Yong Li

List of Publications by Year in descending order

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107	10,894	52	100
	citations	h-index	g-index
papers	Citations	II-IIIQEX	g-muex
111	111	111	13439
all docs	docs citations	times ranked	citing authors

#	Article	lF	Citations
1	Role of the tumor microenvironment in PD-L1/PD-1-mediated tumor immune escape. Molecular Cancer, $2019, 18, 10.$	19.2	810
2	Circular RNAs function as ceRNAs to regulate and control human cancer progression. Molecular Cancer, 2018, 17, 79.	19.2	757
3	MYC/BCL2 protein coexpression contributes to the inferior survival of activated B-cell subtype of diffuse large B-cell lymphoma and demonstrates high-risk gene expression signatures: a report from The International DLBCL Rituximab-CHOP Consortium Program. Blood, 2013, 121, 4021-4031.	1.4	596
4	MicroRNAs in NF-ÂB signaling. Journal of Molecular Cell Biology, 2011, 3, 159-166.	3.3	530
5	Neoantigen vaccine: an emerging tumor immunotherapy. Molecular Cancer, 2019, 18, 128.	19.2	398
6	Emerging role of tumor-related functional peptides encoded by lncRNA and circRNA. Molecular Cancer, 2020, 19, 22.	19.2	330
7	Circular RNAs in human cancer. Molecular Cancer, 2017, 16, 25.	19.2	310
8	Role of metabolism in cancer cell radioresistance and radiosensitization methods. Journal of Experimental and Clinical Cancer Research, 2018, 37, 87.	8.6	288
9	Targeting the IDO1 pathway in cancer: from bench to bedside. Journal of Hematology and Oncology, 2018, 11, 100.	17.0	277
10	Circular RNAs (circRNAs) in cancer. Cancer Letters, 2018, 425, 134-142.	7.2	229
11	Apoptosis and the target genes of microRNA-21. Chinese Journal of Cancer, 2011, 30, 371-380.	4.9	227
12	miR-301a as an NF-κB activator in pancreatic cancer cells. EMBO Journal, 2011, 30, 57-67.	7.8	204
13	Loss of the <i>miR-21</i> allele elevates the expression of its target genes and reduces tumorigenesis. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 10144-10149.	7.1	202
14	Apoptosis and the target genes of microRNA-21. Chinese Journal of Cancer, 2011, 30, 371-380.	4.9	197
15	Predictive biomarkers and mechanisms underlying resistance to PD1/PD-L1 blockade cancer immunotherapy. Molecular Cancer, 2020, 19, 19.	19.2	180
16	Mechanisms of vasculogenic mimicry in hypoxic tumor microenvironments. Molecular Cancer, 2021, 20, 7.	19.2	177
17	Natural killer group 2D receptor and its ligands in cancer immune escape. Molecular Cancer, 2019, 18, 29.	19.2	149
18	Long noncoding RNA AFAP1-AS1 acts as a competing endogenous RNA of miR-423-5p to facilitate nasopharyngeal carcinoma metastasis through regulating the Rho/Rac pathway. Journal of Experimental and Clinical Cancer Research, 2018, 37, 253.	8.6	148

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19	Long non-coding RNA PVT1 predicts poor prognosis and induces radioresistance by regulating DNA repair and cell apoptosis in nasopharyngeal carcinoma. Cell Death and Disease, 2018, 9, 235.	6.3	143
20	Patients with diffuse large B-cell lymphoma of germinal center origin with BCL2 translocations have poor outcome, irrespective of MYC status: a report from an International DLBCL rituximab-CHOP Consortium Program Study. Haematologica, 2013, 98, 255-263.	3.5	142
21	Genetic alterations and their clinical implications in DLBCL. Nature Reviews Clinical Oncology, 2019, 16, 634-652.	27.6	136
22	LncRNAs regulate the cytoskeleton and related Rho/ROCK signaling in cancer metastasis. Molecular Cancer, 2018, 17, 77.	19.2	131
23	Singleâ€cell RNA sequencing in cancer research. Journal of Experimental and Clinical Cancer Research, 2021, 40, 81.	8.6	128
24	Application of atomic force microscopy in cancer research. Journal of Nanobiotechnology, 2018, 16, 102.	9.1	127
25	Effects of tumor metabolic microenvironment on regulatory T cells. Molecular Cancer, 2018, 17, 168.	19.2	119
26	Epstein-Barr virus-encoded miR-BART6-3p inhibits cancer cell metastasis and invasion by targeting long non-coding RNA LOC553103. Cell Death and Disease, 2016, 7, e2353-e2353.	6.3	118
27	Co-expression of AFAP1-AS1 and PD-1 predicts poor prognosis in nasopharyngeal carcinoma. Oncotarget, 2017, 8, 39001-39011.	1.8	114
28	Immune Profiling and Quantitative Analysis Decipher the Clinical Role of Immune-Checkpoint Expression in the Tumor Immune Microenvironment of DLBCL. Cancer Immunology Research, 2019, 7, 644-657.	3.4	106
29	Upregulated long non-coding RNA LINC00152 expression is associated with progression and poor prognosis of tongue squamous cell carcinoma. Journal of Cancer, 2017, 8, 523-530.	2.5	105
30	Expression of LINC00312, a long intergenic non-coding RNA, is negatively correlated with tumor size but positively correlated with lymph node metastasis in nasopharyngeal carcinoma. Journal of Molecular Histology, 2013, 44, 545-554.	2.2	104
31	miR-153 suppresses IDO1 expression and enhances CAR T cell immunotherapy. Journal of Hematology and Oncology, 2018, 11, 58.	17.0	98
32	<i>circ<scp>MAN</scp>1A2</i> could serve as a novel serum biomarker for malignant tumors. Cancer Science, 2019, 110, 2180-2188.	3.9	96
33	LOC401317, a p53-Regulated Long Non-Coding RNA, Inhibits Cell Proliferation and Induces Apoptosis in the Nasopharyngeal Carcinoma Cell Line HNE2. PLoS ONE, 2014, 9, e110674.	2.5	93
34	Overexpression long non-coding RNA <i>LINC00673 </i> is associated with poor prognosis and promotes invasion and metastasis in tongue squamous cell carcinoma. Oncotarget, 2017, 8, 16621-16632.	1.8	92
35	<i>MYD88</i> L265P Mutation in Lymphoid Malignancies. Cancer Research, 2018, 78, 2457-2462.	0.9	92
36	miR-21 depletion in macrophages promotes tumoricidal polarization and enhances PD-1 immunotherapy. Oncogene, 2018, 37, 3151-3165.	5.9	90

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37	CTLA-4 in Regulatory T Cells for Cancer Immunotherapy. Cancers, 2021, 13, 1440.	3.7	88
38	KRAS/NF-κB/YY1/miR-489 Signaling Axis Controls Pancreatic Cancer Metastasis. Cancer Research, 2017, 77, 100-111.	0.9	86
39	The emerging role of Epstein-Barr virus encoded microRNAs in nasopharyngeal carcinoma. Journal of Cancer, 2018, 9, 2852-2864.	2.5	83
40	The role of Wnt signaling pathway in tumor metabolic reprogramming. Journal of Cancer, 2019, 10, 3789-3797.	2.5	80
41	The oncogenic microRNA miR-21 promotes regulated necrosis in mice. Nature Communications, 2015, 6, 7151.	12.8	78
42	Epstein-Barr virus encoded miR-BART11 promotes inflammation-induced carcinogenesis by targeting FOXP1. Oncotarget, 2016, 7, 36783-36799.	1.8	78
43	Identification of genomic alterations in nasopharyngeal carcinoma and nasopharyngeal carcinoma-derived Epstein–Barr virus by whole-genome sequencing. Carcinogenesis, 2018, 39, 1517-1528.	2.8	74
44	BPIFB1 (LPLUNC1) inhibits migration and invasion of nasopharyngeal carcinoma by interacting with VTN and VIM. British Journal of Cancer, 2018, 118, 233-247.	6.4	73
45	Genome-Wide Analysis of 18 Epstein-Barr Viruses Isolated from Primary Nasopharyngeal Carcinoma Biopsy Specimens. Journal of Virology, 2017, 91, .	3.4	70
46	BPIFB1 (LPLUNC1) inhibits radioresistance in nasopharyngeal carcinoma by inhibiting VTN expression. Cell Death and Disease, 2018, 9, 432.	6.3	70
47	Regulation network and expression profiles of Epstein-Barr virus-encoded microRNAs and their potential target host genes in nasopharyngeal carcinomas. Science China Life Sciences, 2014, 57, 315-326.	4.9	66
48	Long non-coding RNAs in cancer. Science China Life Sciences, 2012, 55, 1120-1124.	4.9	65
49	Epstein–Barr Virus–Encoded Circular RNA CircBART2.2 Promotes Immune Escape of Nasopharyngeal Carcinoma by Regulating PD-L1. Cancer Research, 2021, 81, 5074-5088.	0.9	65
50	An integrative transcriptomic analysis reveals p53 regulated miRNA, mRNA, and lncRNA networks in nasopharyngeal carcinoma. Tumor Biology, 2016, 37, 3683-3695.	1.8	61
51	Interaction of the oncogenic miR-21 microRNA and the p53 tumor suppressor pathway. Carcinogenesis, 2013, 34, 1216-1223.	2.8	60
52	Circular RNA circRNF13 inhibits proliferation and metastasis of nasopharyngeal carcinoma via SUMO2. Molecular Cancer, 2021, 20, 112.	19.2	60
53	Assessment of CD37 B-cell antigen and cell of origin significantly improves risk prediction in diffuse large B-cell lymphoma. Blood, 2016, 128, 3083-3100.	1.4	59
54	High Expression of LINC01420 indicates an unfavorable prognosis and modulates cell migration and invasion in nasopharyngeal carcinoma. Journal of Cancer, 2017, 8, 97-103.	2.5	59

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55	LncRNAs regulate cancer metastasis via binding to functional proteins. Oncotarget, 2018, 9, 1426-1443.	1.8	55
56	Long non-coding RNAs are involved in alternative splicing and promote cancer progression. British Journal of Cancer, 2022, 126, 1113-1124.	6.4	53
57	LncRNA AATBC regulates Pinin to promote metastasis in nasopharyngeal carcinoma. Molecular Oncology, 2020, 14, 2251-2270.	4.6	52
58	EBV miRNAs BART11 and BART17-3p promote immune escape through the enhancer-mediated transcription of PD-L1. Nature Communications, 2022, 13, 866.	12.8	51
59	Long non-coding RNA LOC284454 promotes migration and invasion of nasopharyngeal carcinoma via modulating the Rho/Rac signaling pathway. Carcinogenesis, 2019, 40, 380-391.	2.8	49
60	Clinical features, tumor biology, and prognosis associated with MYC rearrangement and Myc overexpression in diffuse large B-cell lymphoma patients treated with rituximab-CHOP. Modern Pathology, 2015, 28, 1555-1573.	5.5	48
61	Clinical and Biologic Significance of <i>MYC</i> Genetic Mutations in <i>De Novo</i> Diffuse Large B-cell Lymphoma. Clinical Cancer Research, 2016, 22, 3593-3605.	7.0	48
62	Targeting PD-L1 in non-small cell lung cancer using CAR T cells. Oncogenesis, 2020, 9, 72.	4.9	48
63	The influence of circular RNAs on autophagy and disease progression. Autophagy, 2022, 18, 240-253.	9.1	48
64	LPLUNC1 Inhibits Nasopharyngeal Carcinoma Cell Growth via Down-Regulation of the MAP Kinase and Cyclin D1/E2F Pathways. PLoS ONE, 2013, 8, e62869.	2.5	47
65	The role of exosomal non-coding RNAs in cancer metastasis. Oncotarget, 2018, 9, 12487-12502.	1.8	47
66	Emerging role of metabolic reprogramming in tumor immune evasion and immunotherapy. Science China Life Sciences, 2021, 64, 534-547.	4.9	47
67	CircARHGAP12 promotes nasopharyngeal carcinoma migration and invasion via ezrin-mediated cytoskeletal remodeling. Cancer Letters, 2021, 496, 41-56.	7.2	46
68	Single nucleotide variation in the TP53 3′ untranslated region in diffuse large B-cell lymphoma treated with rituximab-CHOP: a report from the International DLBCL Rituximab-CHOP Consortium Program. Blood, 2013, 121, 4529-4540.	1.4	41
69	miR-301a promotes lung tumorigenesis by suppressing Runx3. Molecular Cancer, 2019, 18, 99.	19.2	39
70	Long non-coding RNA AFAP1-AS1 accelerates lung cancer cells migration and invasion by interacting with SNIP1 to upregulate c-Myc. Signal Transduction and Targeted Therapy, 2021, 6, 240.	17.1	39
71	Cloning and characterization of the putative AFAP1-AS1 promoter region. Journal of Cancer, 2019, 10, 1145-1153.	2.5	37
72	miR-25 Promotes Cell Proliferation, Migration, and Invasion of Non-Small-Cell Lung Cancer by Targeting the LATS2/YAP Signaling Pathway. Oxidative Medicine and Cellular Longevity, 2019, 2019, 1-14.	4.0	36

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73	Modulation of tumorigenesis by the pro-inflammatory microRNA miR-301a in mouse models of lung cancer and colorectal cancer. Cell Discovery, 2015, 1, 15005.	6.7	34
74	Epsteinâ€Barr virusâ€encoded miRâ€BART6â€3p inhibits cancer cell proliferation through the LOC553103â€STM axis. FASEB Journal, 2020, 34, 8012-8027.	N1 0.5	34
75	What are the applications of single-cell RNA sequencing in cancer research: a systematic review. Journal of Experimental and Clinical Cancer Research, 2021, 40, 163.	8.6	33
76	Abnormal X chromosome inactivation and tumor development. Cellular and Molecular Life Sciences, 2020, 77, 2949-2958.	5.4	32
77	Proteomic Analysis of the Molecular Mechanism of Lovastatin Inhibiting the Growth of Nasopharyngeal Carcinoma Cells. Journal of Cancer, 2019, 10, 2342-2349.	2.5	31
78	Herpesvirus acts with the cytoskeleton and promotes cancer progression. Journal of Cancer, 2019, 10, 2185-2193.	2.5	31
79	Prognostic factors, therapeutic approaches, and distinct immunobiologic features in patients with primary mediastinal large B-cell lymphoma on long-term follow-up. Blood Cancer Journal, 2020, 10, 49.	6.2	31
80	Long non-coding RNA AFAP1-AS1 is a novel biomarker in various cancers: a systematic review and meta-analysis based on the literature and GEO datasets. Oncotarget, 2017, 8, 102346-102360.	1.8	30
81	Therapeutic cancer vaccines: From biological mechanisms and engineering to ongoing clinical trials. Cancer Treatment Reviews, 2022, 109, 102429.	7.7	30
82	Upregulation of long non-coding RNA LOC284454 may serve as a new serum diagnostic biomarker for head and neck cancers. BMC Cancer, 2020, 20, 917.	2.6	28
83	Cancer/testis antigens: from serology to mRNA cancer vaccine. Seminars in Cancer Biology, 2021, 76, 218-231.	9.6	27
84	N6-methyladenosine-dependent signalling in cancer progression and insights into cancer therapies. Journal of Experimental and Clinical Cancer Research, 2021, 40, 146.	8.6	26
85	Glyphosate induces benign monoclonal gammopathy and promotes multiple myeloma progression in mice. Journal of Hematology and Oncology, 2019, 12, 70.	17.0	25
86	Leucovorin Enhances the Anti-cancer Effect of Bortezomib in Colorectal Cancer Cells. Scientific Reports, 2017, 7, 682.	3.3	24
87	Immunoglobulin somatic hypermutation has clinical impact in DLBCL and potential implications for immune checkpoint blockade and neoantigen-based immunotherapies. , 2019, 7, 272.		22
88	A refined cell-of-origin classifier with targeted NGS and artificial intelligence shows robust predictive value in DLBCL. Blood Advances, 2020, 4, 3391-3404.	5.2	22
89	The long noncoding RNA AATBC promotes breast cancer migration and invasion by interacting with YBX1 and activating the YAP1/Hippo signaling pathway. Cancer Letters, 2021, 512, 60-72.	7.2	22
90	Genetic Subtyping and Phenotypic Characterization of the Immune Microenvironment and MYC/BCL2 Double Expression Reveal Heterogeneity in Diffuse Large B-cell Lymphoma. Clinical Cancer Research, 2022, 28, 972-983.	7.0	22

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91	Malignant Transformation of Human Bronchial Epithelial Cells Induced by Arsenic through STAT3/miR-301a/SMAD4 Loop. Scientific Reports, 2018, 8, 13291.	3.3	21
92	MYD88 L265P elicits mutation-specific ubiquitination to drive NF- \hat{I}^0B activation and lymphomagenesis. Blood, 2021, 137, 1615-1627.	1.4	21
93	Cancer testis antigen SPAG9 is a promising marker for the diagnosis and treatment of lung cancer. Oncology Reports, 2016, 35, 2599-2605.	2.6	20
94	EBVâ€miRâ€BART12 accelerates migration and invasion in EBVâ€associated cancer cells by targeting tubulin polymerizationâ€promoting protein 1. FASEB Journal, 2020, 34, 16205-16223.	0.5	19
95	Tissue-specific microRNA expression alters cancer susceptibility conferred by a TP53 noncoding variant. Nature Communications, 2019, 10, 5061.	12.8	18
96	A high-throughput screening identifies histone deacetylase inhibitors as therapeutic agents against medulloblastoma. Experimental Hematology and Oncology, 2019, 8, 30.	5.0	17
97	Ubiquitination of the DNA-damage checkpoint kinase CHK1 by TRAF4 is required for CHK1 activation. Journal of Hematology and Oncology, 2020, 13, 40.	17.0	16
98	Expanding anti-CD38 immunotherapy for lymphoid malignancies. Journal of Experimental and Clinical Cancer Research, 2022, 41, .	8.6	15
99	MR1-Restricted T Cells in Cancer Immunotherapy. Cancers, 2020, 12, 2145.	3.7	13
100	AFAP1-AS1: a rising star among oncogenic long non-coding RNAs. Science China Life Sciences, 2021, 64, 1602-1611.	4.9	11
101	Posttranslational Modifications in PD-L1 Turnover and Function: From Cradle to Grave. Biomedicines, 2021, 9, 1702.	3.2	11
102	2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD) induces peripheral blood abnormalities and plasma cell neoplasms resembling multiple myeloma in mice. Cancer Letters, 2019, 440-441, 135-144.	7.2	10
103	Determining clinical course of diffuse large B-cell lymphoma using targeted transcriptome and machine learning algorithms. Blood Cancer Journal, 2022, 12, 25.	6.2	7
104	Genomic complexity is associated with epigenetic regulator mutations and poor prognosis in diffuse large B-cell lymphoma. Oncolmmunology, 2021, 10, 1928365.	4.6	6
105	Long nonâ€'coding RNA expression profiles and related regulatory networks in areca nut chewingâ€'induced tongue squamous cell carcinoma. Oncology Letters, 2020, 20, 1-1.	1.8	4
106	Long non-coding RNA expression profiles and related regulatory networks in areca nut chewing-induced tongue squamous cell carcinoma. Oncology Letters, 2020, 20, 302.	1.8	3
107	The role of alternative splicing in human cancer progression. American Journal of Cancer Research, 2021, 11, 4642-4667.	1.4	3