

Jiachun Lu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5358189/publications.pdf>

Version: 2024-02-01

127
papers

5,713
citations

87723

38
h-index

88477

70
g-index

133
all docs

133
docs citations

133
times ranked

8787
citing authors

#	ARTICLE	IF	CITATIONS
1	Circular RNA ITCH has inhibitory effect on ESCC by suppressing the Wnt/ β -catenin pathway. <i>Oncotarget</i> , 2015, 6, 6001-6013.	0.8	626
2	Prevalence of Chronic Obstructive Pulmonary Disease in China. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2007, 176, 753-760.	2.5	600
3	A genome-wide association study identifies two new lung cancer susceptibility loci at 13q12.12 and 22q12.2 in Han Chinese. <i>Nature Genetics</i> , 2011, 43, 792-796.	9.4	340
4	Increased Levels of the Long Intergenic Non-coding Protein Coding RNA POU3F3 Promote DNA Methylation in Esophageal Squamous Cell Carcinoma Cells. <i>Gastroenterology</i> , 2014, 146, 1714-1726.e5.	0.6	169
5	Micropeptide <sc>CIP</sc> 2A encoded by <sc>LINC</sc> 00665 inhibits triple-negative breast cancer progression. <i>EMBO Journal</i> , 2020, 39, e102190.	3.5	138
6	LncRNA-encoded polypeptide ASRPS inhibits triple-negative breast cancer angiogenesis. <i>Journal of Experimental Medicine</i> , 2020, 217, .	4.2	136
7	Changes in Electronic Cigarette Use Among Adults in the United States, 2014-2016. <i>JAMA - Journal of the American Medical Association</i> , 2018, 319, 2039.	3.8	118
8	COVID-19: asymptomatic carrier transmission is an underestimated problem. <i>Epidemiology and Infection</i> , 2020, 148, e116.	1.0	114
9	Overexpression of lncRNA IGFBP4-1 reprograms energy metabolism to promote lung cancer progression. <i>Molecular Cancer</i> , 2017, 16, 154.	7.9	103
10	Exosomal FMR1-AS1 facilitates maintaining cancer stem-like cell dynamic equilibrium via TLR7/NF κ B/c-Myc signaling in female esophageal carcinoma. <i>Molecular Cancer</i> , 2019, 18, 22.	7.9	93
11	The evidence of indirect transmission of SARS-CoV-2 reported in Guangzhou, China. <i>BMC Public Health</i> , 2020, 20, 1202.	1.2	91
12	A Novel Micropeptide Encoded by Y-Linked LINC00278 Links Cigarette Smoking and AR Signaling in Male Esophageal Squamous Cell Carcinoma. <i>Cancer Research</i> , 2020, 80, 2790-2803.	0.4	91
13	Epigenetic silencing of O6-methylguanine DNA methyltransferase gene in NiS-transformed cells. <i>Carcinogenesis</i> , 2008, 29, 1267-1275.	1.3	86
14	A functional polymorphism at microRNA-629-binding site in the 3'-untranslated region of NBS1 gene confers an increased risk of lung cancer in Southern and Eastern Chinese population. <i>Carcinogenesis</i> , 2012, 33, 338-347.	1.3	80
15	Genetic Variants at 6p21.1 and 7p15.3 Are Associated with Risk of Multiple Cancers in Han Chinese. <i>American Journal of Human Genetics</i> , 2012, 91, 928-934.	2.6	76
16	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016, 7, 11478.	5.8	73
17	Community based integrated intervention for prevention and management of chronic obstructive pulmonary disease (COPD) in Guangdong, China: cluster randomised controlled trial. <i>BMJ: British Medical Journal</i> , 2010, 341, c6387-c6387.	2.4	65
18	MicroRNA-152 targets DNA methyltransferase 1 in NiS-transformed cells via a feedback mechanism. <i>Carcinogenesis</i> , 2013, 34, 446-453.	1.3	65

#	ARTICLE	IF	CITATIONS
19	A Sequence Polymorphism in <i>miR-608</i> Predicts Recurrence after Radiotherapy for Nasopharyngeal Carcinoma. <i>Cancer Research</i> , 2013, 73, 5151-5162.	0.4	64
20	A Functional Copy-Number Variation in MAPKAPK2 Predicts Risk and Prognosis of Lung Cancer. <i>American Journal of Human Genetics</i> , 2012, 91, 384-390.	2.6	61
21	A genetic polymorphism in <i>lincRNA-uc003opf.1</i> is associated with susceptibility to esophageal squamous cell carcinoma in Chinese populations. <i>Carcinogenesis</i> , 2013, 34, 2908-2917.	1.3	61
22	NF κ B1 and NF κ BIA Polymorphisms Are Associated with Increased Risk for Sporadic Colorectal Cancer in a Southern Chinese Population. <i>PLoS ONE</i> , 2011, 6, e21726.	1.1	59
23	The polymorphism and haplotypes of PIN1 gene are associated with the risk of lung cancer in southern and eastern chinese populations. <i>Human Mutation</i> , 2011, 32, 1299-1308.	1.1	59
24	Polymorphisms and haplotypes of the NBS1 gene are associated with risk of sporadic breast cancer in non-Hispanic white women ≤ 55 years. <i>Carcinogenesis</i> , 2006, 27, 2209-2216.	1.3	58
25	A novel functional variant (-842C>G) in the PIN1 promoter contributes to decreased risk of squamous cell carcinoma of the head and neck by diminishing the promoter activity. <i>Carcinogenesis</i> , 2009, 30, 1717-1721.	1.3	58
26	A functional copy number variation in the WWOX gene is associated with lung cancer risk in Chinese. <i>Human Molecular Genetics</i> , 2013, 22, 1886-1894.	1.4	58
27	Overdiagnosis of COPD in Subjects With Unobstructed Spirometry. <i>Chest</i> , 2019, 156, 277-288.	0.4	57
28	Promoter polymorphism (T>C) in the endothelial nitric oxide synthase gene is associated with risk of sporadic breast cancer in non-Hispanic white women age younger than 55 years. <i>Cancer</i> , 2006, 107, 2245-2253.	2.0	56
29	Functional genetic variations in the IL-23 receptor gene are associated with risk of breast, lung and nasopharyngeal cancer in Chinese populations. <i>Carcinogenesis</i> , 2012, 33, 2409-2416.	1.3	55
30	Long non-coding RNA LINC00672 contributes to p53 protein-mediated gene suppression and promotes endometrial cancer chemosensitivity. <i>Journal of Biological Chemistry</i> , 2017, 292, 5801-5813.	1.6	54
31	Identification of Three Circular RNA Cargoes in Serum Exosomes as Diagnostic Biomarkers of Non-Small-Cell Lung Cancer in the Chinese Population. <i>Journal of Molecular Diagnostics</i> , 2020, 22, 1096-1108.	1.2	54
32	CD44 rs13347 C>T polymorphism predicts breast cancer risk and prognosis in Chinese populations. <i>Breast Cancer Research</i> , 2012, 14, R105.	2.2	52
33	Functional <i>NBS1</i> polymorphism is associated with occurrence and advanced disease status of nasopharyngeal carcinoma. <i>Molecular Carcinogenesis</i> , 2011, 50, 689-696.	1.3	48
34	MicroRNA-218 acts by repressing TNFR1-mediated activation of NF κ B, which is involved in MUC5AC hyper-production and inflammation in smoking-induced bronchiolitis of COPD. <i>Toxicology Letters</i> , 2017, 280, 171-180.	0.4	46
35	Association between Chronic Obstructive Pulmonary Disease and Lung Cancer: A Case-Control Study in Southern Chinese and a Meta-Analysis. <i>PLoS ONE</i> , 2012, 7, e46144.	1.1	46
36	172G>T variant in the 5' untranslated region of DNA repair gene RAD51 reduces risk of squamous cell carcinoma of the head and neck and interacts with a P53 codon 72 variant. <i>Carcinogenesis</i> , 2006, 28, 988-994.	1.3	45

#	ARTICLE	IF	CITATIONS
37	A non-synonymous polymorphism Thr115Met in the EpCAM gene is associated with an increased risk of breast cancer in Chinese population. <i>Breast Cancer Research and Treatment</i> , 2011, 126, 487-495.	1.1	45
38	The Association between BMI and COPD: The Results of Two Population-based Studies in Guangzhou, China. <i>COPD: Journal of Chronic Obstructive Pulmonary Disease</i> , 2013, 10, 567-572.	0.7	45
39	A functional variant (â~1304T>C) in the MKK4 promoter contributes to a decreased risk of lung cancer by increasing the promoter activity. <i>Carcinogenesis</i> , 2010, 31, 1405-1411.	1.3	40
40	Feedback circuitry via let-7c between lncRNA CCAT1 and c-Myc is involved in cigarette smoke extract-induced malignant transformation of HBE cells. <i>Oncotarget</i> , 2017, 8, 19285-19297.	0.8	39
41	Sequence Variation in Mature MicroRNA-499 Confers Unfavorable Prognosis of Lung Cancer Patients Treated with Platinum-Based Chemotherapy. <i>Clinical Cancer Research</i> , 2015, 21, 1602-1613.	3.2	37
42	MicroRNA-191, regulated by HIF-2Î±, is involved in EMT and acquisition of a stem cell-like phenotype in arsenite-transformed human liver epithelial cells. <i>Toxicology in Vitro</i> , 2018, 48, 128-136.	1.1	37
43	Association between the NBS1E185Q polymorphism and cancer risk: a meta-analysis. <i>BMC Cancer</i> , 2009, 9, 124.	1.1	36
44	Identification of chimeric TSNAXâ€“DISC1 resulting from intergenic splicing in endometrial carcinoma through high-throughput RNA sequencing. <i>Carcinogenesis</i> , 2014, 35, 2687-2697.	1.3	36
45	CircLRP6 Regulation of ZEB1 via miR-455 Is Involved in the Epithelial-Mesenchymal Transition During Arsenite-Induced Malignant Transformation of Human Keratinocytes. <i>Toxicological Sciences</i> , 2018, 162, 450-461.	1.4	36
46	Effects of a Functional Variant c.353T>C in <i>Snai1</i> on Risk of Two Contextual Diseases. Chronic Obstructive Pulmonary Disease and Lung Cancer. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2014, 189, 139-148.	2.5	36
47	Functional Polymorphisms of CHRNA3 Predict Risks of Chronic Obstructive Pulmonary Disease and Lung Cancer in Chinese. <i>PLoS ONE</i> , 2012, 7, e46071.	1.1	36
48	A Polymorphism rs12325489C>T in the LincRNA-ENST00000515084 Exon Was Found to Modulate Breast Cancer Risk via GWAS-Based Association Analyses. <i>PLoS ONE</i> , 2014, 9, e98251.	1.1	36
49	The functional promoter polymorphism (â~842G>C) in the PIN1 gene is associated with decreased risk of breast cancer in non-Hispanic white women 55Ayears and younger. <i>Breast Cancer Research and Treatment</i> , 2010, 122, 243-249.	1.1	35
50	A microRNA-135a/b binding polymorphism in CD133 confers decreased risk and favorable prognosis of lung cancer in Chinese by reducing CD133 expression. <i>Carcinogenesis</i> , 2013, 34, 2292-2299.	1.3	34
51	Polymorphism in mature microRNA-608 sequence is associated with an increased risk of nasopharyngeal carcinoma. <i>Gene</i> , 2015, 565, 180-186.	1.0	34
52	Chronic obstructive pulmonary disease in the absence of chronic bronchitis in China. <i>Respirology</i> , 2010, 15, 1072-1078.	1.3	32
53	The protective role of polymorphism <i>MKK4â€“/i> 1304 T>C in nasopharyngeal carcinoma is modulated by Epsteinâ€“Barr virus' infection status. <i>International Journal of Cancer</i> , 2012, 130, 1981-1990.	2.3	32
54	Functional polymorphisms in NFÎ±B1/Î±BÎ± predict risks of chronic obstructive pulmonary disease and lung cancer in Chinese. <i>Human Genetics</i> , 2013, 132, 451-460.	1.8	31

#	ARTICLE	IF	CITATIONS
55	Upregulation of long non-coding RNA RAB1A-2 induces FGF1 expression worsening lung cancer prognosis. <i>Cancer Letters</i> , 2018, 438, 116-125.	3.2	31
56	Analysis of Survival-Related lncRNA Landscape Identifies A Role for LINC01537 in Energy Metabolism and Lung Cancer Progression. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3713.	1.8	31
57	The bioinformatics tools for the genome assembly and analysis based on third-generation sequencing. <i>Briefings in Functional Genomics</i> , 2019, 18, 1-12.	1.3	31
58	Association between the Cytotoxic T-Lymphocyte Antigen 4 +49G > A polymorphism and cancer risk: a meta-analysis. <i>BMC Cancer</i> , 2010, 10, 522.	1.1	30
59	A genome-wide gene-gene interaction analysis identifies an epistatic gene pair for lung cancer susceptibility in Han Chinese. <i>Carcinogenesis</i> , 2014, 35, 572-577.	1.3	29
60	Long non-coding RNA <i>AGER-1</i> functionally upregulates the innate immunity gene <i>AGER</i> and approximates its anti-tumor effect in lung cancer. <i>Molecular Carcinogenesis</i> , 2018, 57, 305-318.	1.3	29
61	Impaired autophagic flux and p62-mediated EMT are involved in arsenite-induced transformation of L-02 cells. <i>Toxicology and Applied Pharmacology</i> , 2017, 334, 75-87.	1.3	28
62	Polymorphisms of cytosolic serine hydroxymethyltransferase and risk of lung cancer: A case-control analysis. <i>Lung Cancer</i> , 2007, 57, 143-151.	0.9	26
63	Heterozygous Genetic Variations of <i>FOXP3</i> in Xp11.23 Elevate Breast Cancer Risk in Chinese Population via Skewed X-Chromosome Inactivation. <i>Human Mutation</i> , 2013, 34, n/a-n/a.	1.1	26
64	Risk factors shared by COPD and lung cancer and mediation effect of COPD: two center case-control studies. <i>Cancer Causes and Control</i> , 2015, 26, 11-24.	0.8	26
65	Exposure to Polycyclic Aromatic Hydrocarbons, Plasma Cytokines and Heart Rate Variability. <i>Scientific Reports</i> , 2016, 6, 19272.	1.6	25
66	Genetic variants in the H2AFX promoter region are associated with risk of sporadic breast cancer in non-Hispanic white women aged ≥55 years. <i>Breast Cancer Research and Treatment</i> , 2008, 110, 357-366.	1.1	24
67	IL-21 gene polymorphism is associated with the prognosis of breast cancer in Chinese populations. <i>Breast Cancer Research and Treatment</i> , 2013, 137, 893-901.	1.1	23
68	The association between T304T>G polymorphism in the promoter of <i>MKK4</i> gene and the risk of sporadic colorectal cancer in southern Chinese population. <i>International Journal of Cancer</i> , 2009, 125, 1876-1883.	2.3	22
69	A Common Genetic Variant (97906C>A) of DAB2IP/AIP1 Is Associated with an Increased Risk and Early Onset of Lung Cancer in Chinese Males. <i>PLoS ONE</i> , 2011, 6, e26944.	1.1	21
70	Functional polymorphisms in PIN1 promoter and esophageal carcinoma susceptibility in Chinese population. <i>Molecular Biology Reports</i> , 2013, 40, 829-838.	1.0	21
71	NF-κB-regulated miR-155, via repression of QKI, contributes to the acquisition of CSC-like phenotype during the neoplastic transformation of hepatic cells induced by arsenite. <i>Molecular Carcinogenesis</i> , 2018, 57, 483-493.	1.3	21
72	Duplicated copy of CHRNA7 increases risk and worsens prognosis of COPD and lung cancer. <i>European Journal of Human Genetics</i> , 2015, 23, 1019-1024.	1.4	20

#	ARTICLE	IF	CITATIONS
73	Involvement of HIF-1 α -regulated miR-21, acting via the Akt/NF- κ B pathway, in malignant transformation of HBE cells induced by cigarette smoke extract. <i>Toxicology Letters</i> , 2018, 289, 14-21.	0.4	20
74	Functional genetic variants of <i>c-Jun</i> and their interaction with smoking and drinking increase the susceptibility to lung cancer in southern and eastern Chinese. <i>International Journal of Cancer</i> , 2012, 131, E744-58.	2.3	19
75	The functional polymorphism of NBS1 p.Glu185Gln is associated with an increased risk of lung cancer in Chinese populations: Case-control and a meta-analysis. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2014, 770, 61-68.	0.4	18
76	MALAT1 via microRNA-17 regulation of insulin transcription is involved in the dysfunction of pancreatic β cells induced by cigarette smoke extract. <i>Journal of Cellular Physiology</i> , 2018, 233, 8862-8873.	2.0	17
77	Long noncoding RNA DNAJC3-AS1 promotes osteosarcoma progression via its sense cognate gene DNAJC3. <i>Cancer Medicine</i> , 2019, 8, 761-772.	1.3	17
78	A functional CNVR_3425.1 damping lincRNA FENDRR increases lifetime risk of lung cancer and COPD in Chinese. <i>Carcinogenesis</i> , 2018, 39, 347-359.	1.3	16
79	The effect of functional <i>MAPKAPK2</i> copy number variation CNV-30450 on elevating nasopharyngeal carcinoma risk is modulated by EBV infection. <i>Carcinogenesis</i> , 2014, 35, 46-52.	1.3	15
80	Cumulative evidence for relationships between multiple variants in the <i>VTI1A</i> and <i>TCF7L2</i> genes and cancer incidence. <i>International Journal of Cancer</i> , 2018, 142, 498-513.	2.3	15
81	Functional annotation of melanoma risk loci identifies novel susceptibility genes. <i>Carcinogenesis</i> , 2020, 41, 452-457.	1.3	15
82	A micropeptide XBP1SBM encoded by lncRNA promotes angiogenesis and metastasis of TNBC via XBP1s pathway. <i>Oncogene</i> , 2022, 41, 2163-2172.	2.6	15
83	The polymorphisms and haplotypes of WWOX gene are associated with the risk of lung cancer in southern and eastern chinese populations. <i>Molecular Carcinogenesis</i> , 2013, 52, 19-27.	1.3	14
84	<i>Sipa1</i> promoter polymorphism predicts risk and metastasis of lung cancer in Chinese. <i>Molecular Carcinogenesis</i> , 2013, 52, 110-117.	1.3	14
85	Genetic variant in the 3'-untranslated region of VEGFR1 gene influences chronic obstructive pulmonary disease and lung cancer development in Chinese population. <i>Mutagenesis</i> , 2014, 29, 311-317.	1.0	14
86	The MKK7 p.Glu116Lys Rare Variant Serves as a Predictor for Lung Cancer Risk and Prognosis in Chinese. <i>PLoS Genetics</i> , 2016, 12, e1005955.	1.5	14
87	Upregulation of lncRNA FEZF-AS1 is associated with advanced clinical stages and family history of cancer in patients with NSCLC. <i>Pathology Research and Practice</i> , 2018, 214, 857-861.	1.0	14
88	Willingness and influential factors of parents to vaccinate their children with novel inactivated enterovirus 71 vaccines in Guangzhou, China. <i>Vaccine</i> , 2018, 36, 3772-3778.	1.7	14
89	Long noncoding RNA PANDAR inhibits the development of lung cancer by regulating autophagy and apoptosis pathways. <i>Journal of Cancer</i> , 2020, 11, 4783-4790.	1.2	14
90	Long non-coding RNA AGER-1 inhibits colorectal cancer progression through sponging miR-182. <i>International Journal of Biological Markers</i> , 2020, 35, 10-18.	0.7	14

#	ARTICLE	IF	CITATIONS
91	A functional polymorphism in the promoter of ERK5 gene interacts with tobacco smoking to increase the risk of lung cancer in Chinese populations. <i>Mutagenesis</i> , 2013, 28, 561-567.	1.0	13
92	Identification of novel prognosis-related genes associated with cancer using integrative network analysis. <i>Scientific Reports</i> , 2018, 8, 3233.	1.6	12
93	X chromosome-linked long noncoding RNA <i>lincXLEC1</i> regulates <i>c-Myc</i> -dependent cell growth by collaborating with MBP1 in endometrial cancer. <i>International Journal of Cancer</i> , 2019, 145, 927-940.	2.3	12
94	Discovery of a novel linc01125 isoform in serum exosomes as a promising biomarker for NSCLC diagnosis and survival assessment. <i>Carcinogenesis</i> , 2021, 42, 831-841.	1.3	12
95	Functional Genetic Polymorphisms in PP2A Subunit Genes Confer Increased Risks of Lung Cancer in Southern and Eastern Chinese. <i>PLoS ONE</i> , 2013, 8, e77285.	1.1	12
96	Functional variants of -1318T > G and -673C > T in c-Jun promoter region associated with increased colorectal cancer risk by elevating promoter activity. <i>Carcinogenesis</i> , 2011, 32, 1043-1049.	1.3	11
97	A genetic variation of the p38 ^{Î²} promoter region is correlated with an increased risk of sporadic colorectal cancer. <i>Oncology Letters</i> , 2013, 6, 3-8.	0.8	11
98	A Functional Polymorphism in the 3'-UTR of PXR Interacts with Smoking to Increase Lung Cancer Risk in Southern and Eastern Chinese Smoker. <i>International Journal of Molecular Sciences</i> , 2014, 15, 17457-17468.	1.8	11
99	In type 2 diabetes induced by cigarette smoking, activation of p38 MAPK is involved in pancreatic Î²-cell apoptosis. <i>Environmental Science and Pollution Research</i> , 2018, 25, 9817-9827.	2.7	11
100	Salt-inducible kinases inhibitor HG-9-91-01 targets RIPK3 kinase activity to alleviate necroptosis-mediated inflammatory injury. <i>Cell Death and Disease</i> , 2022, 13, 188.	2.7	11
101	Association of nsv823469 copy number loss with decreased risk of chronic obstructive pulmonary disease and pulmonary function in Chinese. <i>Scientific Reports</i> , 2017, 7, 40060.	1.6	10
102	Upregulation of long noncoding RNA RAB11B-AS1 promotes tumor metastasis and predicts poor prognosis in lung cancer. <i>Annals of Translational Medicine</i> , 2020, 8, 582-582.	0.7	9
103	The Functional Copy Number Variation-67048 in <i>WWOX</i> Contributes to Increased Risk of COPD in Southern and Eastern Chinese. <i>COPD: Journal of Chronic Obstructive Pulmonary Disease</i> , 2015, 12, 494-501.	0.7	8
104	The role of <i>WWOX</i> polymorphisms on COPD susceptibility and pulmonary function traits in Chinese: a case-control study and family-based analysis. <i>Scientific Reports</i> , 2016, 6, 21716.	1.6	8
105	Study Design and Interim Outcomes of Guangzhou Institute of Respiratory Disease COPD Biobank. <i>COPD: Journal of Chronic Obstructive Pulmonary Disease</i> , 2016, 13, 203-213.	0.7	8
106	CMGene: A literature-based database and knowledge resource for cancer metastasis genes. <i>Journal of Genetics and Genomics</i> , 2017, 44, 277-279.	1.7	8
107	Imputation-based association analyses identify new lung cancer susceptibility variants in <i>CDK6</i> and <i>SH3RF1</i> and their interactions with smoking in Chinese populations. <i>Carcinogenesis</i> , 2013, 34, 2010-2016.	1.3	7
108	Polymorphisms of <i>NFÎ±1</i> and <i>Î±2</i> and Their Synergistic Effect on Nasopharyngeal Carcinoma Susceptibility. <i>BioMed Research International</i> , 2015, 2015, 1-9.	0.9	7

#	ARTICLE	IF	CITATIONS
109	Variants in the <i>PSCA</i> gene associated with risk of cancer and nonneoplastic diseases: systematic research synopsis, meta-analysis and epidemiological evidence. <i>Carcinogenesis</i> , 2019, 40, 70-83.	1.3	7
110	A Newfound Association between MDC1 Functional Polymorphism and Lung Cancer Risk in Chinese. <i>PLoS ONE</i> , 2014, 9, e106794.	1.1	7
111	Transmission dynamics and successful control measures of SARS-CoV-2 in the mega-size city of Guangzhou, China. <i>Medicine (United States)</i> , 2021, 100, e27846.	0.4	7
112	Clinical Features of Patients with Bronchiectasis with Comorbid Chronic Obstructive Pulmonary Disease in China. <i>Medical Science Monitor</i> , 2019, 25, 6805-6811.	0.5	6
113	Down expression of Inc-BMP1-1 decreases that of Caveolin-1 is associated with the lung cancer susceptibility and cigarette smoking history. <i>Aging</i> , 2020, 12, 462-480.	1.4	6
114	Functional Polymorphism in the MSI1 Gene Promoter Confers a Decreased Risk of Lung Cancer in Chinese by Reducing MSI1 Expression. <i>Current Genomics</i> , 2018, 19, 375-383.	0.7	6
115	Long non-coding RNA LSAMP-1 is down-regulated in non-small cell lung cancer and predicts a poor prognosis. <i>Cancer Cell International</i> , 2022, 22, 181.	1.8	6
116	A functional genetic variant in fragile-site gene FATS modulates the risk of breast cancer in triparous women. <i>BMC Cancer</i> , 2015, 15, 559.	1.1	5
117	Transmission dynamics of SARS-CoV-2 in a mid-size city of China. <i>BMC Infectious Diseases</i> , 2021, 21, 793.	1.3	5
118	Mitogen/Extracellular Signal-Regulated Kinase Kinase-5 Promoter Region Polymorphisms Affect the Risk of Sporadic Colorectal Cancer in a Southern Chinese Population. <i>DNA and Cell Biology</i> , 2012, 31, 342-349.	0.9	4
119	Integrative analysis to identify oncogenic gene expression changes associated with copy number variations of enhancer in ovarian cancer. <i>Oncotarget</i> , 2017, 8, 91558-91567.	0.8	4
120	Effect of EME1 exon variant Ile350Thr on risk and early onset of breast cancer in southern Chinese women. <i>Journal of Biomedical Research</i> , 2013, 27, 193-201.	0.7	4
121	Long Noncoding RNA PRRG4-4 Promotes Viability, Cell Cycle, Migration, and Invasion in Lung Cancer Cells. <i>DNA and Cell Biology</i> , 2018, 37, 953-966.	0.9	3
122	The pan-cancer analysis of gain-of-functional mutations to identify the common oncogenic signatures in multiple cancers. <i>Gene</i> , 2019, 697, 57-66.	1.0	2
123	Rare variant of <i>MAP2K7</i> is associated with increased risk of COPD in southern and eastern Chinese. <i>Respirology</i> , 2017, 22, 691-698.	1.3	1
124	CIGene: a literature-based online resource for cancer initiation genes. <i>BMC Genomics</i> , 2018, 19, 552.	1.2	1
125	COPD. <i>Chest</i> , 2020, 157, 473-475.	0.4	1
126	Prognostic Value of Germline Copy Number Variants and Environmental Exposures in Non-small Cell Lung Cancer. <i>Frontiers in Genetics</i> , 2021, 12, 681857.	1.1	0

#	ARTICLE	IF	CITATIONS
127	The Polymorphism of EME1 Gene is Associated with an Increased Risk of Lung Cancer: A Case-Control Study from Chinese Population. Journal of Cancer Research Updates, 2014, 3, 174-181.	0.3	0