131.	6.3	66
34	miR-375-3p suppresses tumorigenesis and partially reverses chemoresistance by targeting YAP1 and SP1 in colorectal cancer cells. Aging, 2019, 11, 7357-7385.	3.1	66
35	and genetic variations and gastric cancer risk in the Chinese population. American Journal of Translational Research (discontinued), 2019, 11, 3698-3706.	0.0	6
36	Increased expression of tight junction proteinÃ ⁻ ¿½20ccludin is associated with the protective effect of mosapride against aspirin‑induced gastric injury. Experimental and Therapeutic Medicine, 2018, 15, 1626-1632.	1.8	7

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37	CircHIPK3 promotes colorectal cancer growth and metastasis by sponging miR-7. Cell Death and Disease, 2018, 9, 417.	6.3	497
38	Polymorphisms of TGFBR1, TLR4 are associated with prognosis of gastric cancer in a Chinese population. Cancer Cell International, 2018, 18, 191.	4.1	21
39	The pro-metastasis effect of circANKS1B in breast cancer. Molecular Cancer, 2018, 17, 160.	19.2	219
40	The inhibitory role of miR‑485‑5p in colorectal cancer proliferation and invasion via targeting of CD147. Oncology Reports, 2018, 39, 2201-2208.	2.6	5
41	Genetic variations in PRKAA1 predict the risk and progression of gastric Cancer. BMC Cancer, 2018, 18, 923.	2.6	8
42	The long noncoding RNA SNHG1 regulates colorectal cancer cell growth through interactions with EZH2 and miR-154-5p. Molecular Cancer, 2018, 17, 141.	19.2	259
43	SP1-induced lncRNA-ZFAS1 contributes to colorectal cancer progression via the miR-150-5p/VEGFA axis. Cell Death and Disease, 2018, 9, 982.	6.3	165
44	DNA-methylation-mediated silencing of miR-486-5p promotes colorectal cancer proliferation and migration through activation of PLAGL2/IGF2/β-catenin signal pathways. Cell Death and Disease, 2018, 9, 1037.	6.3	70
45	MiR-490-3p Functions As a Tumor Suppressor by Inhibiting Oncogene VDAC1 Expression in Colorectal Cancer. Journal of Cancer, 2018, 9, 1218-1230.	2.5	50
46	Ginsenoside Rd ameliorates colitis by inducing p62-driven mitophagy-mediated NLRP3 inflammasome inactivation in mice. Biochemical Pharmacology, 2018, 155, 366-379.	4.4	83
47	Exosomal IncRNA 91H is associated with poor development in colorectal cancer by modifying HNRNPK expression. Cancer Cell International, 2018, 18, 11.	4.1	90
48	Serum and exosome long non coding RNAs as potential biomarkers for hepatocellular carcinoma. Journal of Cancer, 2018, 9, 2631-2639.	2.5	97
49	LACTB, a novel epigenetic silenced tumor suppressor, inhibits colorectal cancer progression by attenuating MDM2-mediated p53 ubiquitination and degradation. Oncogene, 2018, 37, 5534-5551.	5.9	62
50	miR-150-5p suppresses tumor progression by targeting VEGFA in colorectal cancer. Aging, 2018, 10, 3421-3437.	3.1	87
51	Meta-analysis of prognostic value of inflammation parameter in breast cancer. Journal of Cancer Research and Therapeutics, 2018, 14, S85-S89.	0.9	10
52	Microfluidics Cell Loadingâ€Dock System: Ordered Cellular Array for Dynamic Lymphocyteâ€Communication Study. Advanced Biology, 2017, 1, e1700085.	3.0	27
53	Fast, Sensitive, and Quantitative Point-of-Care Platform for the Assessment of Drugs of Abuse in Urine, Serum, and Whole Blood. Analytical Chemistry, 2017, 89, 8273-8281.	6.5	28
54	Integrated analysis of long non-coding RNAs in human gastric cancer: An in silico study. PLoS ONE, 2017, 12, e0183517.	2.5	7

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55	A nomogram based on serum bilirubin and albumin levels predicts survival in gastric cancer patients. Oncotarget, 2017, 8, 41305-41318.	1.8	35
56	MicroRNA expression profiles predict progression and clinical outcome in lung adenocarcinoma. OncoTargets and Therapy, 2016, Volume 9, 5679-5692.	2.0	35
57	A systematic review on the association between the <i>Helicobacter pylori vacA i</i> genotype and gastric disease. FEBS Open Bio, 2016, 6, 409-417.	2.3	13
58	Nanoporous Glass Integrated in Volumetric Bar-Chart Chip for Point-of-Care Diagnostics of Non-Small Cell Lung Cancer. ACS Nano, 2016, 10, 1640-1647.	14.6	67
59	Circulating miR-148/152 family as potential biomarkers in hepatocellular carcinoma. Tumor Biology, 2016, 37, 4945-4953.	1.8	27
60	Different effects of the three polymorphisms on 15q25.1 onlung cancer risk: Evidence from published literatures. Journal of Cancer Research and Therapeutics, 2016, 12, 12.	0.9	3
61	Association of Clostridium difficile infection in hospital mortality: A systematic review and meta-analysis. American Journal of Infection Control, 2015, 43, 1316-1320.	2.3	26
62	Inhibition of CD147 expression by RNA interference reduces proliferation, invasion and increases chemosensitivity in cancer stem cell-like HT-29 cells. International Journal of Oncology, 2015, 47, 1476-1484.	3.3	9
63	Gene therapy for human colorectal cancer cell lines with recombinant adenovirus 5 based on loss of the insulin-like growth factor 2 imprinting. International Journal of Oncology, 2015, 46, 1759-1767.	3.3	12
64	Gene therapy for colorectal cancer by adenovirus-mediated siRNA targeting CD147 based on loss of the IGF2 imprinting system. International Journal of Oncology, 2015, 47, 1881-1889.	3.3	10
65	Associations of polymorphisms in microRNAs with female breast cancer risk in Chinese population. Tumor Biology, 2015, 36, 4575-4582.	1.8	44
66	Prognostic value of pre-operative inflammatory response biomarkers in gastric cancer patients and the construction of a predictive model. Journal of Translational Medicine, 2015, 13, 66.	4.4	172
67	Prognostic value of neutrophilâ€ŧoâ€ŀymphocyte ratio in breast cancer. FEBS Open Bio, 2015, 5, 502-507.	2.3	104
68	Long non oding RNA 91H contributes to the occurrence and progression of esophageal squamous cell carcinoma by inhibiting IGF2 expression. Molecular Carcinogenesis, 2015, 54, 359-367.	2.7	53
69	Association of the Polymorphisms in the Fas/FasL Promoter Regions with Cancer Susceptibility: A Systematic Review and Meta-Analysis of 52 Studies. PLoS ONE, 2014, 9, e90090.	2.5	34
70	Prognostic Value of Long Non-Coding RNA HOTAIR in Various Cancers. PLoS ONE, 2014, 9, e110059.	2.5	32
71	Increased CD59 protein expression is associated with the outcome of patients with diffuse large B-cell lymphoma treated with R-CHOP. Medical Oncology, 2014, 31, 56.	2.5	12
72	The diplotype Fas â~'1377A/â~'670G as a genetic marker to predict a lower risk of breast cancer in Chinese women. Tumor Biology, 2014, 35, 9147-9161.	1.8	8

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73	Up-Regulation of 91H Promotes Tumor Metastasis and Predicts Poor Prognosis for Patients with Colorectal Cancer. PLoS ONE, 2014, 9, e103022.	2.5	72
74	Deregulated Expression of miR-224 and its Target Gene: CD59 Predicts Outcome of Diffuse Large B-cell Lymphoma Patients Treated with R-CHOP. Current Cancer Drug Targets, 2014, 14, 659-670.	1.6	20
75	Differential effects of insulin-like growth factor-1 CA repeat polymorphism on breast cancer risk along with race: A meta-analysis. Gene, 2013, 525, 92-98.	2.2	8

The Association between Four Genetic Variants in MicroRNAs (rs11614913, rs2910164, rs3746444,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf

77	Interleukin 1 beta (IL1B) promoter polymorphism and cancer risk: evidence from 47 published studies. Mutagenesis, 2011, 26, 637-642.	2.6	44
78	Clinical Efficacy of Convalescent Plasma Therapy on Treating COVID-19 Patients: Evidence from Matched Study and a Meta-Analysis. SSRN Electronic Journal, 0, , .	0.4	0