

Mulong Du

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

118
papers

2,003
citations

23
h-index

40
g-index

126
ext. papers

2,639
ext. citations

7.8
avg, IF

4.9
L-index

#	Paper	IF	Citations
118	Pan-Cancer Analysis Revealed as a New Biomarker for Prognosis and Immunotherapy.. <i>Journal of Oncology</i> , 2022 , 2022, 3477148	4.5	3
117	LncRNA-422 suppresses the proliferation and growth of colorectal cancer cells by targeting SFPQ.. <i>Clinical and Translational Medicine</i> , 2022 , 12, e664	5.7	1
116	Association of genetic variants in autophagy-lysosome pathway genes with susceptibility and survival to prostate cancer. <i>Gene</i> , 2022 , 808, 145953	3.8	2
115	Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-eIF3h interaction.. <i>Molecular Cancer</i> , 2022 , 21, 49	42.1	6
114	Genetic variants in choline metabolism pathway are associated with the risk of bladder cancer in the Chinese population.. <i>Archives of Toxicology</i> , 2022 , 1	5.8	0
113	Evaluation of genetic variants in nucleosome remodeling and deacetylase (NuRD) complex subunits encoding genes and gastric cancer susceptibility.. <i>Archives of Toxicology</i> , 2022 , 1	5.8	0
112	Association between circulating vitamin E and ten common cancers: evidence from large-scale Mendelian randomization analysis and a longitudinal cohort study.. <i>BMC Medicine</i> , 2022 , 20, 168	11.4	2
111	Genetic variants in the Hedgehog signaling pathway genes are associated with gastric cancer risk in a Chinese Han population.. <i>Journal of Biomedical Research</i> , 2021 , 36, 22-31	1.5	
110	Fine Particulate Matter Induces Childhood Asthma Attacks via Extracellular Vesicle-Packaged Let-7i-5p-Mediated Modulation of the MAPK Signaling Pathway. <i>Advanced Science</i> , 2021 , e2102460	13.6	2
109	Global internet search trends related to gastrointestinal symptoms predict regional COVID-19 outbreaks. <i>Journal of Infection</i> , 2021 ,	18.9	2
108	Functional variants of RPS6KB1 and PIK3R1 in the autophagy pathway genes and risk of bladder cancer. <i>Archives of Toxicology</i> , 2021 , 1	5.8	0
107	Genetic variations in the CTLA-4 immune checkpoint pathway are associated with colon cancer risk, prognosis, and immune infiltration via regulation of IQCB1 expression. <i>Archives of Toxicology</i> , 2021 , 95, 2053-2063	5.8	0
106	Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. <i>Archives of Toxicology</i> , 2021 , 95, 2523-2532	5.8	1
105	Integrative omics provide biological and clinical insights into acute respiratory distress syndrome. <i>Intensive Care Medicine</i> , 2021 , 47, 761-771	14.5	2
104	Vaping-induced metabolomic signatures in the circulation of mice are driven by device type, e-liquid, exposure duration and sex. <i>ERJ Open Research</i> , 2021 , 7,	3.5	1
103	Genetic variants of CHEK1, PRIM2 and CDK6 in the mitotic phase-related pathway are associated with nonsmall cell lung cancer survival. <i>International Journal of Cancer</i> , 2021 , 149, 1302-1312	7.5	2
102	The biogenesis and biological function of PIWI-interacting RNA in cancer. <i>Journal of Hematology and Oncology</i> , 2021 , 14, 93	22.4	3

101	Genetic variants in Hippo signalling pathway-related genes affect the risk of colorectal cancer. <i>Archives of Toxicology</i> , 2021 , 95, 271-281	5.8	2
100	Plasma Insulin-like Growth Factor Binding Protein 7 Contributes Causally to ARDS 28-Day Mortality: Evidence From Multistage Mendelian Randomization. <i>Chest</i> , 2021 , 159, 1007-1018	5.3	3
99	Metabolomics identifying biomarkers of PM exposure for vulnerable population: based on a prospective cohort study. <i>Environmental Science and Pollution Research</i> , 2021 , 28, 14586-14596	5.1	5
98	Systematic evaluation of the effects of genetic variants on PIWI-interacting RNA expression across 33 cancer types. <i>Nucleic Acids Research</i> , 2021 , 49, 90-97	20.1	8
97	A prospective study of the associations among fine particulate matter, genetic variants, and the risk of colorectal cancer. <i>Environment International</i> , 2021 , 147, 106309	12.9	4
96	Genetic variants in N6-methyladenosine are associated with bladder cancer risk in the Chinese population. <i>Archives of Toxicology</i> , 2021 , 95, 299-309	5.8	5
95	Circular RNAs in body fluids as cancer biomarkers: the new frontier of liquid biopsies. <i>Molecular Cancer</i> , 2021 , 20, 13	42.1	64
94	Identification of low-frequency variants of UGT1A3 associated with bladder cancer risk by next-generation sequencing. <i>Oncogene</i> , 2021 , 40, 2382-2394	9.2	3
93	Performance of a Machine Learning Algorithm Using Electronic Health Record Data to Identify and Estimate Survival in a Longitudinal Cohort of Patients With Lung Cancer. <i>JAMA Network Open</i> , 2021 , 4, e2114723	10.4	11
92	Identification of common genetic variants associated with serum concentrations of p, pTDDE in non-occupational populations in eastern China. <i>Environment International</i> , 2021 , 152, 106507	12.9	0
91	SARS-CoV-2 Impairs Dendritic Cells and Regulates DC-SIGN Gene Expression in Tissues. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	4
90	Genetic variants in mA regulators are associated with gastric cancer risk. <i>Archives of Toxicology</i> , 2021 , 95, 1081-1088	5.8	7
89	Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020 , 41, 8-17	4.6	30
88	Functional genetic variant of HSD17B12 in the fatty acid biosynthesis pathway predicts the outcome of colorectal cancer. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 14160-14170	5.6	1
87	Remote modulation of lncRNA by risk variant at 16p13 underlying genetic susceptibility to gastric cancer. <i>Science Advances</i> , 2020 , 6, eaay5525	14.3	10
86	Genetic variants in Ras/Raf/MEK/ERK pathway are associated with gastric cancer risk in Chinese Han population. <i>Archives of Toxicology</i> , 2020 , 94, 2683-2690	5.8	2
85	Genetic Variant in Long Non-Coding RNA Modulates Its Expression and Predicts Renal Cell Carcinoma Susceptibility and Mortality. <i>Frontiers in Oncology</i> , 2020 , 10, 785	5.3	3
84	Identification of genetic features associated with fine particulate matter (PM2.5) modulated DNA damage using improved random forest analysis. <i>Gene</i> , 2020 , 740, 144570	3.8	0

83	MUC1 is associated with TFF2 methylation in gastric cancer. <i>Clinical Epigenetics</i> , 2020 , 12, 37	7.7	3
82	Genetic variations in Hippo pathway genes influence bladder cancer risk in a Chinese population. <i>Archives of Toxicology</i> , 2020 , 94, 785-794	5.8	8
81	A transcriptomic study for identifying cardia- and non-cardia-specific gastric cancer prognostic factors using genetic algorithm-based methods. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 9457-9465	5.6	2
80	Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. <i>Environment International</i> , 2020 , 137, 105543	12.9	7
79	Epigenomic analysis of 5-hydroxymethylcytosine (5hmC) reveals novel DNA methylation markers for lung cancers. <i>Neoplasia</i> , 2020 , 22, 154-161	6.4	8
78	Novel CpG-SNPs in the gastric acid secretion pathway GNAI3 and susceptibility to gastric cancer. <i>Gene</i> , 2020 , 736, 144447	3.8	3
77	Multiomics Evaluation of Gastrointestinal and Other Clinical Characteristics of COVID-19. <i>Gastroenterology</i> , 2020 , 158, 2298-2301.e7	13.3	83
76	Urethral Lift as a Safe and Effective Procedure for Prostatic Hyplasia Population: A Systematic Review and Meta-Analysis. <i>Frontiers in Surgery</i> , 2020 , 7, 598728	2.3	3
75	Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. <i>Archives of Toxicology</i> , 2020 , 94, 325-333	5.8	2
74	Explaining the Genetic Causality for Complex Phenotype via Deep Association Kernel Learning. <i>Patterns</i> , 2020 , 1, 100057	5.1	3
73	Tumor evolutionary trajectories during the acquisition of invasiveness in early stage lung adenocarcinoma. <i>Nature Communications</i> , 2020 , 11, 6083	17.4	5
72	The value of renal score in both determining surgical strategies and predicting complications for renal cell carcinoma: A systematic review and meta-analysis. <i>Cancer Medicine</i> , 2020 , 9, 3944-3953	4.8	4
71	Long non-coding RNA FLJ22763 is involved in the progression and prognosis of gastric cancer. <i>Gene</i> , 2019 , 693, 84-91	3.8	8
70	Genetic variants in p53 signaling pathway genes predict chemotherapy efficacy in colorectal cancer. <i>Cancer Medicine</i> , 2019 , 8, 3428-3436	4.8	4
69	Genetic variants in a long noncoding RNA related to Sunitinib Resistance predict risk and survival of patients with renal cell carcinoma. <i>Cancer Medicine</i> , 2019 , 8, 2886-2896	4.8	4
68	Genetic variants in RPA1 associated with the response to oxaliplatin-based chemotherapy in colorectal cancer. <i>Journal of Gastroenterology</i> , 2019 , 54, 939-949	6.9	8
67	Combinations of single nucleotide polymorphisms identified in genome-wide association studies determine risk for colorectal cancer. <i>International Journal of Cancer</i> , 2019 , 145, 2661-2669	7.5	12
66	A genetic variation in the CpG island of pseudogene GBAP1 promoter is associated with gastric cancer susceptibility. <i>Cancer</i> , 2019 , 125, 2465-2473	6.4	11

65	A genetic variant located in the miR-532-5p-binding site of TGFBR1 is associated with the colorectal cancer risk. <i>Journal of Gastroenterology</i> , 2019 , 54, 141-148	6.9	9
64	Vitamin B intake reduces the risk for colorectal cancer: a dose-response analysis. <i>European Journal of Nutrition</i> , 2019 , 58, 1591-1602	5.2	5
63	Decoding tumor mutation burden and driver mutations in early stage lung adenocarcinoma using CT-based radiomics signature. <i>Thoracic Cancer</i> , 2019 , 10, 1904-1912	3.2	18
62	Robotic-Assisted Sentinel Lymph Node Mapping With Indocyanine Green in Pelvic Malignancies: A Systematic Review and Meta-Analysis. <i>Frontiers in Oncology</i> , 2019 , 9, 585	5.3	5
61	Efficacy of Ahmed Glaucoma Valve Implantation on Neovascular Glaucoma. <i>International Journal of Medical Sciences</i> , 2019 , 16, 1371-1376	3.7	11
60	Diagnostic Accuracy of Multi-Parametric Magnetic Resonance Imaging for Tumor Staging of Bladder Cancer: Meta-Analysis. <i>Frontiers in Oncology</i> , 2019 , 9, 981	5.3	8
59	The association between plasma fibrinogen levels and lung cancer: a meta-analysis. <i>Journal of Thoracic Disease</i> , 2019 , 11, 4492-4500	2.6	4
58	Genetic variants in SLC22A3 contribute to the susceptibility to colorectal cancer. <i>International Journal of Cancer</i> , 2019 , 145, 154-163	7.5	15
57	Genome-wide long non-coding RNAs identified a panel of novel plasma biomarkers for gastric cancer diagnosis. <i>Gastric Cancer</i> , 2019 , 22, 731-741	7.6	19
56	Genetic variant in miR-21 binding sites is associated with colorectal cancer risk. <i>Journal of Cellular and Molecular Medicine</i> , 2019 , 23, 2012-2019	5.6	6
55	Tagging SNPs in the HOTAIR gene are associated with bladder cancer risk in a Chinese population. <i>Gene</i> , 2018 , 664, 22-26	3.8	8
54	A Genetic Variant Located in Promoter Region Is Associated with Prognosis of Gastric Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018 , 27, 822-828	4	6
53	Evaluation of vulnerable PM-exposure individuals: a repeated-measure study in an elderly population. <i>Environmental Science and Pollution Research</i> , 2018 , 25, 11833-11840	5.1	6
52	Hypermethylation of EIF4E promoter is associated with early onset of gastric cancer. <i>Carcinogenesis</i> , 2018 , 39, 66-71	4.6	12
51	Polymorphism rs2682818 in miR-618 is associated with colorectal cancer susceptibility in a Han Chinese population. <i>Cancer Medicine</i> , 2018 , 7, 1194-1200	4.8	13
50	Genetic variants in XDH are associated with prognosis for gastric cancer in a Chinese population. <i>Gene</i> , 2018 , 663, 196-202	3.8	1
49	Probabilistic natural mapping of gene-level tests for genome-wide association studies. <i>Briefings in Bioinformatics</i> , 2018 , 19, 545-553	13.4	5
48	LncRNA and its genetic variant rs1902432 are associated with prostate cancer risk. <i>Journal of Cancer</i> , 2018 , 9, 1414-1420	4.5	22

47	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018 , 33, 82-88	8.8	5
46	LncRNA MT1JP functions as a ceRNA in regulating FBXW7 through competitively binding to miR-92a-3p in gastric cancer. <i>Molecular Cancer</i> , 2018 , 17, 87	42.1	166
45	Association of Antioxidative Enzymes Polymorphisms with Efficacy of Platin and Fluorouracil-Based Adjuvant Therapy in Gastric Cancer. <i>Cellular Physiology and Biochemistry</i> , 2018 , 48, 2247-2257	3.9	6
44	Body mass index (BMI) trajectories and risk of colorectal cancer in the PLCO cohort. <i>British Journal of Cancer</i> , 2018 , 119, 130-132	8.7	15
43	Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. <i>Gene</i> , 2018 , 673, 174-180	3.8	9
42	Mutant-allele fraction heterogeneity is associated with non-small cell lung cancer patient survival. <i>Oncology Letters</i> , 2018 , 15, 795-802	2.6	9
41	A functional polymorphism in TFF1 promoter is associated with the risk and prognosis of gastric cancer. <i>International Journal of Cancer</i> , 2018 , 142, 1805-1816	7.5	18
40	Exosome-transmitted long non-coding RNA PTENP1 suppresses bladder cancer progression. <i>Molecular Cancer</i> , 2018 , 17, 143	42.1	134
39	Genetic variants in PI3K/Akt/mTOR pathway genes contribute to gastric cancer risk. <i>Gene</i> , 2018 , 670, 130-135	3.8	12
38	Association study of genetic variants in estrogen metabolic pathway genes and colorectal cancer risk and survival. <i>Archives of Toxicology</i> , 2018 , 92, 1991-1999	5.8	12
37	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018 , 92, 2681-2689	5.8	24
36	KCNMA1 cooperating with PTK2 is a novel tumor suppressor in gastric cancer and is associated with disease outcome. <i>Molecular Cancer</i> , 2017 , 16, 46	42.1	24
35	A fast algorithm for Bayesian multi-locus model in genome-wide association studies. <i>Molecular Genetics and Genomics</i> , 2017 , 292, 923-934	3.1	4
34	Evaluation of genome-wide genotyping concordance between tumor tissues and peripheral blood. <i>Genomics</i> , 2017 , 109, 108-112	4.3	6
33	A seven-gene prognostic signature for rapid determination of head and neck squamous cell carcinoma survival. <i>Oncology Reports</i> , 2017 , 38, 3403-3411	3.5	21
32	Information transduction capacity reduces the uncertainties in annotation-free isoform discovery and quantification. <i>Nucleic Acids Research</i> , 2017 , 45, e143	20.1	6
31	Plasma Mesothelin as a Novel Diagnostic and Prognostic Biomarker in Colorectal Cancer. <i>Journal of Cancer</i> , 2017 , 8, 1355-1361	4.5	10
30	An inverse association between tea consumption and colorectal cancer risk. <i>Oncotarget</i> , 2017 , 8, 37367-37376	3.0	30

29	Rare variants in BRCA2 and CHEK2 are associated with the risk of urinary tract cancers. <i>Scientific Reports</i> , 2016 , 6, 33542	4.9	16
28	Choline and betaine consumption lowers cancer risk: a meta-analysis of epidemiologic studies. <i>Scientific Reports</i> , 2016 , 6, 35547	4.9	22
27	A functional variant in TP63 at 3q28 associated with bladder cancer risk by creating an miR-140-5p binding site. <i>International Journal of Cancer</i> , 2016 , 139, 65-74	7.5	19
26	Functional annotation of colorectal cancer susceptibility loci identifies MLH1 rs1800734 associated with MSI patients. <i>Gut</i> , 2016 , 65, 1227-8	19.2	11
25	Pri-miR-34b/c rs4938723 polymorphism contributes to acute lymphoblastic leukemia susceptibility in Chinese children. <i>Leukemia and Lymphoma</i> , 2016 , 57, 1436-41	1.9	23
24	Association Between MIF-AS rs755622 and Nephrolithiasis Risk in a Chinese Population. <i>Medical Science Monitor</i> , 2016 , 22, 563-8	3.2	2
23	Association of genetic variants in lncRNA H19 with risk of colorectal cancer in a Chinese population. <i>Oncotarget</i> , 2016 , 7, 25470-7	3.3	68
22	The association analysis of hOGG1 genetic variants and gastric cancer risk in a Chinese population. <i>Oncotarget</i> , 2016 , 7, 66061-66068	3.3	7
21	Common genetic variation in ETV6 is associated with colorectal cancer susceptibility. <i>Nature Communications</i> , 2016 , 7, 11478	17.4	45
20	Genome-Wide Association Study of Bladder Cancer in a Chinese Cohort Reveals a New Susceptibility Locus at 5q12.3. <i>Cancer Research</i> , 2016 , 76, 3277-84	10.1	29
19	Genetic variants in lncRNA H19 are associated with the risk of bladder cancer in a Chinese population. <i>Mutagenesis</i> , 2016 , 31, 531-8	2.8	60
18	Identification of a novel susceptibility locus at 16q23.1 associated with childhood acute lymphoblastic leukemia in Han Chinese. <i>Human Molecular Genetics</i> , 2016 , 25, 2873-2880	5.6	6
17	Genetic variants in noncoding PIWI-interacting RNA and colorectal cancer risk. <i>Cancer</i> , 2015 , 121, 2044-50	5.4	43
16	Genetic variants of H2AX gene were associated with PM2.5-modulated DNA damage levels in Chinese Han populations. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2015 , 778, 41-5	3.3	3
15	Personal exposure to PM2.5, genetic variants and DNA damage: a multi-center population-based study in Chinese. <i>Toxicology Letters</i> , 2015 , 235, 172-8	4.4	26
14	Genome-wide analysis of long noncoding RNA signature in human colorectal cancer. <i>Gene</i> , 2015 , 556, 227-34	3.8	53
13	Identification of novel piRNAs in bladder cancer. <i>Cancer Letters</i> , 2015 , 356, 561-7	9.9	91
12	Folic acid supplements and colorectal cancer risk: meta-analysis of randomized controlled trials. <i>Scientific Reports</i> , 2015 , 5, 12044	4.9	39

11	The association analysis of lncRNA HOTAIR genetic variants and gastric cancer risk in a Chinese population. <i>Oncotarget</i> , 2015 , 6, 31255-62	3.3	91
10	The prognostic significance of HOTAIR for predicting clinical outcome in patients with digestive system tumors. <i>Journal of Cancer Research and Clinical Oncology</i> , 2015 , 141, 2139-45	4.9	29
9	Circulating miR-497 and miR-663b in plasma are potential novel biomarkers for bladder cancer. <i>Scientific Reports</i> , 2015 , 5, 10437	4.9	87
8	MDM2 SNP309 polymorphism is associated with colorectal cancer risk. <i>Scientific Reports</i> , 2014 , 4, 4851	4.9	12
7	Hsa-miR-196a2 polymorphism increases the risk of acute lymphoblastic leukemia in Chinese children. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2014 , 759, 16-21	3.3	28
6	Clinical potential role of circulating microRNAs in early diagnosis of colorectal cancer patients. <i>Carcinogenesis</i> , 2014 , 35, 2723-30	4.6	51
5	Genetic variants in SMARC genes are associated with DNA damage levels in Chinese population. <i>Toxicology Letters</i> , 2014 , 229, 327-32	4.4	5
4	Association study between XPG Asp1104His polymorphism and colorectal cancer risk in a Chinese population. <i>Scientific Reports</i> , 2014 , 4, 6700	4.9	20
3	Clinical significance of POU5F1P1 rs10505477 polymorphism in Chinese gastric cancer patients receiving cisplatin-based chemotherapy after surgical resection. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 12764-77	6.3	15
2	Associations of NR5A2 gene polymorphisms with the clinicopathological characteristics and survival of gastric cancer. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 22902-17	6.3	8
1	Genetic variations in microRNAs and the risk and survival of renal cell cancer. <i>Carcinogenesis</i> , 2014 , 35, 1629-35	4.6	41