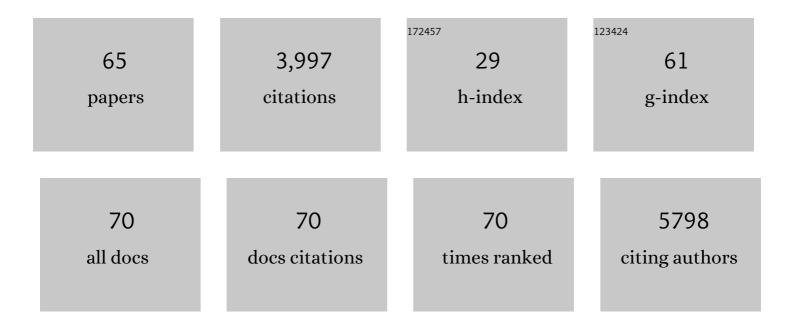
List of Publications by Year in descending order

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HULLING SUN

#	Article	IF	CITATIONS
1	SNHG15 is a negative regulator of inflammation by mediating TRAF2 ubiquitination in stroke-induced immunosuppression. Journal of Neuroinflammation, 2022, 19, 1.	7.2	28
2	Novel insights into the interaction between N6-methyladenosine modification and circular RNA. Molecular Therapy - Nucleic Acids, 2022, 27, 824-837.	5.1	19
3	Dual Antiplatelet Therapy in Patients With Minor Stroke Receiving Intravenous Thrombolysis. Frontiers in Neurology, 2022, 13, 819896.	2.4	4
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	Admission blood cell counts are predictive of stroke-associated infection in acute ischemic stroke patients treated with endovascular therapy. Neurological Sciences, 2021, 42, 2397-2409.	1.9	12
7	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
8	Laboratory Testing Implications of Risk-Stratification and Management of COVID-19 Patients. Frontiers in Medicine, 2021, 8, 699706.	2.6	3
9	Inflammatory Factors as Potential Markers of Early Neurological Deterioration in Acute Ischemic Stroke Patients Receiving Endovascular Therapy – The AISRNA Study. Journal of Inflammation Research, 2021, Volume 14, 4399-4407.	3.5	12
10	A Novel Nomogram for Predicting Prognosis after Mechanical Thrombectomy in Patients with Acute Ischemic Stroke. Current Neurovascular Research, 2021, 18, 479-488.	1.1	3
11	MiRâ€142â€3p functions as a tumor suppressor by targeting RAC1/PAK1 pathway in breast cancer. Journal of Cellular Physiology, 2020, 235, 4928-4940.	4.1	28
12	Susceptibility of <em>PON1</em> / <em>PON2</em> Genetic Variations to Ischemic Stroke Risk in a Chinese Han Population. Pharmacogenomics and Personalized Medicine, 2020, Volume 13, 563-570.	0.7	7
13	MiR-485-5p as a potential biomarker and tumor suppressor in human colorectal cancer. Biomarkers in Medicine, 2020, 14, 239-248.	1.4	20
14	Serum exosomal miR-122 as a potential diagnostic and prognostic biomarker of colorectal cancer with liver metastasis. Journal of Cancer, 2020, 11, 630-637.	2.5	65
15	Analysis of METTL3 and METTL14 in hepatocellular carcinoma. Aging, 2020, 12, 21638-21659.	3.1	44
16	<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
17	Identification of Serum Exosomal hsa-circ-0004771 as a Novel Diagnostic Biomarker of Colorectal Cancer. Frontiers in Genetics, 2019, 10, 1096.	2.3	157
18	lncRNA 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

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19	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
20	Circulating miR-1290 and miR-320d as Novel Diagnostic Biomarkers of Human Colorectal Cancer. Journal of Cancer, 2019, 10, 43-50.	2.5	53
21	Low triglyceride to high-density lipoprotein cholesterol ratio predicts hemorrhagic transformation in large atherosclerotic infarction of acute ischemic stroke. Aging, 2019, 11, 1589-1601.	3.1	16
22	Polymorphisms of IL-23R predict survival of gastric cancer patients in a Chinese population. Cytokine, 2019, 117, 79-83.	3.2	6
23	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
24	Triglyceride-to-high density lipoprotein cholesterol ratio predicts clinical outcomes in patients with gastric cancer. Journal of Cancer, 2019, 10, 6829-6836.	2.5	21
25	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
26	CircHIPK3 promotes colorectal cancer growth and metastasis by sponging miR-7. Cell Death and Disease, 2018, 9, 417.	6.3	497
27	Prognostic significance of long noncoding <scp>RNA</scp> Z38 as a candidate biomarker in breast cancer. Journal of Clinical Laboratory Analysis, 2018, 32, .	2.1	25
28	Polymorphisms of TGFBR1, TLR4 are associated with prognosis of gastric cancer in a Chinese population. Cancer Cell International, 2018, 18, 191.	4.1	21
29	The pro-metastasis effect of circANKS1B in breast cancer. Molecular Cancer, 2018, 17, 160.	19.2	219
30	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
31	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
32	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
33	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
34	microRNA-485-5p Functions as a Tumor Suppressor in Colorectal Cancer Cells by Targeting CD147. Journal of Cancer, 2018, 9, 2603-2611.	2.5	47
35	Differential long noncoding RNA expressions in peripheral blood mononuclear cells for detection of acute ischemic stroke. Clinical Science, 2018, 132, 1597-1614.	4.3	46
36	Serum and exosome long non coding RNAs as potential biomarkers for hepatocellular carcinoma. Journal of Cancer, 2018, 9, 2631-2639.	2.5	97

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37	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
38	miR-150-5p suppresses tumor progression by targeting VEGFA in colorectal cancer. Aging, 2018, 10, 3421-3437.	3.1	87
39	Meta-analysis of prognostic value of inflammation parameter in breast cancer. Journal of Cancer Research and Therapeutics, 2018, 14, S85-S89.	0.9	10
40	Combination of preoperative NLR, PLR and CEA could increase the diagnostic efficacy for I-III stage CRC. Journal of Clinical Laboratory Analysis, 2017, 31, e22075.	2.1	39
41	Association of Genetic Polymorphisms in the LncRNAs with Gastric Cancer Risk in a Chinese Population. Journal of Cancer, 2017, 8, 531-536.	2.5	30
42	The Predictive and Prognostic Role of Stromal Tumor-infiltrating Lymphocytes in HER2-positive Breast Cancer with Trastuzumab-based Treatment: a Meta-analysis and Systematic Review. Journal of Cancer, 2017, 8, 3838-3848.	2.5	16
43	Association between SNPs in Long Non-coding RNAs and the Risk of Female Breast Cancer in a Chinese Population. Journal of Cancer, 2017, 8, 1162-1169.	2.5	23
44	A nomogram based on serum bilirubin and albumin levels predicts survival in gastric cancer patients. Oncotarget, 2017, 8, 41305-41318.	1.8	35
45	Prognostic performance of lymphocyte-to-monocyte ratio in diffuse large B-cell lymphoma: an updated meta-analysis of eleven reports. OncoTargets and Therapy, 2016, 9, 3017.	2.0	16
46	Association of the DISC1 and NRG1 genetic polymorphisms with schizophrenia in a Chinese population. Gene, 2016, 590, 293-297.	2.2	18
47	The association of Phosphatase and tensin homolog (PTEN) deletion and prostate cancer risk: A meta-analysis. Biomedicine and Pharmacotherapy, 2016, 83, 114-121.	5.6	14
48	Circulating miR-148/152 family as potential biomarkers in hepatocellular carcinoma. Tumor Biology, 2016, 37, 4945-4953.	1.8	27
49	MiR-608, pre-miR-124-1 and pre-miR26a-1 polymorphisms modify susceptibility and recurrence-free survival in surgically resected CRC individuals. Oncotarget, 2016, 7, 75865-75873.	1.8	44
50	Nucleotide excision repair pathway gene polymorphisms are linked to breast cancer risk in a Chinese population. Oncotarget, 2016, 7, 84872-84882.	1.8	25
51	The effect of BIM deletion polymorphism on intrinsic resistance and clinical outcome of cancer patient with kinase inhibitor therapy. Scientific Reports, 2015, 5, 11348.	3.3	23
52	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
53	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
54	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

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55	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
56	Estrogen Receptor 1 (ESR1) genetic variations in cancer risk: A systematic review and meta-analysis. Clinics and Research in Hepatology and Gastroenterology, 2015, 39, 127-135.	1.5	11
57	Circulating vitamin D binding protein, total, free and bioavailable 25-hydroxyvitamin D and risk of colorectal cancer. Scientific Reports, 2015, 5, 7956.	3.3	38
58	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
59	Evaluation the susceptibility of five polymorphisms in microRNA-binding sites to female breast cancer risk in Chinese population. Gene, 2015, 573, 160-165.	2.2	11
60	Prognostic value of neutrophilâ€ŧo″ymphocyte ratio in breast cancer. FEBS Open Bio, 2015, 5, 502-507.	2.3	104
61	Upregulated IncRNA-UCA1 contributes to progression of hepatocellular carcinoma through inhibition of miR-216b and activation of FGFR1/ERK signaling pathway. Oncotarget, 2015, 6, 7899-7917.	1.8	329
62	Association between estrogen receptor 1 (ESR1) genetic variations and cancer risk: a meta-analysis. Journal of B U on, 2015, 20, 296-308.	0.4	12
63	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
64	Prognostic Value of Long Non-Coding RNA HOTAIR in Various Cancers. PLoS ONE, 2014, 9, e110059.	2.5	32
65	Up-Regulation of 91H Promotes Tumor Metastasis and Predicts Poor Prognosis for Patients with Colorectal Cancer, PLoS ONF, 2014, 9, e103022	2.5	72